Supplementary Materials: A Phylogenetic Survey on the Structure of the HIV-1 Leader RNA Domain That Encodes the Splice Donor Signal

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                                        5          15         25         35         45         55         65         75         85         95
B.FR.83.HXB2_LAI_IIIB_BRU.K034      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
A1.CD.02.02CD_KTB035.AM000055       TAGTGTGTGC CCTTCTGTTG TGTGC----- ---------C TTTGGTAA-C TAAGGATCCC TCAGCCCCCT --TTAAAAGA A-GTGAA--- AATCTCTACC
A1.CD.97.97CD_KCC2.AM000053         TGGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACC ---TTAGCAG A-GTGAAA-- AATCTCTAGC
A1.ES.05.X1608_8.FJ670519           TAGTGTGTGC CCGTCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --GGAGACC- --GTGCGAA- AATCTCTAGC
A1.IT.02.60000.EU861977             TAGTGTGTGC CCGTCTGTTG TGTRA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG T-GTAAAA-- AATCTCTAGC
A1.KE.01.ML1901PCR2.EU110095        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1.KE.01.ML1945.EU110088            ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1.KE.02.ML1990.EU110092            ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1.KE.02.ML2014.EU110094            ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1.KE.86.ML170_1986.AF539405        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1.KE.94.Q23_17.AF004885            TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGG T-GTGAA--- AATCTCTAGC
A1.KE.97.ML013_2.AY322185           TAGTGTGTGC CCGTCTGTTG TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGG T-GTAAA--- AATCTCTAGC
A1.KE.97.ML605_3.AY322190           TAGTGTGTGC CCGTCTGTTG TCTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG T-GTAAA--- AATCTCTAGC
A1.KE.97.ML752.AY322193             TAGTGTGTGC CCATTTGTTG TAAGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGG T-GTAAA--- AATCTCTAGC
A1.RU.00.RU00051.EF545108           ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1.RU.05.RU_560_1125_JA.JQ2928      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1.RU.06.RU_915_1016.JQ292896       TAGTGTGTGC CCGTCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGCGG T-GTAAA--- AATCTCTAGC
A1.RU.06.RU_915_1035.JQ292897       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGCGG T-GTAAA--- AATCTCTAGC
A1.RU.06.RU_915_1038.JQ292898       TAGTGTGTGC CCGTCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGCGG T-GTAAA--- AATCTCTAGC
A1.RU.06.RU_915_1041.JQ292899       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGCGG T-GTAAA--- AATCTCTAGC
A1.RU.06.RU_SP_B_049.JQ292900       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGCGG T-GTAAA--- AATCTCTAGC
A1.RU.07.Irkutsk_5.JQ292891         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1.RU.08.PokA1Ru.FJ864679           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGCGG T-GTAAA--- AATCTCTAGC
A1.RU.08.RUA001.JQ292893            TAGTGTGTGC CCGTCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
A1.RU.08.RUA007.JQ292894            TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TTTGGTAA-C TAGAGATCCC TCAGACCACT --YTAGGCGG T-GTAAA--- AATCTCTAGC
A1.RU.10.10RU6617.JX500696          TAGTGTGTGC CCATCTGTTG TATGA----- ---------C TCTGGTGA-C TAGAGATCCC TCAGACCACT --CTAGGCGG T-GTAA---- AATCTCTAGC
A1.RU.10.10RU6792.JX500695          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGCAG T-GTAAA--- AATCTCTAGC
A1.RU.11.11RU6950.JX500694          TAGTGTGTGC CCGTCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGCGG T-GTAAA--- AATCTCTAGC
A1.RW.07.pR463F.JX236677            TAGTGTGTGC CCGTCTGTTG TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
A1.RW.07.pR880F.JX236678            CAGTGTGTGC CCGTCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG T-GCAAA--- AATCTCTAGC
A1.RW.92.92RW008.AB253421           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGATTG A-GTAAA--- AATCTCTAGC
A1.RW.92.92RW025A.AB287376          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
A1.RW.93.93RW037A.AB287379          TAGTGTGTGC CCGTCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
A1.SE.95.UGSE8131.AF107771          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----A----- AATCTCTAGC
A1.TZ.97.97TZ03.AF361873            ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1.UA.00.98UA0116.AF413987          TAGTGTGTGC CCGTCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTACT --CTAGGCAG T-GTAAA--- AATCTCTAGC
A1.UG.07.p191084.JX236669           TAGTGTGTGC CCGTCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTACT --CTAGACAG T-GTAAA--- AATCTCTAGC
A1.UG.07.p191845.JX236671           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG T-ATAAA--- AATCTCTAGC
A1.UG.07.p9004SDM.JX236676          CAGTGTGTGC CCGTCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTT A-GTAAA--- AATCTCTAGC
A1.UG.85.U455_U455A.M62320          TAGTGTGTGC CCGTCTGTTA TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --ATAGACTG T-GTAAA--- AATCTCTAGC
A1.UG.92.92UG037.AB253429           TAGTGTGTGC CCGTCTGTTT TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
A1.UG.92.UG029.AB098333             TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGAAGG T-GTAAA--- AATCTCTAGC
A1.UG.x.UG031.AB098331              TAGTGTGTGC CCGTCTGTTG TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
A1.UG.x.UG275.AB485632              CAGTGTGTGC CCGTCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGATCACT --CTAGACTG T-GTAAA--- AATCTCTAGC
A1.ZA.04.04ZASK162B1.DQ396400       ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAGA--- AATCTCTAGC
A2.CD.97.97CDKTB48.AF286238         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A2.CY.94.94CY017_41.AF286237        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ------TTGA
B.AU.86.MBC200.AF042100             TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.AU.87.MBC925.AF042101             TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.AU.95.C24.AF538304                TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAG-- AATCTCTAGC
B.AU.95.C42.AF538305                TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCAAT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.AU.95.C76.AF538306                TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.AU.95.C92.AF538307                TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAG-- AATCTCTAGC
B.AU.95.MBCC54.AF042103             ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.AU.96.MBCC98.AF042104             ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.AU.96.MBCD36.AF042105             ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.AU.99.1181.AF538302               TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCCTT --TAATTCCG T-GTAGAA-- AATCTCTAGC
B.AU.x.2870718.AY857022             ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.AU.x.9125091.AY857165             ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.AU.x.VH_VHPCR.AF146728            ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.BO.09.DEMB09BO001.JX140656        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.BR.02.02BR008.DQ358808            ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------AG T-GTGGAA-- AATCTCTAGC
B.BR.02.02BR011.DQ358809            TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TAAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.02.02BR013.DQ358810            TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTCGTATAC TAGAGATCCC TCAGACC-TT --TTAGTCAG T-GTGGAAA- AATCTCTAGC
B.BR.02.02BR1013.JN692432           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --AAAGTCAG T-GTGGAAA- AATCTCTAGC
B.BR.02.02BR2025.JN692435           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.02.02BR2032.JN692439           ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.BR.02.02BR2033.JN692440           AGGTGTGTGC CCGTCTGTTG TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATC --TTAGACAG T-GGAAA--- AATCTCTAGC
B.BR.02.02BR2041.JN692443           ATAGTAGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCC-- --TATTAGTC A-GTGTGGAA AATCTCTAGC
B.BR.02.02BR2042.JN692444           ---------C CCGTCTGATT TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACTT --TTAGTTCG T-GTGGAA-- AATCTCTAGC
B.BR.02.04BR1064.JN692433           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACCT --TTAGTCAG T-GTGGAAA- AATCTCTAGC
B.BR.02.05BR1094.JN692470           ------GTGC CCGTCTGATT TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACTT --TTAGTTCG T-GTGGAA-- AATCTCTAGC
B.BR.03.03BR1020.JN692445           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --AAAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.03.03BR1046.JN692447           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCAC --TTAGTCCG A-GTGAAA-- AATCTCTAGC
B.BR.03.BREPM1023.EF637057          ---------- ---------- ---------- ---------- TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.03.BREPM1024.EF637056          ---------- ---------- ---------- ---------- ---------- ---------- -------AAT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.03.BREPM1027.EF637054          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- -ATCTCTAGC
B.BR.03.BREPM1028.EF637053          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.BR.03.BREPM1032.EF637051          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.BR.03.BREPM1033.EF637050          ---------- ---------- ---------- ---------- ---------- ---------- -----CCATT --TAGATCAG T-GTGGAA-- AATCTCTAGC
B.BR.03.BREPM1035.EF637049          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.BR.03.BREPM1038.EF637048          ---------- ---------- ---------- ---------- -ATGGTAA-C TAGAGATCCC TCAGACCATT --CTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.03.BREPM1040.EF637047          ---------- ---------- ---------- ---------- ---------- ---------- --AGACCAAT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.03.BREPM2012.EF637046          ---------- ---------- ---------- ---------- ---------- ---------- -------TT- --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.04.04BR1047.JN692450           ---------- ---------- ---------- ---------- ---------- ---------- -CAGACCATT --TTAGTCAG T-GTGAA--- AATCTCTAGC
B.BR.04.04BR1049.JN692451           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
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B.BR.04.04BR1051.JN692452        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACTT --GTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.04.04BR1054.JN692453        ---------- ---------- ---------- ---------- ---------C TGGAGATCCC TCAGACCCAT --TTAGTCA- --GTGTGGAA AATCTCTAGC
B.BR.04.04BR1055.JN692454        TAGTGTGTGC CCGTCTGTTG TGTAA----- ---------C TCTAGTAA-C TAAAAATCCC TCAAACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.04.04BR1057.JN692455        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --CAGTCAGT --GCGGAAA- AATCTCTAGC
B.BR.04.04BR1068.JN692457        ---------- ---------- ---------- ---------- ---------C TAGAGATCCC TCAGACATTG --TAGTCAGT --GTGGAAA- AATCTCTAGC
B.BR.04.BREPM1066.FJ195090       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACT- --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.04.BREPM1070.FJ195086       ---------- --TTCGGTGG TGTGA----- ---------T TCTGGTAC-C TAGAGTTCCT TCAGACCTTA --TTGTTCAG T-TGGGAA-- ATTCTTTGCC
B.BR.05.05BR1077.JN692460        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.BR.05.05BR1078.JN692461        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.BR.05.05BR1079.JN692462        ---------C CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.05.05BR1080.JN692463        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.BR.05.05BR1082.JN692465        TAGTGTGTGC CCGTCTGTTG TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.05.05BR1089.JN692467        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.BR.05.05BR1092.JN692468        TTATGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.05.05BR1095.JN692471        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --GTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.05.05BR1101.JN692473        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTCTT --CTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.05.05BR1104.JN692474        TTATGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TAAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.05.05BR1107.JN692475        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TAAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.05.BREPM1081.FJ195091       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACTT --TTAATCCG T-GTGGAA-- AATCTCTAGC
B.BR.05.BREPM1084.FJ195088       ---------- ---------- ---------- ---------- ---------- -------CCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.05.BREPM1093.FJ195089       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.BR.06.06BR1115.JN692479        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG A-GTGGAA-- AATCTCTAGC
B.BR.06.06BR1119.JN692480        ---------- ---------- ---------- ---------- ---------- --GAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CA.06.502_1027_wg01.JF320413   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAAAC TAGAGATCCC TCAGACCC-- --TTTTTAGT CAGTGTGGAA AATCTCTAGC
B.CA.06.502_1799_FL02.JF320427   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCTTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
B.CA.07.502_1191_03.JF320424     ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CA.96.WC10C_4.AY314056         -AGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CA.97.CANB3FULL.AY779553       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.CH.03.HIV_CH_BID_V4470_2003.   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.CN.01.CNHN24.AY180905          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----GATT-- AATCTCTAGC
B.CN.02.02HNsc11.DQ007903        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGAACCTA --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.03.SHXDC0081.JF932492       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCCTA --ATAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.05.05CNHB_hp3.DQ990880      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT ---TAGTTTG T-GTGGAA-- AATCTCTAGC
B.CN.06.CC056.JF932482           ---------C CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.07.AH070011.JF932468        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.CN.07.AH070014.JF932469        ---------- CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.07.AH070017.JF932470        ---------- ---------- ----A----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.07.AH070018.JF932471        --------GC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGAACAGT --TTAGTTAG T-GTGGAA-- AATCTCTAGC
B.CN.07.AH070057.JF932472        ---------- -CGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.07.BJ070030.JF932473        --GTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACTT --CTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.07.CBJC261.JF932474         --------GC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGAACCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.07.CBJC392.JF932475         --------GC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCAT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.07.CBJC394.JF932476         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
B.CN.07.CBJC396.JF932477         ---------C CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGAACCTT --TTAGTTAG T-GTGGAA-- AATCTCTAGC
B.CN.07.GS070017.JF932484        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.CN.07.GZ070002.JF932485        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.CN.07.GZ070030.JF932486        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.07.HB070006.JF932487        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.CN.07.HB070022.JF932488        --------GC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.07.HB070035.JF932489        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAAA- AATCTCTAGC
B.CN.07.JL070038.JF932490        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.CN.07.SX070080.JF932493        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.CN.07.hb070025.JF932499        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.07.hen1345.JF932500         ------GTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.08.CBJC476.JF932478         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCTTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.08.CBJC489.JF932479         ---------C CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGAACCGT --GTAGTTAG T-GTGGAA-- AATCTCTAGC
B.CN.08.CBJC500.JF932480         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.08.CBJC503.JF932481         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTCTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.08.cbjc468.JF932498         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCCT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.09.YN09P0014.JF932494       ---------C CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.09.ZK042.JF932497           ---------C CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGAACCTT --TTAGTCTG T-GTGGAA-- AATCTCTAGC
B.CN.10.DEMB10CN002.JX140658     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.CN.98.YN9802.JF932495          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.CN.98.YN9838.JF932496          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.CN.99.plwj.GU177863            TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
B.CN.x.RL42.U71182               ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- AATCTCTAGC
B.CU.99.Cu19.AY586542            TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTC --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.CU.99.Cu43.AY586543            TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTC --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.DE.86.D31.U43096               TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.DE.86.HAN.U43141               TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGATCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.DK.01.CTL_016.EF514704         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.01.CTL_017.EF514705         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------G
B.DK.01.CTL_018.EF514706         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.01.CTL_023.EF514707         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.01.CTL_030.EF514708         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------TT
B.DK.01.CTL_033.EF514709         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.01.CTL_035.EF514710         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.01.CTL_041.EF514711         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.01.CTL_043.EF514712         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.04.PMVL_012.EF514699        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.04.PMVL_013.EF514700        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.04.PMVL_018.EF514697        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.04.PMVL_025.EF514702        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.04.PMVL_027.EF514698        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.04.PMVL_039.EF514703        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.04.PMVL_049.EF514701        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.07.PMVL_011.FJ694790        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.ES.05.X1890.EU786672           ------GTGC CCGTCTGTTG TTTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCCAT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.ES.06.X1959.EU786675           ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----TCTAGC
B.ES.06.X2102.EU786677           -------TGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.ES.07.X2149.EU786678           ------GTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.ES.07.X2210_3.EU786679         -------TGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACAG --CTAGTCAG T-GTGGAA-- TATCTCTAGC
B.ES.07.X2231.EU786680           -------TGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
B.ES.08.ES_X2515_3.GQ372988      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --T-AGTCAG T-GTGGAA-- AATCTCTAGC
B.ES.08.ES_X2556_3.GQ372990      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.ES.08.P2008.FJ670531           -AAGGGTTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TAAGTCAG T-GTGGAA-- AATCTCTAGC
B.ES.08.X2425_2.FJ670525         -------TGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACTT --GTAGTCTG T-GTAGAA-- AATCTCTAGC
B.ES.08.X2510_2.FJ853622         -----TGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.ES.08.X2555_2.GU362883         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.ES.08.X2574_2.GU362886         -AGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.ES.09.P2149_3.GU362881         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TAGTCAG T-GTGGAAA- AATCTCTAGC
B.ES.09.X2689_2.GU362885         -AGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTCTT --GTAGTTAG T-GTGGAA-- AATCTCTAGC
B.ES.89.U61.DQ854716             TAGTGTGTGC CCGTCTATTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TTCAGACCAA ATTTAGTCAG T-GTGAAA-- AATCTCTAGC
B.FR.05.DEMB05FR001.JX140652     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.FR.08.DEMB08FR002.JX140654     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
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B.GA.88.OYI_397.M26727          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.GB.05.MM45d213_GN1.HM586212   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
B.GB.83.CAM1.D10112             TAGTGTGTGC CCATCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.GB.86.GB8_46R.AJ271445        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTAGAA-- AATCTCTAGC
B.GB.x.MANC.U23487              TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.HK.06.HK003.FJ460500          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.HK.06.HK004.FJ460501          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.IN.x.11807.EF694037           ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.IT.05.SG1.DQ672623            TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GTAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.00.DR2508.AB289588         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTCG --GTAGTCAG T-GTAGAA-- AATCTCTAGC
B.JP.01.DR388.AB289590          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TAAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.04.04JPDR6075B.AB221125    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.04.DR5913.AB480696         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACACTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.04.DR6089.AB286955         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TAAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.04.DR6174.AB480692         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TAAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.04.DR6175.AB480694         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACATTT --ATAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.05.DR6538.AB287363         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC CCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.05.DR6737.AB287364         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
B.JP.05.DR7060.AB287366         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACATTT --GTAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.05.DR7065.AB287368         TAGTGTGTGC CCATCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGATATCT --TTAGTCCG T-GTGGAAA- AATCTCTAGC
B.JP.98.DR1120.AB480698         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.99.DR1348.AB287370         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.x.DR1673.AB564745          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACC-TT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.x.DR1712.AB604946          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCA-T --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.x.DR1777.AB604948          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTATT --TTAGTCAG T-GTGAAA-- AATCTCTAGC
B.JP.x.JRC03B.AB565496          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.x.JRC05B.AB565497          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACTT --GTAGTCAG T-GTGAAA-- AATCTCTAGC
B.JP.x.JRC65B.AB565502          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.x.pJPDR0796B02.AB565478    TAGTGTGTGC CCGTCTGGTA TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCCT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.JP.x.pJRC57B09.AB641836       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.KR.02.02OSG1.JQ429433         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACTT --GTAGTCAG T-GTGGAA-- AATCTCTAGC
B.KR.03.03HJY8.JQ316131         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.KR.03.03KDE11.JQ316128        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.KR.03.03KGS5.JQ316132         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
B.KR.03.03LSH1.JQ316127         TAGTGTGTGC CCATCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAAACCCTT --TTAGTCAG T-GTGAAA-- AATCTCTAGC
B.KR.03.03YGS3.JQ316135         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.KR.04.04CWS5.JQ316133         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.KR.04.04KJS8.JQ316130         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.KR.04.04KMH5.DQ295193         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.KR.04.04KMK5.JQ316126         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.KR.04.04LHS6.AY839827         TAGTGTGTGC CCATCTGTTG TGTGACTCCG GTACTTAGAC TCTGGTAA-C TAGAGATCCC TCAGAC-CTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.KR.04.04LSK7.DQ295192         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.KR.04.04WK7_HIV_1_wk.DQ29519  TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.KR.05.05CSR3.DQ837381         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACAATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.KR.05.05YJN2.JQ316134         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.KR.07.07KYY4.JQ341411         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACTT --TTAGTTAG T-GTGGAA-- AATCTCTAGC
B.KR.99.99HYH2.JQ316129         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCACACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.MM.99.mSTD101.AB097870        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
B.NL.00.671_00T36.AY423387      TAGTGTGTGC CCGTCTGTTA TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.NL.86.3202A21_ACH3202A21.U34  TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GAGGAA-- AATCTCTAGC
B.NL.96.H434_42_A1.AY970948     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-ATGGAA-- AATCTCTAGC
B.PE.06.502_0491_wg5.JF320183   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCTGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.PE.06.502_0524_FL04.JF320008  ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.PE.06.502_0648_FL02.JF320215  ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.PE.06.502_0841_FL04.JF320208  ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.PE.06.502_2794_FL05.JF320244  ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCCTT --AAAGTCAG T-GTGGAA-- AATCTCTAGC
B.PE.07.502_0525_wg5.JF320191   --GTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --AAAGTCAA T-GTGGAA-- AATCTCTAGC
B.PE.07.502_1047_wg5.JF320226   ---------- ---------- ---------- ---------- --GGGTAA-C TAGAGATCCC TCAGACCT-- --TTAAAGTC A-GTGTGGAA AATCTCTAGC
B.PE.07.502_1399_wg4.JF320013   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.PE.07.502_2254_FL6.JF320018   ---------- ---------- ---------- ---------- --TGGTAG-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.PE.07.502_2349_wg2.JF320028   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCCTT --AAAGTCAG T-GTGGAA-- AATCTCTAGT
B.PE.07.502_2622_wg1.JF320189   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCCG A-GAGAA--- AATCTCTAGC
B.PE.07.502_2649_wg8.JF320019   ---------- ---------- ---------- ---------- ---------C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.RU.04.04RU128005.AY682547     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTRGAA-- AATCTCTAGC
B.RU.04.04RU129005.AY751406     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.RU.04.04RU139089.AY751407     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.RU.04.04RU139095.AY819715     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.RU.09.09RU4457.JX500709       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TAAGTCAG T-GTGGAA-- AATCTCTAGC
B.RU.10.10RU6629.JX500707       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.RU.11.11RU21n.JX500708        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.TH.06.AA010a_WG3.JX446800     ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCCTT --AAAGTCAG T-GTGGAA-- AATCTCTAGC
B.TH.07.AA040a_WG11.JX447156    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.TH.90.BK132.AY173951          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.TW.94.TWCYS_LM49.AF086817     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.00.14294_1.DQ853436        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGATCCTT --TTAGTTAG C-GTAGAA-- AATCTCTAGC
B.US.00.ES1_20.EF363123         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.00.RHPA_TF1.JN944917       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.00.THRO_TF1.JN944930       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCCTA --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.00.WITO_flB3.FJ496169      ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.01.REJO_TF1.JN944911       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.01.TRJO_flA1.FJ496151      ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGATCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.02.04013226_2_flH11.FJ496  ---------- -------TGG TAATA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.04.ES10_53.EF363127        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.04.ES4_24.EF363124         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.04.ES8_43.EF363126         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.04.SAMI_WGA1.EU547186      ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCTTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.04.USPI71101EI7y04051pcWG  ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTTAG T-GTGGAA-- AATCTCTAGC
B.US.04.USPI83747EI6y04121pcWG  --GTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TAGTCAG T-GTGGAAA- AATCTCTAGC
B.US.05.04013396_0_flE6.FJ4960  ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.05.502_0223_wg1.JF320059   ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------AG T-GTGGAA-- AATCTCTAGC
B.US.05.502_1400_FL02.JF320043  ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- -TGTGGAA-- AATCTCTAGC
B.US.05.502_2136_FL02.JF320185  TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.05.502_2495_wg02.JF320054  ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---TGGAAA- AATCTCTAGC
B.US.05.CR0307R.FJ469717        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----TCTAGC
B.US.05.USPI38417EI33y05051pcW  ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCATC --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.05.USPI88403EI14y05121pcW  ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGAC--TC --TAAGTCAG T-GTGGAAA- AATCTCTAGC
B.US.06.502_0341_FL05.JF320003  ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.06.502_1046_FL04.JF320564  --GTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATC --TTAGACAG T-GTAGAA-- AATCTCTAGC
B.US.06.502_1174_FL09.JF320053  ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.06.502_1211_FL01.JF320151  ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACC --TTAGTCAG T-GGGAAA-- AATCTCTAGC
B.US.06.502_1619_FL06.JF320126  ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T--TGGAA-- AATCTCTAGC
B.US.06.502_2667_FL03.JF320145  ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------GT --GTGGAA-- AATCTCTAGC
B.US.06.CH106_TF1.JN944897      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --TTAGTCCG G-GTGGAA-- AATCTCTAGC
B.US.06.CH40E_flA1.FJ495818     ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.06.CH58E_flA9.FJ495941     ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
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B.US.06.CH77E_flA1.FJ496000      ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCTTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.07.502_0364_wg2.JF320563    ---------- ---------- ---------- ---------- ----GTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.07.502_1115_wg1.JF320045    ---------- ---------- ---------- ---------- --TGGTAG-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.07.502_1478_wg4.JF320150    ---------C CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.07.502_1897_wg6.JF320182    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTAGTCAG G-GTGGAA-- AATCTCTAGC
B.US.07.502_2289_05.JF320197     ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.09.C1P.GU733713             TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.09.DEMB09US002.JX140657     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.US.10.CP1.JN397365             TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.10.VC1.JN397364             TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.11.ES38.JN397362            TAGTGTGTGC CCGTCTGTTG TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTCTC --GTAGTCAG T-GTGAAA-- AATCTCTAGC
B.US.83.5018_83.AY835777         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.83.5157_83.AY835781         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.83.RF_HAT3.M17451           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTCT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.83.SF2_LAV2_ARV2.K02007     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAAA- AATCTCTAGC
B.US.84.5019_84.AY835779         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.84.MNCG_MN.M17449           TAGTGTGTGC CCGTCTGTTA TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGATCCTT --TTAGGCAG T-GTGGAA-- AATCTCTAGC
B.US.84.NY5CG.M38431             TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.84.SF33.AY352275            TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.85.5077_85.AY835769         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.85.Ba_L.AB221005            TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.86.5084_86.AY835775         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.86.5096_86.AY835749         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.86.5127_86.AY835774         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.86.AD87_ADA.AF004394        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.86.JRFL_JR_FL.U63632        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.US.86.YU_2.M93258              TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.87.5113_87.AY835758         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.87.BC_BCSG3.L02317          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCATACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.88.5160_88.AY835763         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.88.WR27.AF286365            TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.89.P896_89_6.U39362         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCATC --CTAGTTAG T-GTAGAA-- AATCTCTAGC
B.US.90.US1.AY173952             ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.US.90.US2.AY173953             ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.US.90.US3.AY173954             ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.US.90.US4.AY173955             ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.US.90.WCIPR.U69591             TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.90.WEAU160_GHOSH.U21135     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.91.5048_91.AY835761         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.91.DH12_3.AF069140          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.91.SUMA_flC5.FJ496145       ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCAA- --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.93.WCD32P0793.DQ487188      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCATA ----AGTCAG T-GTGGAA-- AATCTCTAGC
B.US.93.WCM32P0793.DQ487190      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCCTT ----AGTCAG T-GTGGAA-- AATCTCTAGC
B.US.94.5082_94.AY835773         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.95.5073_95.AY835768         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.95.USPI90770EI72y95091pcW   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCTTT --ATAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.96.5155_96.AY835753         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.96.USPI55751EI32y96071pcW   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCCTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.97.ARES2.AB078005           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.98.15384_1.DQ853463         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCCTT --TTAGTTAG C-GTAGAA-- AATCTCTAGC
B.US.98.98USHVTN1925c1.AY56010   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.98.98USHVTN3605c9.AY56010   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
B.US.98.98USHVTN8229c6.AY56010   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACC-TT --TTAGTCAG T-ATAGAAA- AATCTCTAGC
B.US.98.98USHVTN941c1.AY560110   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-ATGGAA-- AATCTCTAGC
B.US.98.WC3_0498_4.EF175212      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTCTT --TTAGTCAG T-GTGAAA-- AATCTCTAGC
B.US.x.CR0059T.FJ469694          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----TCTAGC
B.US.x.sample_C_BID_D617.JX503   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---GGGGTGT
B.VE.10.DEMB10VE001.JX140659     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.ZA.03.03ZAPS045MB2.DQ396398    ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---AAAGTCA G-TAGAAA-- AATCTCTAGC
C.BR.02.02BR2022.JN692434        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACCT --ATTGTCAG T-GTGGAA-- AATCTCTAGC
C.BR.04.04BR013.AY727522         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGATACCT --TTTGTCAG T-GTGGAA-- AATCTCTAGC
C.BR.04.04BR021.AY727523         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACGCCT --TTCGTCAG T-GTGGAA-- AATCTCTAGC
C.BR.04.04BR038.AY727524         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGATACTT --TTTGTCCG T-GTGGAA-- AATCTCTAGC
C.BR.04.04BR073.AY727525         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACAT --TTTGTCCG T-GTGGAA-- AATCTCTAGC
C.BR.07.DEMC07BR003.JX140663     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.BR.92.BR025_d.U52953           ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.BR.98.98BR004.AF286228         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.BW.00.00BW07621.AF443088       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW076820.AF443089      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW087421.AF443090      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW147127.AF443091      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW16162.AF443092       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW1686.AF443093        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW17593.AF443094       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW17732.AF443095       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- TCTCTAGCG-
C.BW.00.00BW17835.AF443096       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW17956.AF443097       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW18113.AF443098       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW18595.AF443099       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW18802.AF443100       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW192113.AF443101      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW20361.AF443102       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW20636.AF443103       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW20872.AF443104       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW2127214.AF443105     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW22767.AF443107       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW38193.AF443108       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW38428.AF443109       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW38713.AF443110       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW3876_9.AF443111      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW3886_8.AF443112      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW3891_6.AF443113      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW3970_2.AF443114      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.00.00BW5031_1.AF443115      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- TCTCTAGCAG
C.BW.96.96BW01B03.AF110959       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAAAGATCCC TCAAACCCTT ---TTGGTTG T-GTGGAA-- AATCTCTAGC
C.BW.96.96BW0402.AF110962        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --CGCGGTAG T-GTGGAA-- AATCTCTAGC
C.BW.96.96BW0502.AF110967        TAGTGTGTGC CCGTCTGATG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGAACCTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.BW.96.96BW06.AF290028          TAGTGTGTGC CCATCTGTTA TATGG--A-- ---------G TCTGGTAA-C TAGAGATCCC TCAGGCCCTT ---TTGGTAG T-GAGGAA-- AATCTCTAGC
C.BW.96.96BW1104.AF110969        ---------- ---------- ---------- ---------- ---------- ---------- ---------- -----GGTAG T-GTGGAA-- AATCTCTAGC
C.BW.96.96BW1210.AF110972        ---------- ---------- ---------- ---------- ---------- ---------- ---------- -----GGTAG T-GTGGAA-- AATTTCTAGC
C.BW.96.96BW15C02.AF110974       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAAAGATCCC TCAGACCCTT --TTTGGTAT T-GTGGAA-- AATCTCTAGC
C.BW.96.96BW16B01.AF110976       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTT- --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.BW.96.96BW17.AF110980          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.96.96BWM032.AF443075        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.96.96BWMO1_5.AF443074       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
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C.BW.98.98BWMC122.AF443076      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.98.98BWMC134.AF443077      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.98.98BWMC14A3.AF443078     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.98.98BWMO1410.AF443079     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.98.98BWMO18D5.AF443080     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.98.98BWMO36A5.AF443081     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.98.98BWMO37D5.AF443082     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.99.99BW393212.AF443083     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.99.99BW46424.AF443084      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.99.99BW4745.AF443085       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.BW.99.99BW47547.AF443086      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTTTAGC
C.BW.99.99BWMC168.AF443087      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.DJ.x.DJ259.AB485643           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTTCT --TTTGTCCG T-GTGGAA-- AAATCTCTAGC
C.DK.01.CTL_015.EF514713        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------T
C.ES.06.X1936.EU786673          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT ---GTGGTAG T-GTGGAA-- AATCTCTAGC
C.ES.07.X2118_2.EU884500        -----TGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT ---TTGGTTG T-GTGGAA-- AATCTCTAGC
C.ES.08.X2363_2.EU786681        ---------- ----CTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGATCAAT --TTTGCTCT T-GTGGAA-- AATCTCTAGC
C.ET.86.ETH2220.U46016          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
C.IL.98.98IS002.AF286233        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.IN.03.D24.EF469243            CAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCCTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.IN.93.93IN101.AB023804        CAGTGTGTGC CCGTCTATTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.IN.93.93IN904.AF067157        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
C.IN.93.93IN999.AF067154        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
C.IN.94.94IN11246.AF067159      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
C.IN.94.94IN476.AF286223        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.IN.95.95IN21068.AF067155      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
C.IN.98.98IN012.AF286231        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.IN.98.98IN022.AF286232        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.IN.99.01IN565_10.AY049708     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- -----AAA-- AATCTCTAGC
C.IN.x.VB39.EF694033            ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------C
C.IN.x.VB49.EF694036            ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
C.MM.99.mIDU101_3.AB097871      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
C.TZ.98.98TZ013.AF286234        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.TZ.98.98TZ017.AF286235        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.ZA.00.1069MB.AY838567         ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TGTGGTAG T-GTGGAA-- AATCTCCAGC
C.ZA.00.1119MB.AY463229         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCATT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.00.1134MB.AY463217         TAGTGTGTGC CCGTCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --TTGGT-AG T-GTGGAA-- AATCTCTAGC
C.ZA.00.1157M3M.AY585266        ---------- ---------- ---------- ---------- ---------- ---------- ---------- --CTGGTAAC T-AGAGAA-- AATCTCTAGC
C.ZA.00.1162MB.AY463224         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.00.1165MB.AY463230         TAGTGTGTGC CCGTCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --ATTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.00.1168MB.AY463231         TAGTGTGTGC CCGTCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.00.1170MB.AY463225         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TTGGCAG T-ATGGAA-- AATCTCTAGC
C.ZA.00.1171MB.AY463232         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.00.1176MB.AY463218         ---------- ---------- ---------- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTAT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.00.1178MB.AY463233         TAGTGTGTGC CCGTCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GTTGGTAG T-GAA----- AATCTCTAGC
C.ZA.00.1184MB.AY838566         ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.00.1189MB.AY838565         ---------- ---TCTGTTG TGTGG----- --------AC TCTGGTAA-C TAGAGATCCC TCAGACCCTT -TATTGGTAG T-GTGGAAA- AATCTCTAGC
C.ZA.00.1192M3M.AY463219        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.00.1195MB.AY463220         TAGTGTGTGC CCGTCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GTTGGTAG T-GCGGAA-- AATCTCTAGC
C.ZA.00.1197MB.AY463234         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.00.1210MB.AY463221         ---------- ---------- ---------- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --TGTGGTAG A-GTGGAA-- AATCTCTAGC
C.ZA.00.1214MB.AY463236         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --CGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.00.1217MB.AY463226         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.00.1225MB.AY463227         TAGTGTGTGC CCGTCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GTTGGTAT T-GTGGCA-- AATCTCTAGC
C.ZA.00.1228MB.AY463222         TAGTGTGTGC CCGTCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TTGGTAG T-GTGGAA-- AATCTTTTAGC
C.ZA.00.J112MA.AY838568         ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAC --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.00.J38MA.AY463228          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TTGGAAG T-GTGGAA-- AATCTCTAGC
C.ZA.01.01ZATM45.AY228557       TAGTGTGTGC CCATCTGTTG TGTGACTCTG GTCTGTGA-C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.01.2134MB.AY463237         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.01.J54Ma.AY463223          TAGTGTGTGC CCGTCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTCTT --TGTGGTAG T-GCGGAA-- AATCTCTAGC
C.ZA.02.02ZAPS001MB1.DQ275648   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.02.02ZAPS005MB1.DQ351235   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAG-C TAGAGATCCC TCAGATCCTT --AGTGGTAG T-GTGGAAA- AATCTCTAGC
C.ZA.02.02ZAPS006MB1.DQ351220   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --CGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.02.02ZAPS008MB1.DQ275647   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.02.02ZAPS013MB1.DQ351222   ---------- ---TCTGTTG TGTGA----- ---------C TCTGGGAA-C TAGAGATCCC TCAGACCTTT ---GCGGCTG T-GTGGAA-- AATCTCTAGC
C.ZA.02.02ZAPS014MB1.DQ351218   ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.02.02ZAPS015MB1.DQ369995   ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.03.03ZAPS017MB1.DQ351224   ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAG --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.03.03ZAPS020MB1.DQ275653   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGACACTT --GTTGGTTG T-GTGGAA-- AGTCTCTAGC
C.ZA.03.03ZAPS021MB1.DQ369978   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTTG T-GTGGAA-- AATCTCTAGC
C.ZA.03.03ZAPS023MB1.DQ351225   ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAC --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.03.03ZAPS024MB1.DQ396367   ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCCTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.03.03ZAPS025MB1.DQ351226   ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.03.03ZAPS026MB1.DQ369985   ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.03.03ZAPS027MB1.DQ351223   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --AGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.03.03ZAPS030MB1.DQ369996   ---------- ---TCTGTTG TGTGA----- ---------C TCCGGTAA-C TAGAGATCCC TCAGACCCTT --GTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.03.03ZAPS032MB1.DQ445633   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
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C.ZA.03.03ZAPS044MB1.DQ396384   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT ---TTGGTTG T-GTGGAA-- AATCTCTAGC
C.ZA.03.03ZAPS046MB1.DQ369984   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
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C.ZA.03.03ZAPS049MB1.DQ369986   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAG --TTTGGTAG C-GTGGAA-- AATCTCTAGC
C.ZA.03.03ZAPS050MB1.DQ369980   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TGGGTTG T-GTGGAA-- AATCTCTAGC
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C.ZA.03.03ZAPS052MB1.DQ369987   ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --ATTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.03.03ZAPS054MB2.DQ369988   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCAT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
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C.ZA.03.03ZAPS057MB2.DQ369989   ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
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C.ZA.03.03ZAPS063MB1.DQ396388   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAC --TTTGGTAG T-GTGGAA-- AATCTCTAGC
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C.ZA.03.03ZAPS079B1.DQ093592    ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.03.03ZAPS081MB1.DQ351219   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTGG T-GTGGAA-- AATCTCTAGC
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C.ZA.03.03ZAPS086MB1.DQ275654   ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --GTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.03.03ZAPS088MB1.DQ275651   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTGGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.03.03ZAPS089MB1.DQ351216   ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
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C.ZA.03.03ZAPS136MB1.DQ351231    ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
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C.ZA.03.SK116B1.AY772696         ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGCAG T-GTGGAA-- AATCTCTAGA
C.ZA.04.04ZAPS157MB1.DQ351232    ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS160B1.DQ164107     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGATCCTT --TTTGGTAG T-GTGAAA-- AATCTCTAGC
C.ZA.04.04ZAPS161B1.DQ164115     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --TGTGTTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS165MB1.DQ351221    ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTAT --ATTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS168B1.DQ164121     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCTTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS169MB1.DQ396393    ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGAACCTT ---TTGGCAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS172MB1.DQ275659    ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCCTT --GTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS177MB1.DQ396394    ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --CTTGGTGG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS188B1.DQ164122     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --TGAGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS189B1.DQ164117     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAGT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS190B1.DQ093602     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --TGAGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS195B1.DQ164118     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS197MB1.DQ369997    ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --CGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS198MB1.DQ445637    ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGATCAAT --TTTGGTCG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS199B1.DQ164110     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TGTGGCAG T-GTGGAA-- AATTTCTAGC
C.ZA.04.04ZAPS202B1.DQ093598     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS205B1.DQ093599     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS206B1.DQ164126     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS208B1.DQ093603     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS214B1.DQ093588     ---------- ---TCTGTTG TGTGA----- ---------C TCCGGTAA-C TAGAGATCCC TCAGACCCTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS216B1.DQ093600     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZAPS217B1.DQ164119     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
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C.ZA.04.04ZASK031B2.AY901974     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK083B2.DQ093585     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK127B1.DQ011172     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK128B1.DQ093586     ---------- ---TCTGTTG TTTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCAAT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK131B1.DQ056412     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK132B1.DQ093594     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK135B1.DQ011166     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
C.ZA.04.04ZASK136B1.DQ011177     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK139B1.AY878072     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK142B1.DQ011167     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK145B1.AY901976     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GTTGATAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK146.AY772699       ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTC --AGTGTTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK147B1.DQ164114     ---------- ---------- ----A----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCAT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK148B1.DQ093590     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK150B1.DQ011178     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTCCT --TTTGGTAG T-GTGAAA-- AATCTCTAGC
C.ZA.04.04ZASK151B1.AY901977     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCCT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK154B1.AY878071     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK155B1.AY901978     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAACT AGAGATCCCT CAGACCCCAA --GTGGGTAA GTGTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK156B1.DQ011171     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK159B1.DQ011179     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TTGGCCG T-GTGGAA-- AATCTCCAGC
C.ZA.04.04ZASK160B1.DQ011173     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK161B1.DQ011170     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK163B1.AY901979     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK164B1.DQ056405     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK165B1.DQ396387     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK167B1.DQ164127     ---------- ---TCTGGTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCAAC --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK168B1.AY878058     --TCTGTTGT GTGACTCTGG TAATA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GTTGGTTG C-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK169B1.DQ396381     ---------- ---TCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK170B1.DQ093595     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK171B1.DQ351217     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCAAC --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK172B1.DQ369998     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TGGAGATCCC TCAGACCCTT --GTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK173B1.DQ093604     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK174B1.AY901980     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK175B1.DQ164129     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTT- --TTTGGTTG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK176B1.DQ056416     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAC --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK178B1.DQ093587     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK180B1.AY878059     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGATCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK181B1.AY878062     ---------- ---TCTGATG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT ---GTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK182B1.AY878054     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK183B1.AY878055     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK184B1.DQ056418     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAC --TTCGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK185B1.DQ011174     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK190B1.DQ056409     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAGT --TTTGGCAG T-GTGGA--- AATCTCTAGC
C.ZA.04.04ZASK191B1.DQ369993     ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTATC --TTTAGTTG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK192B1.DQ396382     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK193B1.DQ396396     ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGGGATCCC TCAGACCTTT ----TGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK196B1.DQ056413     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK200B1.DQ396383     ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTTG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK201B1.DQ396397     ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAC --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK202B1.DQ011180     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TGGAGATCCC TCAGACCCTT --TTTGGTTG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK204B1.DQ056414     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK206B1.DQ056415     --TCTGTTGT GTGACTCTGG TAATA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCT- --AGTGGTAT T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK208B1.DQ056406     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK217B1.DQ056417     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TTGGCAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.04ZASK234B1.DQ093605     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.CAP30_5w_F4.GQ999973     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.CAP61_8w_F3.GQ999975     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --GTGGTAGT G-TGGAA--- AATCTCTAGC
C.ZA.04.SK133B1.AY772698         ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTACT --TGTGGTCG T-GTGGAA-- AATCTCTAGC
C.ZA.04.SK134B1.AY703909         ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.SK140B1.AY901973         ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.SK143B1.AY703910         ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT ---GTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.04.SK144B1.AY703911         ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TTGGTAG T-GCGGAA-- AATCTCTAGC
C.ZA.05.05ZAPSK240B1.DQ369991    ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.05.05ZASK243B1.DQ396372     ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.05.05ZASK244B1.DQ369992     ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.05.05ZASK245B1.DQ369982     ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.05.05ZASK246B1.DQ369983     ---------- ---TCTGTTG TGTGG----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.05.05ZASK247B1.DQ369994     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGATCATT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.05.CAP174_4w.GQ999981       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAC --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.05.CAP206_8w_F1.GQ999982    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTGT G-TGGAA--- AATCTCTAGC
C.ZA.05.CAP210_5w.GQ999983       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGATCCTT --TTGTCAGA G-TGGAA--- AATCTCTAGC
C.ZA.05.CAP228_8w_F2.GQ999984    ---------- ---------- ---------- ---------- ---------- ---------- ---------- T-TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.05.CAP229_7w.GQ999985       -AGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.05.CAP239_5w_F1.GQ999991    TAGTGTGTGC CCCTTTGATG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.05.CAP244_8w_F1.GQ999986    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTGGTAGT G-TGGAA--- AATCTCTAGC
C.ZA.05.CAP248_9w.GQ999987       -AGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GTTGGTAG T-GTGGAA-- AATCTCTAGC
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C.ZA.05.CAP257_7w_F1.GQ999990    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAGT --TTGGTAGT G-CGGAA--- AATCTCTAGC
C.ZA.05.CAP45_5w_F1.GQ999974     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.05.CAP63_5w_F4.GQ999976     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGATCCTT T--TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.05.CAP65_6w.GQ999977        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAC --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.05.CAP84_3w_F2.GQ999978     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTCT --TTGGTAGT G-TGGAA--- AATCTCTAGC
C.ZA.05.CAP85_5w_F1.GQ999979     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --TTGGTAGT G-TGGAA--- AATCTCTAGC
C.ZA.05.CAP88_5w_F2.GQ999980     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC G--------- ---------- --GAA----- AATCTCTAGC
C.ZA.05.CAP8_3w_F2.GQ999972      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.07.DEMC07ZA011.JX140664     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.ZA.08.DEMC08ZA011.JX140666     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.ZA.09.DEMC09ZA008.JX140667     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.ZA.09.DEMC09ZA009.JX140668     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.ZA.10.DEMC10ZA001.JX140669     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.ZA.90.pZAC_R3714.JN188292      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTA --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.97.97ZA003.AY118165         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.ZA.97.97ZA009.AY118166         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.ZA.97.97ZA012.AF286227         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.ZA.98.98ZA445.AY158533         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.ZA.98.98ZA502.AY158534         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.ZA.98.98ZA528.AY158535         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.ZA.98.TV001.AY162223           TAGTGTGTGC CCATCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.98.TV002.AY162224           TAGTGTGTGC CCATCTGTTG TGTGACTCTG GTAA-----C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.98.TV012.AY162225           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --TGTGGTGT --GTGGAA-- AATCTCTAGC
C.ZA.99.99ZACM9.AF411967         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.99.99ZALT21.EU293446        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.99.99ZALT39.EU293447        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.99.99ZALT42.EU293448        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
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C.ZA.99.99ZALT45.EU293449        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --CGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.99.99ZALT46.EU293450        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT ---TTTGGTG T-GTGGAA-- AATCTCTAGC
C.ZA.99.99ZALT4.EU293444         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAC --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.99.99ZALT5.EU293445         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTTGGCAG T-GTGGAA-- AATCTCTAGC
C.ZA.99.99ZATM10.AY228556        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZA.99.ZASW7.AF411966           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZM.02.02ZM108.AB254141         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --TTTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZM.02.02ZM110.AB254142         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---GTGTTAG T-GTGGAA-- AATCTCTAGC
C.ZM.02.02ZM114.AB254146         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZM.02.02ZM115.AB254148         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GT-GGCAT T-GTGGAAA- AATCTCTAGC
C.ZM.02.02ZMBC.AB254149          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---GTGGTAG T-GCGGAA-- AATCTCTAGC
C.ZM.02.02ZMDB.AB254153          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
C.ZM.02.02ZMJC.AB254155          TAGTGTGTGC CCATCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAAACCAAA --AGGTGTGC --GTGGAA-- AATCTCTAGC
C.ZM.03.ZM246F_flA10.FJ496186    ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCATT --TTGG-CAG T-GTGGAA-- AATCTCTAGC
C.ZM.03.ZM247F_flA1.FJ496195     ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCATT --TTGG-TAG T-GTGGAA-- AATCTCTAGC
C.ZM.03.ZM249M_flC1.FJ496209     ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCTTT ---TAGGTAG T-GTGGAA-- AATCTCTAGC
C.ZM.89.ZAM18.AB485645           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TGTGGTAG T-GTGAAA-- AATCTCTAGC
C.ZM.96.96ZM651.AF286224         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.ZM.96.96ZM751.AF286225         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
C.ZM.x.HIV1084i.AY805330         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCGT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
D.CD.83.ELI.K03454               TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCCT --TTAGTCAG A-GTGGAA-- AATCTCTAGC
D.CD.83.NDK.M27323               TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGATCCGT --TTAGTCCG A-GTGAAA-- AATCTCTAGC
D.CD.84.84ZR085.U88822           TAGTGTGTGC CCGTCTGTTT TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACAGCT --TTAGTCCG G-GTGGAA-- AATCTCTAGC
D.CD.85.Z2Z6_Z2_CDC_Z34.M22639   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCCT --TTAGTCAG A-GTGGAA-- AATCTCTAGC
D.CM.10.DEMD10CM009.JX140670     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
D.KE.93.MB2059.AF133821          CAGTGTGTGC CCATCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTCT --TTAGTCAG T-GTAAAA-- AATCTCTAGC
D.KE.97.ML415_2.AY322189         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTCT --GTAGTCAG T-GTGAA--- AATCTCTAGC
D.KR.04.04KBH8.DQ054367          TAGTGTATGC CCGTCTGTTG TGTAA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTCG --TTTGTCAG T-GTAAAA-- AATCTCTCGC
D.SN.90.SE365.AB485648           TAGTGTGTGC CCATCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
D.UG.05.p190049.JX236668         TAGTGTGTGC CCGTCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGAA--- AATCTCTAGC
D.UG.07.p191647.JX236670         TAGTGTGTGC CCATCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTCT --TTGGTCAG T-GTAGAA-- AATCTCTAGC
D.UG.07.p191882.JX236673         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTCT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
D.UG.07.pSC191727.JX236679       TAGTGTGTGC CCATCTGTTG TGTGA----- ---------C TCTGGTAAAC TAGAGATCCC TCAGACCTCT --TTAGTCAG T-GTAGTCAG AATCTCTAGC
D.UG.08.p191859.JX236672         TAGTGTGTGC CCATCAATTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCTCT --TTAGTCCG G-GTGGAA-- AATCTCTAGC
D.UG.91.UG270.AB485650           TAGTGTGTGC CCATCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTCT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
D.UG.92.92UG001.AJ320484         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTCT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
D.UG.94.94UG114.U88824           ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
D.ZA.84.R2.AY773338              ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
D.ZA.85.R286.AY773340            ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
D.ZA.86.R482.AY773341            ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
D.ZA.90.R1.EF633445              ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
F1.AR.02.ARE933.DQ189088         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
F1.BE.93.VI850.AF077336          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
F1.BR.02.02BR082.FJ771006        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
F1.BR.02.02BR170.FJ771007        ---------- --GTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
F1.BR.06.06BR564.FJ771008        TAATGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATC --TTAGTCAG T-GTGGAA-- AATCTCTAGC
F1.BR.06.06BR579.FJ771009        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
F1.BR.07.07BR844.FJ771010        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
F1.BR.89.BZ126.AY173957          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------C
F1.BR.93.93BR020_1.AF005494      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
F1.DE.x.MVP_30846.EU446022       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
F1.ES.02.ES_X845_4.FJ670516      TAAGGGGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG A-GTGTAA-- AATCTCTAGC
F1.ES.11.DEMF110ES001.JX140671   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
F1.ES.x.P1146.DQ979023           ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
F1.ES.x.X1093_2.DQ979025         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
F1.ES.x.X1670.DQ979024           -------TGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
F1.FI.93.FIN9363.AF075703        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
F1.RO.96.BCI_R07.AB485658        TAGTGTGTGC CCGTCTGTTG TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
F1.RU.08.D88_845.GQ290462        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
F2.CM.10.DEMF210CM007.JX140673   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
G.BE.96.DRCBL.AF084936           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGATAG T-GTAAA--- AATCTCTAGC
G.CM.01.A1786.FJ389367           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
G.CM.04.178_15.FJ389363          TAGTGTGTGC CCGTCTGTTG TGCGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGATAG T-GTAAA--- AATCTCTAGC
G.CM.04.314_40.FJ389364          TAGTGTGTGC CCGTCTGTTT TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTARACAG T-GTAAA--- AATCTCTAGC
G.CM.04.515_28.FJ389365          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
G.CM.04.944_5.FJ389366           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CRAGACAC T-GTAAA--- AATCTCTAGC
G.CM.10.DEMG10CM008.JX140676     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
G.CM.96.96CMABB55.AY772535       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
G.CN.06.sh52.HM067749            TAGTGTGGGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGGACCAC T-CTAGACAG T-GTAAA--- AATCTCTAGC
G.CU.99.Cu74.AY586547            TGGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGG T-GTAAA--- AATCTCTAGC
G.CU.99.Cu85.AY586548            TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGAYAG T-GTAAA--- AATCTCTAGC
G.CU.99.Cu87.AY586549            TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
G.ES.05.P962.EU786670            -------TGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG A-ATAAA--- AATCTCTAGC
G.ES.08.P1981_2.FJ670530         TAAGGGGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGTA- --GTGTAAA- AATCTCTAGC
G.ES.09.X2634_2.GU362882         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGTGA- --GTGTAAA- AATCTCTAGC
G.ES.99.X138.AF450098            ---------- -------TTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGATAG T-GTAAA--- AATCTCTAGC
G.GH.03.03GH175G.AB287004        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
G.KE.93.HH8793_12_1.AF061641     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --GGCTTA-- AATCTCTAGC
G.NG.92.92NG083_JV10832.U88826   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
G.PT.x.PT2695.AY612637           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGATAG T-GTAAA--- AATCTCTAGA
G.PT.x.PT3037.FR846408           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
G.PT.x.PT3306.FR846409           TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAAACAG T-GTAAA--- AATCTCTAGC
G.PT.x.PT988.FR846410            TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACC --CTAGATAG T-GTAAA--- AATCTCTAGC
G.SE.93.SE6165_G6165.AF061642    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --GGCTTA-- AATCTCTAGC
H.BE.93.VI991.AF190127           ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
H.BE.93.VI997.AF190128           ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
H.CF.90.056.AF005496             ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
H.GB.00.00GBAC4001.FJ711703      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAW-C TAGAGATCCC TCAGACCATT --YTAGTCAG T-GTGGAA-- AATCTCTAGC
J.CM.04.04CMU11421.GU237072      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACC-TT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
J.SE.93.SE9280_7887.AF082394     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
J.SE.94.SE9173_7022.AF082395     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.CF.90.90CF11697.AF197340   TGGTGTGTGC CCGTCTGTGA TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.CF.90.90CF4071.AF197341    TGGTGTGTGC CCGTCTGTGA TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CAAGACTG A-GTAAA--- AATCTCTAGC
01_AE.CF.90.90CR402_CAR_E_4002   TGGTGTGTGC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.CN.05.FJ051.DQ859178       CGGTGTGTGC CCATCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.CN.05.FJ053.DQ859179       TGGTGTGTGC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGATTG A-GTAAA--- AATCTCTAGC
01_AE.CN.05.Fj052.EF036528       TGGTGTGTGC CCGTCTGTGT TTGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTA A-GTAAA--- AATCTCTAGC
01_AE.CN.05.Fj055.EF036527       TGGTGTGTGC CCATCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.CN.05.Fj056.EF036529       TGGTGTGTGC CCATCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CCAGACTG A-GTAAA--- AATCTCTAGC
01_AE.CN.05.Fj057.EF036530       TGGTGTGTGC CCATCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.CN.05.Fj065.EF036534       TGGTGTGTGC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.CN.05.Fj066.EF036535       TGGTGTGTGC CCATCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.CN.06.FJ054.DQ859180       TGGTGTGTGC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACTG --CA-GMCTG A-GTAAA--- AATCTCTAGC
01_AE.CN.06.Fj062.EF036531       TGGTGTGTGC CCATCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.CN.06.Fj063.EF036532       TGGTGTGTGC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTA A-GTAAA--- AATCTCTAGC
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01_AE.CN.06.Fj064.EF036533       TAGTGTGTGC CCGTCTGTGT TAGGA----- ---------C TCTAGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTA A-GTAAA--- AATCTCTAGC
01_AE.HK.04.HK001.DQ234790       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.JP.x.DR0492.AB253423       TGGTGTGTGC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.JP.x.DR2594.AB253660       TGGTGTGTGC CCATTTGTGT CAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.JP.x.DR6824.AB253426       TGGTGTGTGC CCATCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.JP.x.JRC77AE.AB565504      TGGTGTGTGC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- ATTCTCTAGC
01_AE.TH.04.AA027a_wg4.JX44702   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.TH.04.AA075a_WG7.JX44754   -------TGC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAA---- AATCTCTAGC
01_AE.TH.04.BKM.DQ314732         TGGTGTGTGC CCGTCTGTGT TTGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.05.AA004a_wg4a.JX4467   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GAAA---- AATCTCTAGC
01_AE.TH.05.AA033a_wg6a.JX4470   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.05.AA049a_WG13.JX4472   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.TH.05.AA064a_WG2.JX44741   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.TH.05.AA079a_WG4.JX44759   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.TH.05.AA101a_WG1.JX44793   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.TH.05.AA107a_wg4.JX44802   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.TH.06.AA002a_WG1.JX44666   --------GC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.06.AA017a_wg1.JX44689   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.TH.06.AA034a_wg2.JX44708   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.06.AA038a_WG3.JX44713   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.TH.06.AA055a_WG4.JX44731   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.TH.06.AA056a_WG5.JX44731   ---------- ---------- ---------- ---------- ---------- ---------- ---------- -----GACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.06.AA059a_WG5.JX44735   -------TGC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.06.AA063a_WG37.JX4474   ---------- ---------- ---------- ---------- ---------- ---------- ----ACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.06.AA068a_14.JX447465   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.TH.06.AA082a_WG9.JX44764   -------TGC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.06.AA085a_wg2.JX44768   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.06.AA088a_wg14.JX4477   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.06.AA099a_WG9.JX44789   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGG A-GTAAA--- AATCTCTAGC
01_AE.TH.07.AA015a_WG4.JX44687   ---------- ---------- ---------- ---------- ---------- ---------- TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.07.AA019a_WG11.JX4469   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.07.AA028a_wg3.JX44702   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.07.AA050a_WG7.JX44728   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.TH.08.AA007a_WG10.JX4467   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.TH.08.AA037a_WG6.JX44712   ---------C CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAA---- AATCTCTAGC
01_AE.TH.08.AA060a_WG1.JX44735   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.TH.08.AA067a_WG12.JX4474   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.08.AA108a_WG6.JX44802   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.09.AA090a_WG11.JX4477   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.09.AA111a_WG11.JX4480   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.90.CM240.U54771         TGGTGTGTGC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTACC
01_AE.TH.93.93TH253.U51189       TGGTGTGTGC CCATCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGATCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.93.93TH9021.AF164485    TGGTGTGTGC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAAAAATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.95.95TNIH022.AB032740   TGGTGTGTGC CCGTCTGTGT TAAGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.95.95TNIH047.AB032741   TGGTGTGTGC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.TH.x.NP03.AB485654         TGGTGTGTGC CCGTCTGTGT TTGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01_AE.VN.97.97VNAG204.FJ185247   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
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01_AE.VN.97.97VNAG207.FJ185249   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNAG210.FJ185251   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
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01_AE.VN.97.97VNAG214.FJ185253   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNAG218.FJ185255   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNHCM314.FJ18524   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNHCM319.FJ18524   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNHCM343.FJ18524   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNHCM345.FJ18524   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.98.98VNND15.FJ185235    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
02_AG.CM.08.DE00208CM001.JX140   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
02_AG.CM.08.DE00208CM004.JX140   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
02_AG.CM.99.pBD6_15.AY271690     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --TTAGACTG T-GTAAA--- AATCTCTAGC
02_AG.ES.06.P1261.EU786671       -------TGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCACT --CTAGACTT A-GTGAAA-- AATCTCTAGC
02_AG.ES.06.P1423.EU884501       ------GTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CGAGACCG T-GTAGAA-- AATCTCTAGC
02_AG.FR.91.DJ263.AF063223       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
02_AG.GH.03.03GH181AG.AB286855   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCACT --GTAGACTG T-GTAAAA-- AATCTCTAGC
02_AG.GH.03.03GH182AG.AB286857   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --TTACTCAG T-GTAAA--- AATCTCTAGC
02_AG.GH.03.03GH189AG.AB286862   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --TTCAGTTT GTGTAAA--- AATCTCTAGC
02_AG.GH.03.03GH197AG.AB286863   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTACT --CTAGACGG T-GTAA---- AATCTCTAGC
02_AG.GH.03.GHNJ185.AB231895     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG T-GTAGAA-- AATCTCTAGC
02_AG.GH.03.GHNJ188.AB231896     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTACT --TTAGTCAG T-GTAAA--- AATCTCTAGC
02_AG.GH.03.GHNJ196.AB231898     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTAGTCTT G-GTAAA--- AATCTCTAGC
02_AG.GH.97.97GH_AG1.AB049811    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTACT --CTAGACTG T-GTAAA--- AATCTCTAGC
02_AG.GH.x.I_2496.AB485633       TAGTGTGTGC CCGTCTATTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCACT --CTAGACTG T-GTAAGAA- AATCTCTAGC
02_AG.GW.04.CC_0030.FJ694791     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
02_AG.GW.05.CC_0048.FJ694792     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
02_AG.KR.07.07MHI10.JQ316136     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGCTTG A-GTAGA--- AATCTCTAGC
02_AG.LR.x.POC44951.AB485636     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG T-ATAAA--- AATCTCTAGC
02_AG.NG.x.IBNG.L39106           TAGTGTGTGC CCGTCTGATT GTGTGA---- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG T-GTAAA--- AATCTCTAGC
02_AG.SE.94.SE7812.AF107770      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
03_AB.BY.00.98BY10443.AF414006   TAGTGTGTGC CCGTCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGG T-GTAAAA-- AATCTCTAGC
03_AB.RU.98.RU98001_98RU001.AF   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- -------A-- AATCTCTAGC
04_cpx.CY.94.94CY032_3.AF04933   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
04_cpx.GR.00.DE00400GR002.JX14   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
04_cpx.GR.91.GR11_97PVCH.AF119   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG T-ATAAAA-- AATCTCTAGT
04_cpx.GR.97.GR84_97PVMY.AF119   TAGTGTGTGC CCGTCTGGTG TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGG T-ATAAA--- AATCTCTAGC
05_DF.BE.93.VI961.AF076998       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
05_DF.BE.x.VI1310.AF193253       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- -ATCTCTAGC
05_DF.ES.99.X492.AY227107        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
06_cpx.AU.96.BFP90.AF064699      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
06_cpx.CD.x.BCF_Dioum.AB485660   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GAAA---- AATCTCTAGC
06_cpx.EE.01.EE0359.AY535659     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGAACACT --TTAGACAG T-GTAAA--- AATCTCTAGC
06_cpx.GH.03.03GH173_06.AB2868   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGAAGG T-GTAAA--- AATCTCTAGC
06_cpx.ML.95.95ML127.AJ288982    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACG --CTAGGCGG T-GTAA---- AATCTCTAGC
06_cpx.ML.95.95ML84.AJ245481     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTTGACAG T-GTAAA--- AATCTCTAGC
06_cpx.SN.97.97SE1078.AJ288981   TAGTGTGTGC CCGTCTGGTG TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGAAGG T-GTAAA--- AATCTCTAGC
07_BC.CN.05.XJDC6431_2.EF36837   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
07_BC.CN.05.XJDC6441.EF368370    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
07_BC.CN.06.Sichuan_2006_SC006   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
07_BC.CN.06.Sichuan_2006_SC008   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
07_BC.CN.06.Xinjiang_2006_709.   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
07_BC.CN.07.CNGZD.JQ423923       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
07_BC.CN.98.98CN009.AF286230     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
08_BC.CN.00.p00CH_HH090_08_BC3   CAGTGTGTGC CCGTCTATTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACC-CT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
08_BC.CN.00.p00CH_WS035_08_BC5   CAGTGTGTGC CCGTCTATTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACC-CT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
08_BC.CN.01.p01CH_DL001_08_BC0   CAGTGTGTGC CCGTCTATTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACC-CT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
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08_BC.CN.06.nx2.HM067748          CAGTGTGTGC CCATCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACC-CT --TGTGGCAG T-GTGGAA-- AATCTCTAGC
08_BC.CN.98.98CN006.AF286229      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
09_cpx.CI.00.00IC_10092.AJ8665    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGA A-GTAAA--- AATCTCTAGC
10_CD.TZ.96.96TZ_BF061.AF28954    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- CCTCTCTAGC
10_CD.TZ.96.96TZ_BF071.AF28954    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- CCTCTCTAGC
10_CD.TZ.96.96TZ_BF110.AF28955    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- CCTCTCTAGC
11_cpx.CM.95.95CM_1816.AF49262    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- -----TA--- AATCTCTAGC
11_cpx.CM.96.96CM_4496.AF49262    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------C
11_cpx.CM.97.MP818.AJ291718       AAGGGGGCCC CCGCCTGTGG GGGGA----- ---------C TTTGGAAC-C TAGAAATCCC TAAAACCTTT --TAGCCCGG --GGGGAA-- AATTTTTAGC
11_cpx.FR.99.MP1298.AJ291719      TAGTGTGTGC CCATCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TAGTCCG T-GTGGAA-- AATCTCTAGC
11_cpx.FR.99.MP1307.AJ291720      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTCT ---TATTACG T-GTGGAAAA TCTATATACA
11_cpx.GR.x.GR17.AF179368         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
12_BF.AR.97.A32879.AF408629       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG G-GTGGAA-- AATCTCTAGC
12_BF.AR.97.A32989.AF408630       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
12_BF.AR.99.ARMA159.AF385936      TAGTGTGTGC CCATTTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --CTAGTCAG T-GTGGAA-- AATCTCTAGC
12_BF.UY.99.URTR23.AF385934       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTATT --TTAGTCAG T-GTGAAA-- AATCTCTAGC
12_BF.UY.99.URTR35.AF385935       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
13_cpx.CM.02.02CM_A1394.DQ8453    TGGTGTGTGC CCATCTGTTG TGTGA----- ---------C CCTGGTAT-C TAGAGATCCC TCAGACATTT A-GTCCGTGC G-GTACGGAA AATCTCTAGC
13_cpx.CM.04.04CM_173_9.DQ8453    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCTT- --TTAGTCCG T-GTGGAA-- AATCTCTAGC
13_cpx.CM.04.04CM_632_28.DQ845    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACTTT- --ATAGTCAG T-GTGGAA-- AATCTCTAGC
13_cpx.CM.96.96CM_1849.AF46097    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
13_cpx.CM.96.96CM_4164.AF46097    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
14_BG.ES.00.X605.AF450096         ---------- -------TTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAAATAG T-GTAAA--- AATCTCTAGC
14_BG.ES.00.X623.AF450097         ---------- -------TTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGATAG T-GTA---A- AATCTCTAGC
14_BG.ES.05.X1870.FJ670522        TAA-GTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGTTG- --TTGTAAA- AATCTCTAGC
14_BG.ES.05.X772_8.FJ670528       -TAAGGGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGATA- --GTGTAAA- AATCTCTAGC
15_01B.TH.99.99TH_MU2079.AF516    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----TAATTC G-CCCTTA-- AATCTCTAGC
15_01B.TH.99.99TH_R2399.AF5305    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
16_A2D.KR.97.97KR004.AF286239     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
18_cpx.CU.99.CU14.AY586541        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
18_cpx.CU.99.CU68.AY894993        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
18_cpx.CU.99.CU76.AY586540        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
19_cpx.CU.99.CU29.AY588971        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAAAAGG A-GTAA---- AATCTCTAGC
20_BG.CU.03.CB134.DQ020274        ---------- -------TTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
20_BG.CU.99.Cu103.AY586545        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
20_BG.ES.99.R77.AY586544          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
22_01A1.CM.02.02CAMLT72.EU7439    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
22_01A1.CM.06.LPH27MF.JN864049    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
22_01A1.CM.10.LB054.JN864059      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
23_BG.CU.03.CB118.AY900571        ---------- -------TTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
23_BG.CU.03.CB347.AY900572        ---------- -------TTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
24_BG.CU.03.CB378.AY900574        ---------- -------TTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACRG T-GTAAA--- AATCTCTAGC
24_BG.CU.03.CB471.AY900575        ---------- -------TTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACAACT --CTAGATAG T-GTAAA--- AATCTCTAGC
24_BG.ES.08.X2456_2.FJ670526      ------GTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
25_cpx.CM.06.06CM_BA_040.EU693    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --GTAGATAT T-GTAA---- AATCTCTAGC
25_cpx.SA.03.J11233.EU697906      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --GTAGCGAG T-GTAA---- AATCTCTAGC
25_cpx.SA.03.J11451.EU697908      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --GTAG-GAG T-GTAA---- AATCTCTAGC
26_AU.CD.02.02CD_KS069.FM87778    TGGTGTGTGC CCGTCTGTTG ATTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTACT --GAAGGTAG T-ATAAAA-- AATCTTTACC
26_AU.CD.02.02CD_LBTB084.FM877    TAGTGTGTGC CCGTCTGTTG TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGATCACT -TTTAGGCAG A-GTAAA--- AATCTCTAGC
26_AU.CD.02.02CD_MBTB047.FM877    TGGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGTAG T-GTAAA--- AATCTCTAGC
26_AU.CD.97.97CD_KTB119.FM8777    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGTAG T-GTAAA--- AATCTCTAGC
27_cpx.FR.04.04CD_FR_KZS.AM851    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
28_BF.BR.05.0679SV.JF804812       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
28_BF.BR.99.BREPM12313.DQ08587    ---------- ---------- ---------- ---------- ---------- ---------T TCTGACGCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
28_BF.BR.99.BREPM12609.DQ08587    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ------TCGC
28_BF.BR.99.BREPM12817.DQ08587    ---------- ---------- ---------- ---------- --------AC TAGAGATCCC TCAGACACTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
29_BF.BR.01.BREPM16704.DQ08587    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ------TAGC
29_BF.BR.05.0264RI.JF804807       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
29_BF.BR.99.BREPM11948.DQ08587    ---------- ---------- ---------- ---------- ---------- ---------- ------TCTT --TTAGTTCG T-GTGGAA-- AATCTCTAGC
31_BC.BR.02.110PA.EF091932        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACCT --TTWGTCAG T-GTGGAA-- AATCTCTAGC
31_BC.BR.04.04BR142.AY727527      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGATACCT --TTTGTCAG T-GTGGAA-- AATCTCTAGC
32_06A1.EE.01.EE0369.AY535660     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGAACACT --TTAGACAG T-GTAAA--- AATCTCTAGC
32_06A1.EE.02.EST2002_1169.DQ1    ---------- --GACTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGAAACAC T-TTAGACAG T-GTAAA--- AATCTCTAGC
33_01B.ID.07.JKT189_C.AB547463    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
33_01B.MY.05.05MYKL007_1.DQ366    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
33_01B.MY.05.05MYKL015_2.DQ366    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
33_01B.MY.05.05MYKL031_1.DQ366    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
33_01B.MY.05.05MYKL045_1.DQ366    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
38_BF1.UY.03.UY03_3389.FJ21378    ---------- --------TG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
38_BF1.UY.04.UY04_3987.FJ21378    ---------- --------TG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
38_BF1.UY.04.UY04_4022.FJ21378    ---------- --------TG TGTGA----- ---------C TCTGGTAC-T A-GAGATCCC TCAGACACTT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
38_BF1.UY.05.UY05_4752.FJ21378    ---------- --------TG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
39_BF.BR.03.03BRRJ103.EU735534    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
39_BF.BR.03.03BRRJ327.EU735536    ---------- ---------- ---------- ---------- ---------- ---------- --------TT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
39_BF.BR.04.04BRRJ179.EU735535    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
40_BF.BR.04.04BRRJ115.EU735538    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
40_BF.BR.04.04BRSQ46.EU735540     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
40_BF.BR.05.05BRRJ055.EU735537    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
40_BF.BR.05.05BRRJ200.EU735539    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCAT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
42_BF.LU.03.luBF_01_03.EU17015    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
42_BF.LU.03.luBF_02_03.EU17015    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
42_BF.LU.03.luBF_04_03.EU17014    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
42_BF.LU.03.luBF_05_03.EU17015    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAYT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
42_BF.LU.03.luBF_06_03.EU17014    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
42_BF.LU.03.luBF_07_03.EU17014    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
42_BF.LU.03.luBF_08_03.EU17014    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCCG A-GTGGAA-- AATCTCTAGC
42_BF.LU.03.luBF_09_03.EU17015    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAG-WCCC TCAGACCWTT --TTAGTCCG K-GKGGAA-- AATCTCTAGC
42_BF.LU.04.luBF_11_04.EU17014    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
42_BF.LU.04.luBF_12_04.EU17014    TGGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
42_BF.LU.05.luBF_13_05.EU17013    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
42_BF.LU.05.luBF_14_05.EU17014    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
42_BF.LU.05.luBF_16_05.EU17015    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
42_BF.LU.05.luBF_17_05.EU17014    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTAGTCCG K-GTGGAA-- AATCTCTAGC
42_BF.LU.06.luBF_18_06.EU17013    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
42_BF.LU.06.luBF_19_06.EU17013    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
43_02G.SA.03.J11223.EU697904      TAGTGTGTGC CCATCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGG T-GTAAA--- AATCTCTAGC
43_02G.SA.03.J11232.EU697905      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG T-GTGAA--- AATCTCTAGC
43_02G.SA.03.J11243.EU697907      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CCAGACTG T-GTGAA--- AATCTCTGGC
43_02G.SA.03.J11456.EU697909      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG T-GTAAA--- AATCTCTAGC
44_BF.CL.00.CH80.FJ358521         ------GTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
44_BF.CL.01.CH12.AY536235         TAGTGTGKGC CCGTCTGTTG TTTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTRGAA-- AATCTCTAGC
45_cpx.CD.97.97CD_MBFE185.FN39    CAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TYTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
45_cpx.CD.97.97CD_MBS30.FN3928    CAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGG T-ATAAA--- AATCTCTAGC
45_cpx.CM.97.97CM_MP814.FN3928    CAGTGTGTGC CCGTYTGTTG TGTGA----- ---------C TTTGGTAA-C TAGAGATCCC TCAGACCACT --YTAGRCGG T-GTAAA--- AATCTYTAGC
45_cpx.FR.04.04FR_AUK.EU448295    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGCGG T-GTAAA--- AATCTCTAGC
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45_cpx.GA.97.97GA_TB45.FN39287   CAGTGTGTGC CCATYTATTG A--------- ---------Y TCTGGTAA-C TAGAGATC-- ---------- ---------- --ATATAAA- AATCTCTAGC
46_BF.BR.01.01BR087.DQ358801     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
46_BF.BR.01.01BR125.DQ358802     TAGTGTGTGC CCGTCTGTTG TATGA----- ---------C TCTGGTAA-C TAAAGATCCC TCAGAACATT --TTAGTCTG T-GTGGAAA- AATCTCTAGC
46_BF.BR.07.07BR_FPS625.HM0264   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGAACATT --TTAGTCCG A-GTGGAA-- AATCTCTAGC
46_BF.BR.07.07BR_FPS742.HM0264   ---------- ---------- ---------- ---------- ---------A CTGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
46_BF.BR.07.07BR_FPS783.HM0264   TTGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGAACATT --TTAGTCCG G-GTGGAA-- AATCTCTAGC
46_BF.BR.07.07BR_FPS810.HM0264   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGAACATT --TTAGTCTG G-GTAGAA-- AATCTCTAGC
46_BF.BR.07.07BR_FPS812.HM0264   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACATTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
47_BF.ES.08.P1942.GQ372987       --AGGGTTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TAGTCAG T-GTGGAAA- AATCTCTAGC
47_BF.ES.08.X2457_2.FJ670529     -AAGGGTTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
49_cpx.GM.02.N18380.HQ385477     ---------- --ATCTGTGG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTCT --CTAGACG- --GTGTAAA- AATCTCTAGC
49_cpx.GM.03.N26677.HQ385479     ---------- --ATCTGTGG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCACT --CTAGACG- --GTGTAAA- AATCTCTAGC
49_cpx.GM.97.N28353.HQ385478     --GTGTGTGC CCATCTGTAG TGTGTGA--- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGAAG- --GTGTAAA- AATCTCTAGC
52_01B.MY.03.03MYKL018_1.DQ366   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
53_01B.MY.04.04MYKL016_1.DQ366   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
53_01B.MY.10.10MYKJ067.JX39061   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
53_01B.MY.10.10MYKJ079.JX39061   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
53_01B.MY.11.11FIR164.JX390610   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
54_01B.MY.07.07MYKLD49.EU03191   ---------- ---------- ---------- ---------- ---------- ---------- ---------- -----GCAAC C-TCAATA-- AATCTCTAGC
54_01B.MY.08.08MYKL044.JX39097   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
54_01B.MY.09.09MYSB023.JX39097   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- -------A-- AATCTCTAGC
55_01B.CN.10.HNCS102056.JX5746   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
55_01B.CN.11.GDDG318.JX574662    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01A1.MM.99.mCSW105.AB097872      TGGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGAAGG T-GTAAA--- AATCTCTAGC
01B.CN.08.08CYM047.JF340054      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01B.JP.x.pHIV_1_Y271B01AE64.AB   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT T-TCAGTTTG T-GTAAA--- AATCTCTAGC
01B.MM.00.mIDU502.AB097865       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- -AATCTCTAGC
01B.MM.99.mCSW104.AB097867       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
01B.MY.04.04MYKL019_1.DQ366665   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01B.MY.05.05MYKL043_1.DQ366666   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01B.MY.06.06MMYKLD46.EF495062    TGGTGTGTGC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-TTAAA--- AATCTCTAGC
01B.MY.07.07MYKLD47.EU031913     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01B.MY.07.07MYKLD48.EU031914     TGGTGTGTGC CCGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-TTAAA--- AATCTCTAGC
01B.TH.05.05TH140456.JN631793    ---------- ---------- ---------- ---------- -----TAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01B.TH.05.AA095a_WG21.JX447830   ---------- -CGTCTGTGT TAGGA----- ---------C TCTGGTAA-C TAGAGTTCCC TCAGACCACT --CTAGACGA A-GTAAA--- AATCTCTAGC
01B.TH.06.AA020a_wg2.JX446927    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01B.TH.06.AA025a_WG13.JX447000   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACC --CTAGACTG A-GTAAA--- AATCTCTAGC
01B.TH.06.AA084a_WG10.JX447668   ---------- ---------- ---------- ---------- --TGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
01B.TH.91.CM237.AY167123         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCTTG --TTAGTCAG T-GTGGAA-- AATCTCTAGC
01BC.MM.00.mCSW503.AB097866      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01BC.MM.99.mIDU107.AB097868      CAGTGTGTGC CCGTCTATTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
01F2G.CM.02.LT31.JN864056        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
0206.DZ.10.DEURF10DZ001.JX1406   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
0206.GH.03.03GH195AG_06.AB2868   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTACT --CTAGACGG T-GTAA---- AATCTCTAGC
0206.NE.00.NE36.AJ508597         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACCG C-GTAAA--- AATCTCTAGC
0206.NE.00.NE95.AJ508596         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --GTTGATAG T-GTAAA--- AATCTCTAGC
0206.NE.97.NE03.AJ508595         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGCGG T-GTAAA--- AATCTCTAGC
0209.CI.01.01IC_17395.AJ866554   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG T-GTAAAA-- AATCTCTAGC
0209.CI.01.01IC_PCI118.AJ86655   TAGTGTGTGC CCGTCTGGTG TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG T-GTAAA--- AATCTCTAGC
0209.CI.01.01IC_PCI127.AJ86655   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGG T-GTAAA--- AATCTCTAGC
0209.CI.97.97IC_PCI3.AJ866555    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGA G-GTAAA--- AATCTCTAGC
0222.CM.02.02CAMLT04.EU743964    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
0222.CM.08.BDSH129.JN864052      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
0222.CM.10.LB045.JN864053        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
02A1.ES.05.X230_10.FJ670515      TTAAGGGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --CTAGACGG T-GTAAA--- AATCTCTAGC
02A1.ES.07.ES_P1751.GQ372986     -AAGGGGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGG T-GTAAA--- AATCTCTAGC
02A1.GH.03.GHNJ176.AB231894      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGTCTG C-GTAAA--- AATCTCTAGC
02A1.GH.97.97GH_AG2.AB052867     TAGTGTGTGC CCGTCTGTTG TGTAA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGTTG T-GTAAA--- AATCTCTAGC
02A1.RU.10.10RU6637.JN230353     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGCAG T-GTAAA--- AATCTCTAGC
02B.ES.99.99SP_11339.DQ926899    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAT-C TAGAGATCCC TCAGACCACT --CGAGATTG A-GTAGAA-- AATCTCTAGC
02B.FR.02.URF4.JN882652          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
02B.FR.09.URF2.JN882654          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
02BD.FR.08.URF1.JN882653         -----TGTGC CCGTCTGTTG TGCGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG T-GTAAA--- AATCTCTAGC
02BG.FR.10.URF5.JN882655         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
02C.BE.93.VI1035.AJ276595        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
02D.GH.03.GHNJ193.AB231897       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGTCTG C-GTAAA--- AATCTCTAGC
02D.GH.03.p03GH194AG09.AB48004   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --GTAGTCAC T-GTAGAA-- AATCTCTAGC
02F2G.CM.02.LT66.JN864057        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
02GK.CI.01.01IC_PCI123.AJ86655   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTTA-C TAGAGATCCC TCAGACCACT --CTAGACTG T-GTAAA--- AATCTCTAGC
06A1.BJ.x.B76.AJ293865           ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
1819.CU.99.CU64.AY894995         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGG --GTGTAAA- AATCTCTAGC
26C.CD.02.02CD_LBTB032.FM87777   TAGTGTGTGC CCGTCTGTTG TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --CTAGGCAG A-GTAAA--- AATCTCTAGC
26C.CD.97.97CD_KFE267.FM877778   TAGTGTGTGC CCGTCTGTTA TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TTAGACCACT --CTAGGCAG T-GTGAAA-- AATCTCTAGC
26C.CD.97.MBFE250.FM877783       TAGTGTGTGC CCGTCTGTTG TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGATCACT --CTAGTTAG T-GTGAA--- AATCTCTAGC
A1B.EE.02.EST2002_394.DQ167216   ---------- ---------G TATGA----- ---------C TCTGGTAA-C TAGAGATCCT TCAGACCACT --CTAGGCGG T-GTAAA--- AATCTCTAGC
A1BD.GB.03.34567.JN417238        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---CTCTAGC
A1C.CA.04.04CA7750.EU220698      TAGTGTGTGC CCGTCTGTTG TTTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG T-GTAAA--- AATCTCTAGC
A1C.IN.01.1579A.DQ083238         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGGCGG T-GTAAA--- AATCTCTAGC
A1C.IN.95.95IN21301.AF067156     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
A1C.RW.92.92RW009_06.U88823      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1CD.KE.02.ML2000.EU110093       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1CD.SE.95.SE8603.AF075702       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1CDGKU.ZA.99.CM4_99ZACM4.AF41   TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT ---TTGGTAG T-GTGGAA-- AATCTCTAGC
A1CG.KE.02.ML1979PCR.EU110096    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1D.DK.96.FSA.DQ912822           ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1D.DK.98.MA.DQ912823            ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1D.GB.00.8179.JN417236          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --TCTCTAGC
A1D.GB.03.33365.JN417239         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----TCTAGC
A1D.GB.10.11762.JN417241         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- -------A-- AATCTCTAGC
A1D.GB.10.12792.JN417240         ---------- ---------- ---------- ---------- ------AA-C TAGAGATCCC TCAGACCACT --CTAGACGG T-GTAAA--- AATCTCTAGC
A1D.KE.02.ML1974.EU110090        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1D.SE.93.SE6954.AF075701        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1D.TZ.96.TZBFL0011.AF442569     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1D.TZ.96.TZBFL0088.AF442570     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1D.TZ.97.TZBFL0086.AF442566     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1D.UG.07.p191947.JX236674       TAGTGTGTGC CCGTCTATTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACTG A-GTAAA--- AATCTCTAGC
A1D.UG.08.p191982.JX236675       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTCT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
A1D.UG.90.UG266.AY352657         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1D.UG.92.UG035.AY352656         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1DHK.NO.97.97NOGIL3.AJ237565    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --ATAGTCAG T-GTGGAA-- AATCTCTAGC
A1DK.CD.85.MAL_MALCG.X04415      CAGTGTGTGC CCATCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGG T-GTAAA--- AATCTCTAGC
A1G.BE.94.VI1197.AJ276596        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1G.NG.92.92NG003.U88825         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1G.RU.11.11RU6900.JX500697      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --CTAGAGTG T-GTAAA--- AATCTCTAGC
A1G.RU.11.11RU6939.JX500706      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --TTAGATTG A-GTAAA--- AATCTCTAGC
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A1GHU.GA.x.VI354.AF076474         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1GJ.BW.98.BW2117.AF192135        ---------- ---------- ---------- ---------- ---------- ---------- ---------- --AGGGGGCC A-AGTCGG-- CCTCTCTAGC
A2C.ZA.98.DU178_98ZADU178.AF41    TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --TGTGGTAG T-GTGGAA-- AATCTCTAGC
A2C.ZM.90.ZAM184.U86780           ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
A2CD.KE.01.ML1956.EU110089        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
AC.KE.97.ML672.AY322191           TAGTGTGTGC CCATCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACGG T-GTAAA--- AATCTCTAGC
AC.ZA.04.04ZAPS204B1.DQ093606     ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --GTTGGTAG T-GTGGAA-- AATCTCTAGC
AGKU.CD.x.BCF_Kita.AB485665       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --TTAGAGCA C-GTAAA--- AATCTCTAGC
BC.AR.01.ARE195FL.AY968312        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
BC.BR.01.01_BR_RGS45.GQ365651     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACTT --GTAGTCAG T-GTGGAA-- AATCTCTAGC
BC.BR.01.01_BR_RGS69.GQ365652     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACCT ---TTGTCAG T-GTGGAA-- AATCTCTAGC
BC.BR.02.02_BR_HF31.GQ365649      TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACCT --TTTGTCAG T-GTGGAA-- AATCTCTAGC
BC.BR.05.05_BR_NSP24.GQ365650     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACCT --TGTGTTAG T-GTGGAA-- AATCTCTAGC
BC.BR.06.06BR1114.JN692478        TAGTGTGTGC CCGTCTGTTT TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACCT --TTTGTCAG T-GTGGAA-- AATCTCTAGC
BC.BR.92.92BR023.HM100716         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACCT --TTTGTCAG T-GTGGAA-- AATCTCTAGC
BC.CN.07.309.HM776938             ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------G
BC.CN.07.341.HM776939             ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
BC.IN.02.INDNARI_0218440.EU000    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
BC.IN.02.NARI7_3.EU000511         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
BC.IN.02.NARI9_3.EU000508         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
BC.IN.99.NARI10_2.EU000516        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------A- AATCTCTAGC
BC.MM.99.mIDU103.AB097873         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- --------TT
BCU.FR.06.06FR_CRN.EU448296       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACCT --TTTGTCAT T-GTGGAA-- AATCTCTAGC
BF.BR.03.BREPM1026.EF637055       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
BF.BR.03.BREPM1029.EF637052       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
BF1.BR.00.BREPM13853.DQ085875     ---------- ---------- ---------- ---------A TCTGGTAC-T A-GAGATCCC TCAGACCATT --TGAGTCAG T-GTGGAA-- AATCTCTAGC
BF1.BR.01.01BR042.DQ358799        ---------- ---------- ---------- ---------- -----AA--C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
BF1.BR.01.01BR047.DQ358800        ---------- ---TCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --GTAGTCAG T-GTGGAA-- AATCTCTAGC
BF1.BR.01.01BR226.DQ358803        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT ---TAGTCAG T-GTGGAA-- AATCTCTAGC
BF1.BR.01.01BR323.DQ358804        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
BF1.BR.02.02BR005.DQ358806        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
BF1.BR.02.02BR006.DQ358807        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
BF1.BR.02.02BR033.DQ358811        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
BF1.BR.02.02BR034.DQ358812        TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
BF1.BR.02.02BR2028.JN692437       ---------- ---------- ---------- ---------- ---------- -----CCCTC AGACCTTTTA -----GTCAG T-GTGGAA-- AATCTCTAGC
BF1.BR.03.03BR2018.JN692449       TAGTGTGTGC CCGTCTGTTG TGTG------ ---------C TCTGGTAA-C TATAGATCCC TCAGACAATT --CTAGTCAG T-GTGGAA-- AATCTATAGC
BF1.BR.03.03BR2019.JN692448       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
BF1.BR.04.04BR1067.JN692456       TAGTGTGTGC CCGTCTGTTG TATGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCAAT --TTAGTCGG A-GTAGAA-- AATCTCTAGC
BF1.BR.05.0008SP.JF804805         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
BF1.BR.05.0632SV.JF804810         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
BF1.BR.06.06BR_FPS561.HM026455    AAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TAGTCAGT G-TGGAA--- AATCTCTAGC
BF1.BR.99.BREPM107.AY771588       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
BF1.BR.99.BREPM108.AY771589       TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATC --TTAGTCAG T-GTGGAA-- AATCTCTAGC
BF1.BR.99.BREPM11931.DQ085869     ---------- ---------- ---------- ---------C TGGTACT--- --GAGATCCC TCTGACCTTT ---TAGTCCG T-GTGGAA-- AATCTCTAGC
BF1.ES.08.ES_X2524_2.GQ372989     TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACTATT --TTAGTCAG A-GTAGAA-- AATCTCTAGC
BF1.ES.08.X2432_2.FJ853621        -AGTGTGTGC CCGTCTGTTG TGTGACTCTG GTAACTAA-C TCTGGTGA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
BF1.ES.09.DEMBF09ES003.JX14066    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
BF1.ES.09.DEMBF09ES006.JX14066    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
BF1.IT.01.53143.GU595149          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
BF1.IT.02.30638.GU595148          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
BF1.IT.02.57954.GU595150          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-A TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
BF1.IT.02.58736.GU595160          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TAAGTCAG T-GTGGAA-- AATCTCTAGC
BF1.IT.02.59211.GU595151          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACACTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
BF1.IT.05.83166.GU595152          TAGTGTGTGC CCATCTGTTG TTTGA----- ---------C TCCGGTTA-C TAGAGATCCC TCAGACCATT --TTAGTCCG T-GTGGAA-- AATCTCTAGC
BF1.IT.06.89072.GU595153          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGACCCC TCAGACCCTT --TTAGCCAG T-GTGGAA-- AATCTCTAGC
BF1.JP.04.DR6082.AB480298         TAGTGTGTGC CCATCTGTTG TGTGA----- ---------C TCTGGTAG-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
BF1.JP.04.DR6190.AB480300         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCATT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
BF1.JP.x.DR0769.AB253430          TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCCTT --TTAGTCAG T-GTGGAA-- AATCTCTAGC
BFG.MO.05.MO108.GU207884          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
BG.CU.x.Cu100.AY586546            TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGACAG T-GTAAA--- AATCTCTAGC
BG.DE.01.9196_01.AY882421         TAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAAATAG T-GTAA---- AATCTCTAGC
CD.KE.01.ML1076.EU110086          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
CU.JP.04.DR5782.AB286849          CAGTGTGTGC CCGTCTGTTG TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCTTT --TAGTCAGT --GTGGAA-- AATCTCTAGC
DF1G.ES.04.X963_4.FJ670527        -TAAGGGTGC CCGTCTGTTA TGTGA----- ---------C TCTGGTAA-C TAGAGATCCC TCAGACCACT --CTAGATAG T-GTAAA--- AATCTCTAGC
DO.FR.08.RBF208.GQ351296          ---------- TCATCTGTT- --CAA----- ---------C CCTGGTAT-C TAGAGATCCC TCAGATCACG --TAGACTGA G-GYAGAA—AATCTCTAGC
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                                  ....|....| ....|....| ....|....| ....|....| ....|....| ....|....| ....|....| ....|....| ....|....| ....|....|
                                     105        115        125        135        145        155        165        175        185        195
B.FR.83.HXB2_LAI_IIIB_BRU.K034    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG GGAAA----- ---------- ---------- ----------
A1.CD.02.02CD_KTB035.AM000055     AT-TGGCGCC CAACCAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.CD.97.97CD_KCC2.AM000053       AG-TGGCGCC CGAACAGGGA CTCG------ ---------- ---AAAGCGA AA-GTAAAAG TT-------- ---------- ---------- ----------
A1.ES.05.X1608_8.FJ670519         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.IT.02.60000.EU861977           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.KE.01.ML1901PCR2.EU110095      ---------- ---------- ---------- ---------- ---------- --------TG TT-------- ---------- ---------- ----------
A1.KE.01.ML1945.EU110088          ---------- ---------- ---------- ---------- ---------- ---------- -T-------- ---------- ---------- ----------
A1.KE.02.ML1990.EU110092          ---------- ---------- ---------- ---------- ---------- ---------- TT-------- ---------- ---------- ----------
A1.KE.02.ML2014.EU110094          ---------- ---------- ---------- ---------- ---------- ---------- -T-------- ---------- ---------- ----------
A1.KE.86.ML170_1986.AF539405      ---------- ---------- ---------- ---------- -------GGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.KE.94.Q23_17.AF004885          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.KE.97.ML013_2.AY322185         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.KE.97.ML605_3.AY322190         AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTAAGC AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.KE.97.ML752.AY322193           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.RU.00.RU00051.EF545108         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1.RU.05.RU_560_1125_JA.JQ2928    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1.RU.06.RU_915_1016.JQ292896     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.RU.06.RU_915_1035.JQ292897     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.RU.06.RU_915_1038.JQ292898     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.RU.06.RU_915_1041.JQ292899     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.RU.06.RU_SP_B_049.JQ292900     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.RU.07.Irkutsk_5.JQ292891       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1.RU.08.PokA1Ru.FJ864679         AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAC AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.RU.08.RUA001.JQ292893          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.RU.08.RUA007.JQ292894          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.RU.10.10RU6617.JX500696        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TG---TT--- ---------- ---------- ----------
A1.RU.10.10RU6792.JX500695        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.RU.11.11RU6950.JX500694        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.RW.07.pR463F.JX236677          AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTTGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.RW.07.pR880F.JX236678          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TG---TT--- ---------- ---------- ----------
A1.RW.92.92RW008.AB253421         AG-TGGCGCC CGAACAGGGA CTTGAAA-AG CGAAAGTAAC AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.RW.92.92RW025A.AB287376        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
A1.RW.93.93RW037A.AB287379        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.SE.95.UGSE8131.AF107771        AG-TGGCGCC CGAACAGGGA CCCGAAA-AG TGAAAGTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.TZ.97.97TZ03.AF361873          ---------- ------GGAC CCTAA---AG TGAAAGTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.UA.00.98UA0116.AF413987        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.UG.07.p191084.JX236669         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TG---TT--- ---------- ---------- ----------
A1.UG.07.p191845.JX236671         AG-TGGCGCC CGAACAGGGA CTTTAAAAGC GAAAGT-AAC AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
A1.UG.07.p9004SDM.JX236676        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.UG.85.U455_U455A.M62320        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.UG.92.92UG037.AB253429         AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGT-AAT AGGGACTCGA AA-ACGAAAG TT-------- ---------- ---------- ----------
A1.UG.92.UG029.AB098333           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.UG.x.UG031.AB098331            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.UG.x.UG275.AB485632            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1.ZA.04.04ZASK162B1.DQ396400     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A2.CD.97.97CDKTB48.AF286238       ---------- ---------- ---------- ---------- ---TTTTTCG AA-GCGAAG- TA-------- ---------- ---------- ----------
A2.CY.94.94CY017_41.AF286237      AA-AGCGAAA GTAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
B.AU.86.MBC200.AF042100           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG GAAAA----- ---------- ---------- ----------
B.AU.87.MBC925.AF042101           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGT AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.AU.95.C24.AF538304              AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.AU.95.C42.AF538305              AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCGGA AA-GCGAAAG AAAAA----- ---------- ---------- ----------
B.AU.95.C76.AF538306              AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GTCTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.AU.95.C92.AF538307              AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.AU.95.MBCC54.AF042103           --------CC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.AU.96.MBCC98.AF042104           ---------- ---------- ---------- ---------- ---------- ---------G GAAA------ ---------- ---------- ----------
B.AU.96.MBCD36.AF042105           ---------- ---ACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG GGAAA----- ---------- ---------- ----------
B.AU.99.1181.AF538302             AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.AU.x.2870718.AY857022           ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.AU.x.9125091.AY857165           ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.AU.x.VH_VHPCR.AF146728          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.BO.09.DEMB09BO001.JX140656      -G-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.02.02BR008.DQ358808          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GTGAAAG TAGAA----- ---------- ---------- ----------
B.BR.02.02BR011.DQ358809          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
B.BR.02.02BR013.DQ358810          AG-TGGCGCG CGAACAGG-- ---------- ---------- ---GACTAGA AA-GCGAAAG TAGAC----- ---------- ---------- ----------
B.BR.02.02BR1013.JN692432         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.02.02BR2025.JN692435         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AAAGCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.02.02BR2032.JN692439         ---------- ---------- ---------- ---------- ---------- ---------- ----T----- ---------- ---------- ----------
B.BR.02.02BR2033.JN692440         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AAAGCGAAAG TAAAA----- ---------- ---------- ----------
B.BR.02.02BR2041.JN692443         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
B.BR.02.02BR2042.JN692444         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.02.04BR1064.JN692433         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTAGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
B.BR.02.05BR1094.JN692470         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.03.03BR1020.JN692445         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTTA AAAGCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.03.03BR1046.JN692447         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
B.BR.03.BREPM1023.EF637057        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG GAGAA----- ---------- ---------- ----------
B.BR.03.BREPM1024.EF637056        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.03.BREPM1027.EF637054        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
B.BR.03.BREPM1028.EF637053        ---------- ---------- ---------- ---------- -----CTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.03.BREPM1032.EF637051        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.BR.03.BREPM1033.EF637050        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.03.BREPM1035.EF637049        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.BR.03.BREPM1038.EF637048        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAACG----- ---------- ---------- ----------
B.BR.03.BREPM1040.EF637047        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTAGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
B.BR.03.BREPM2012.EF637046        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.BR.04.04BR1047.JN692450         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGG AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.04.04BR1049.JN692451         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.BR.04.04BR1051.JN692452         AG-TGGCGCC TGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
B.BR.04.04BR1054.JN692453         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.BR.04.04BR1055.JN692454         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.04.04BR1057.JN692455         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAAGA----- ---------- ---------- ----------
B.BR.04.04BR1068.JN692457         AG-TGGCGCC --GAAAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.04.BREPM1066.FJ195090        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
B.BR.04.BREPM1070.FJ195086        GG-TGGCGCC GGACCAGG-- ---------- ---------- ---GCCTTGA AA-AGGAAAT AAGAA----- ---------- ---------- ----------
B.BR.05.05BR1077.JN692460         ---------- ---------- ---------- ---------- ---------- ---------- ----T----- ---------- ---------- ----------
B.BR.05.05BR1078.JN692461         ---------- ---------- ---------- ---------- ---------- -----AACTC TTAAT----- ---------- ---------- ----------
B.BR.05.05BR1079.JN692462         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG CGGAA----- ---------- ---------- ----------
B.BR.05.05BR1080.JN692463         ---------- ---------- ---------- ---------- ---------- ---------- ----T----- ---------- ---------- ----------
B.BR.05.05BR1082.JN692465         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.BR.05.05BR1089.JN692467         ---------- ---------- ---------- ---------- ---------- ---------- ----T----- ---------- ---------- ----------
B.BR.05.05BR1092.JN692468         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTAA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.05.05BR1095.JN692471         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.BR.05.05BR1101.JN692473         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCGGA AA-ACGAAAG TAAAA----- ---------- ---------- ----------
B.BR.05.05BR1104.JN692474         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAGAA----- ---------- ---------- ----------
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B.BR.05.05BR1107.JN692475        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.BR.05.BREPM1081.FJ195091       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAAAA----- ---------- ---------- ----------
B.BR.05.BREPM1084.FJ195088       AG-TGGCGCC GGAACAGG-- ---------- ---------- ---GACTTGA GA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.05.BREPM1093.FJ195089       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.06.06BR1115.JN692479        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.BR.06.06BR1119.JN692480        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.CA.06.502_1027_wg01.JF320413   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG AGAAA----- ---------- ---------- ----------
B.CA.06.502_1799_FL02.JF320427   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCGGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.CA.07.502_1191_03.JF320424     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.CA.96.WC10C_4.AY314056         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.CA.97.CANB3FULL.AY779553       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.CH.03.HIV_CH_BID_V4470_2003.   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.CN.01.CNHN24.AY180905          AG-TGGCGGC CGAACAGG-- ---------- ---------- ---GACTAGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.02.02HNsc11.DQ007903        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.03.SHXDC0081.JF932492       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.05.05CNHB_hp3.DQ990880      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.06.CC056.JF932482           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TGAGA----- ---------- ---------- ----------
B.CN.07.AH070011.JF932468        ---------- ---------- ---------- ---------- -GGGACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.07.AH070014.JF932469        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TGAGA----- ---------- ---------- ----------
B.CN.07.AH070017.JF932470        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.07.AH070018.JF932471        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
B.CN.07.AH070057.JF932472        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
B.CN.07.BJ070030.JF932473        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
B.CN.07.CBJC261.JF932474         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.07.CBJC392.JF932475         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.07.CBJC394.JF932476         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.07.CBJC396.JF932477         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
B.CN.07.GS070017.JF932484        ---------- ---ACAGG-- ---------- ---------- ---GACGCGA AA-ACGAAAG TGAGA----- ---------- ---------- ----------
B.CN.07.GZ070002.JF932485        ---------- ---ACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.07.GZ070030.JF932486        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
B.CN.07.HB070006.JF932487        ---------- ----CAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.07.HB070022.JF932488        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
B.CN.07.HB070035.JF932489        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.07.JL070038.JF932490        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.CN.07.SX070080.JF932493        ---------- ---------- ---------- ---------- AGGGACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.07.hb070025.JF932499        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
B.CN.07.hen1345.JF932500         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.08.CBJC476.JF932478         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.08.CBJC489.JF932479         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.08.CBJC500.JF932480         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.08.CBJC503.JF932481         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.08.cbjc468.JF932498         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TGAGA----- ---------- ---------- ----------
B.CN.09.YN09P0014.JF932494       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.09.ZK042.JF932497           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.10.DEMB10CN002.JX140658     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.CN.98.YN9802.JF932495          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.98.YN9838.JF932496          ---TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TGAGA----- ---------- ---------- ----------
B.CN.99.plwj.GU177863            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CN.x.RL42.U71182               AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
B.CU.99.Cu19.AY586542            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.CU.99.Cu43.AY586543            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAGAA----- ---------- ---------- ----------
B.DE.86.D31.U43096               AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
B.DE.86.HAN.U43141               AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.DK.01.CTL_016.EF514704         ---------- ---AC-AG-- ---------- ---------- ---GACCGGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.DK.01.CTL_017.EF514705         AG-TGGCGCC CCGAACAG-- ---------- ---------- --GGACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.DK.01.CTL_018.EF514706         AG-TGGCGCC CCGGACAG-- ---------- ---------- --GGACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.DK.01.CTL_023.EF514707         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.01.CTL_030.EF514708         AG-TGGCGCC CGACAGGG-- ---------- ---------- ---AC-GCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.DK.01.CTL_033.EF514709         AG-TGGCGCC CGAAACAG-- ---------- ---------- --GGACYTAA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.DK.01.CTL_035.EF514710         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.01.CTL_041.EF514711         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.01.CTL_043.EF514712         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.DK.04.PMVL_012.EF514699        GT-KGCGCCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.DK.04.PMVL_013.EF514700        -T-GCGGCCC CGAACAGG-- ---------- ---------- --GACCTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.DK.04.PMVL_018.EF514697        ---KGCGCCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.DK.04.PMVL_025.EF514702        GT-KGCGCCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.DK.04.PMVL_027.EF514698        ---GSSGCCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.DK.04.PMVL_039.EF514703        --------CC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG AGAAA----- ---------- ---------- ----------
B.DK.04.PMVL_049.EF514701        ---------- -GAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.DK.07.PMVL_011.FJ694790        ---------C CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG AGAAA----- ---------- ---------- ----------
B.ES.05.X1890.EU786672           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG AGAAA----- ---------- ---------- ----------
B.ES.06.X1959.EU786675           AG-TGGCGCC CGAACASG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.ES.06.X2102.EU786677           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GTGAAAG TTGAA----- ---------- ---------- ----------
B.ES.07.X2149.EU786678           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCGGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.ES.07.X2210_3.EU786679         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG GGAAA----- ---------- ---------- ----------
B.ES.07.X2231.EU786680           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GTGAAAG AGGAA----- ---------- ---------- ----------
B.ES.08.ES_X2515_3.GQ372988      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AAAGCGAAAG TAGAA----- ---------- ---------- ----------
B.ES.08.ES_X2556_3.GQ372990      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.ES.08.P2008.FJ670531           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG GAAAA----- ---------- ---------- ----------
B.ES.08.X2425_2.FJ670525         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.ES.08.X2510_2.FJ853622         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
B.ES.08.X2555_2.GU362883         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.ES.08.X2574_2.GU362886         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.ES.09.P2149_3.GU362881         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
B.ES.09.X2689_2.GU362885         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.ES.89.U61.DQ854716             AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.FR.05.DEMB05FR001.JX140652     ---------- -GAACAGG-- ---------- ---------- --GACCCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.FR.08.DEMB08FR002.JX140654     ---------- ----CAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TTGAA----- ---------- ---------- ----------
B.GA.88.OYI_397.M26727           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTTA AA-GTGAAAG TGGAA----- ---------- ---------- ----------
B.GB.05.MM45d213_GN1.HM586212    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AAAGTGAAAG AGAAA----- ---------- ---------- ----------
B.GB.83.CAM1.D10112              AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
B.GB.86.GB8_46R.AJ271445         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TTGAA----- ---------- ---------- ----------
B.GB.x.MANC.U23487               AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.HK.06.HK003.FJ460500           ---------C CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG AGAA------ ---------- ---------- ----------
B.HK.06.HK004.FJ460501           ---------C CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.IN.x.11807.EF694037            ---------- ---------- ---------- ---------- ---GACTTGA AA-GTGAAAG GAAAA----- ---------- ---------- ----------
B.IT.05.SG1.DQ672623             AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.JP.00.DR2508.AB289588          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.JP.01.DR388.AB289590           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.JP.04.04JPDR6075B.AB221125     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.JP.04.DR5913.AB480696          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.JP.04.DR6089.AB286955          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.JP.04.DR6174.AB480692          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
B.JP.04.DR6175.AB480694          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.JP.05.DR6538.AB287363          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAGAA----- ---------- ---------- ----------
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B.JP.05.DR6737.AB287364          AG-TGGCGCC TGAACAGG-- ---------- ---------- ---GACCTGA AA-ACGAAAG CGGAA----- ---------- ---------- ----------
B.JP.05.DR7060.AB287366          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
B.JP.05.DR7065.AB287368          AG-TGGCGCC CGAACAGG-- ---------- ---------- --GACCTTGA AA-GCGAAAG TGGAA----- ---------- ---------- ----------
B.JP.98.DR1120.AB480698          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-TCGAAAG TAGAA----- ---------- ---------- ----------
B.JP.99.DR1348.AB287370          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
B.JP.x.DR1673.AB564745           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAG----- ---------- ---------- ----------
B.JP.x.DR1712.AB604946           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG CGGAA----- ---------- ---------- ----------
B.JP.x.DR1777.AB604948           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.JP.x.JRC03B.AB565496           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAAGA----- ---------- ---------- ----------
B.JP.x.JRC05B.AB565497           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG CAAAA----- ---------- ---------- ----------
B.JP.x.JRC65B.AB565502           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-ACGAAAG TGAGA----- ---------- ---------- ----------
B.JP.x.pJPDR0796B02.AB565478     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.JP.x.pJRC57B09.AB641836        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
B.KR.02.02OSG1.JQ429433          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGACAG TAGAA----- ---------- ---------- ----------
B.KR.03.03HJY8.JQ316131          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GGCCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.KR.03.03KDE11.JQ316128         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.KR.03.03KGS5.JQ316132          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.KR.03.03LSH1.JQ316127          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.KR.03.03YGS3.JQ316135          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAGAA----- ---------- ---------- ----------
B.KR.04.04CWS5.JQ316133          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
B.KR.04.04KJS8.JQ316130          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.KR.04.04KMH5.DQ295193          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.KR.04.04KMK5.JQ316126          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.KR.04.04LHS6.AY839827          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGGTA----- ---------- ---------- ----------
B.KR.04.04LSK7.DQ295192          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.KR.04.04WK7_HIV_1_wk.DQ29519   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
B.KR.05.05CSR3.DQ837381          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG GAAAA----- ---------- ---------- ----------
B.KR.05.05YJN2.JQ316134          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
B.KR.07.07KYY4.JQ341411          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGTA----- ---------- ---------- ----------
B.KR.99.99HYH2.JQ316129          AG-TGGCGCC CGAACAGG-- ---------- ---------- --GACCGAGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.MM.99.mSTD101.AB097870         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
B.NL.00.671_00T36.AY423387       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AG----AAAG TAAAA----- ---------- ---------- ----------
B.NL.86.3202A21_ACH3202A21.U34   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.NL.96.H434_42_A1.AY970948      AA-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.PE.06.502_0491_wg5.JF320183    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG GGAAA----- ---------- ---------- ----------
B.PE.06.502_0524_FL04.JF320008   ---------- ---------- ---------- ---------- ---------A AA-GCGAAAG CAAAA----- ---------- ---------- ----------
B.PE.06.502_0648_FL02.JF320215   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCGGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.PE.06.502_0841_FL04.JF320208   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.PE.06.502_2794_FL05.JF320244   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAA------ ---------- ---------- ----------
B.PE.07.502_0525_wg5.JF320191    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAAAA----- ---------- ---------- ----------
B.PE.07.502_1047_wg5.JF320226    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGTA----- ---------- ---------- ----------
B.PE.07.502_1399_wg4.JF320013    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.PE.07.502_2254_FL6.JF320018    AG-TGGCGCC CGAACAGG-- ---------- ---------- --GACCTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.PE.07.502_2349_wg2.JF320028    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TTGAA----- ---------- ---------- ----------
B.PE.07.502_2622_wg1.JF320189    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG CAGAA----- ---------- ---------- ----------
B.PE.07.502_2649_wg8.JF320019    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.RU.04.04RU128005.AY682547      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACYYGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.RU.04.04RU129005.AY751406      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG GGAAA----- ---------- ---------- ----------
B.RU.04.04RU139089.AY751407      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.RU.04.04RU139095.AY819715      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.RU.09.09RU4457.JX500709        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AAAGTGAAAG TAAAA----- ---------- ---------- ----------
B.RU.10.10RU6629.JX500707        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.RU.11.11RU21n.JX500708         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAAA------ ---------- ---------- ----------
B.TH.06.AA010a_WG3.JX446800      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAA------ ---------- ---------- ----------
B.TH.07.AA040a_WG11.JX447156     ---------- ---------- ---------- ---------- --GGACTTGA AA-ACGAAAG TGAGA----- ---------- ---------- ----------
B.TH.90.BK132.AY173951           ---------- ---------- ---------- ---------- --GGACCTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.TW.94.TWCYS_LM49.AF086817      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCAAAAG AGAAA----- ---------- ---------- ----------
B.US.00.14294_1.DQ853436         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.US.00.ES1_20.EF363123          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.00.RHPA_TF1.JN944917        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.00.THRO_TF1.JN944930        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTAA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
B.US.00.WITO_flB3.FJ496169       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GTGAAAG TGAAT----- ---------- ---------- ----------
B.US.01.REJO_TF1.JN944911        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG GGAAA----- ---------- ---------- ----------
B.US.01.TRJO_flA1.FJ496151       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.02.04013226_2_flH11.FJ496   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.04.ES10_53.EF363127         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACACGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.04.ES4_24.EF363124          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG GGAAA----- ---------- ---------- ----------
B.US.04.ES8_43.EF363126          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCGGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.04.SAMI_WGA1.EU547186       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.US.04.USPI71101EI7y04051pcWG   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.04.USPI83747EI6y04121pcWG   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.05.04013396_0_flE6.FJ4960   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.05.502_0223_wg1.JF320059    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCGGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.05.502_1400_FL02.JF320043   AG-TGGCGCC TGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.05.502_2136_FL02.JF320185   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.05.502_2495_wg02.JF320054   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.05.CR0307R.FJ469717         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.US.05.USPI38417EI33y05051pcW   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AAAGCGAAAG TAAAA----- ---------- ---------- ----------
B.US.05.USPI88403EI14y05121pcW   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.06.502_0341_FL05.JF320003   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.06.502_1046_FL04.JF320564   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
B.US.06.502_1174_FL09.JF320053   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.US.06.502_1211_FL01.JF320151   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.US.06.502_1619_FL06.JF320126   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GTGAAAG AAGAA----- ---------- ---------- ----------
B.US.06.502_2667_FL03.JF320145   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.06.CH106_TF1.JN944897       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
B.US.06.CH40E_flA1.FJ495818      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG AGAAA----- ---------- ---------- ----------
B.US.06.CH58E_flA9.FJ495941      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.US.06.CH77E_flA1.FJ496000      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.07.502_0364_wg2.JF320563    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.07.502_1115_wg1.JF320045    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG AGAAA----- ---------- ---------- ----------
B.US.07.502_1478_wg4.JF320150    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
B.US.07.502_1897_wg6.JF320182    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCGGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
B.US.07.502_2289_05.JF320197     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.US.09.C1P.GU733713             AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.US.09.DEMB09US002.JX140657     -----GCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.10.CP1.JN397365             AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.10.VC1.JN397364             AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCGGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.11.ES38.JN397362            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTTA AAAGCGAAAG AGAAA----- ---------- ---------- ----------
B.US.83.5018_83.AY835777         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.US.83.5157_83.AY835781         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.US.83.RF_HAT3.M17451           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.US.83.SF2_LAV2_ARV2.K02007     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.US.84.5019_84.AY835779         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.US.84.MNCG_MN.M17449           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AAAAA----- ---------- ---------- ----------
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B.US.84.NY5CG.M38431            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA GA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.US.84.SF33.AY352275           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TTAA------ ---------- ---------- ----------
B.US.85.5077_85.AY835769        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.US.85.Ba_L.AB221005           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.86.5084_86.AY835775        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.US.86.5096_86.AY835749        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG GGAAA----- ---------- ---------- ----------
B.US.86.5127_86.AY835774        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.US.86.AD87_ADA.AF004394       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAGAA----- ---------- ---------- ----------
B.US.86.JRFL_JR_FL.U63632       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
B.US.86.YU_2.M93258             AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG GAAAA----- ---------- ---------- ----------
B.US.87.5113_87.AY835758        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.US.87.BC_BCSG3.L02317         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCGGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.88.5160_88.AY835763        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.US.88.WR27.AF286365           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTAA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.US.89.P896_89_6.U39362        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCGGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.90.US1.AY173952            ---------- ---------- ---------- ---------- --GGACCTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.90.US2.AY173953            ---------- ---------- ---------- ---------- --GGACTTGA AA-GCGAAAG GGAAA----- ---------- ---------- ----------
B.US.90.US3.AY173954            ---------- ---------- ---------- ---------- --GGACTAGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
B.US.90.US4.AY173955            ---------- ---------- ---------- ---------- --GGACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.US.90.WCIPR.U69591            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG GGAAA----- ---------- ---------- ----------
B.US.90.WEAU160_GHOSH.U21135    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG GGAAA----- ---------- ---------- ----------
B.US.91.5048_91.AY835761        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.US.91.DH12_3.AF069140         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCGGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.91.SUMA_flC5.FJ496145      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCGGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.93.WCD32P0793.DQ487188     AG-TGGCGCC CGAACAGG-- ---------- ---------- --GACGCTGA AA-GTGAAAG AGAAA----- ---------- ---------- ----------
B.US.93.WCM32P0793.DQ487190     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG GAAAA----- ---------- ---------- ----------
B.US.94.5082_94.AY835773        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.US.95.5073_95.AY835768        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.US.95.USPI90770EI72y95091pcW  AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG AGAAA----- ---------- ---------- ----------
B.US.96.5155_96.AY835753        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAG----- ---------- ---------- ----------
B.US.96.USPI55751EI32y96071pcW  AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.US.97.ARES2.AB078005          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.98.15384_1.DQ853463        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
B.US.98.98USHVTN1925c1.AY56010  AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.98.98USHVTN3605c9.AY56010  AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.98.98USHVTN8229c6.AY56010  AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GTGAAAG TAGGTGAA-- ---------- ---------- ----------
B.US.98.98USHVTN941c1.AY560110  AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.US.98.WC3_0498_4.EF175212     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.US.x.CR0059T.FJ469694         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
B.US.x.sample_C_BID_D617.JX503  TT-------- -GAACAGG-- ---------- ---------- ---GACCTGA AA-GTGAAAG TAGA------ ---------- ---------- ----------
B.VE.10.DEMB10VE001.JX140659    ---------- ------GG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
B.ZA.03.03ZAPS045MB2.DQ396398   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG AGAAA----- ---------- ---------- ----------
C.BR.02.02BR2022.JN692434       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BR.04.04BR013.AY727522        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.BR.04.04BR021.AY727523        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BR.04.04BR038.AY727524        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.BR.04.04BR073.AY727525        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BR.07.DEMC07BR003.JX140663    --------CC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BR.92.BR025_d.U52953          ---------- ---------- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BR.98.98BR004.AF286228        ---------- ---------- ---------- ---------- ------TTGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW07621.AF443088      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW076820.AF443089     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGGAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW087421.AF443090     AG-TGGCGCC CGACCAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW147127.AF443091     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.BW.00.00BW16162.AF443092      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW1686.AF443093       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.BW.00.00BW17593.AF443094      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW17732.AF443095      ---TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.BW.00.00BW17835.AF443096      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW17956.AF443097      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.BW.00.00BW18113.AF443098      AG-TGGCGCC CGAACGGG-- ---------- ---------- ---GACCTGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW18595.AF443099      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GGCTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW18802.AF443100      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW192113.AF443101     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-AGCGAAA GTAAA----- ---------- ---------- -----A----
C.BW.00.00BW20361.AF443102      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.BW.00.00BW20636.AF443103      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AG-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW20872.AF443104      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW2127214.AF443105    AG-TGGCGCC CGACCAGG-- ---------- ---------- ---GACCCGA AA-ACGAAAG TGAGA----- ---------- ---------- ----------
C.BW.00.00BW22767.AF443107      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG CAAGA----- ---------- ---------- ----------
C.BW.00.00BW38193.AF443108      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GC-CTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW38428.AF443109      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCAAAAG TGAAG----- ---------- ---------- ----------
C.BW.00.00BW38713.AF443110      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW3876_9.AF443111     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW3886_8.AF443112     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW3891_6.AF443113     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW3970_2.AF443114     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.00.00BW5031_1.AF443115     G--TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.96.96BW01B03.AF110959      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.96.96BW0402.AF110962       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
C.BW.96.96BW0502.AF110967       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
C.BW.96.96BW06.AF290028         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.96.96BW1104.AF110969       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.BW.96.96BW1210.AF110972       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.BW.96.96BW15C02.AF110974      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGG----- ---------- ---------- ----------
C.BW.96.96BW16B01.AF110976      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.BW.96.96BW17.AF110980         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.BW.96.96BWM032.AF443075       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-ACGAAAG TGAGA----- ---------- ---------- ----------
C.BW.96.96BWMO1_5.AF443074      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.BW.98.98BWMC122.AF443076      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.98.98BWMC134.AF443077      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.BW.98.98BWMC14A3.AF443078     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AG-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.98.98BWMO1410.AF443079     AG-TGGCGCC CGAAAGGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.98.98BWMO18D5.AF443080     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.98.98BWMO36A5.AF443081     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.BW.98.98BWMO37D5.AF443082     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.99.99BW393212.AF443083     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAAGA----- ---------- ---------- ----------
C.BW.99.99BW46424.AF443084      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.BW.99.99BW4745.AF443085       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.BW.99.99BW47547.AF443086      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.BW.99.99BWMC168.AF443087      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGG----- ---------- ---------- ----------
C.DJ.x.DJ259.AB485643           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.DK.01.CTL_015.EF514713        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-RCGAAAG TAAGA----- ---------- ---------- ----------
C.ES.06.X1936.EU786673          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ES.07.X2118_2.EU884500        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ES.08.X2363_2.EU786681        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
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C.ET.86.ETH2220.U46016          AG-TGGCGCC CGAACAGG-- ---------- ---------- --GGACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.IL.98.98IS002.AF286233        ---------- ---------- ---------- ---------- --------GA AA-ACGAAAG TGAGA----- ---------- ---------- ----------
C.IN.03.D24.EF469243            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-ACGAAAG TGAGA----- ---------- ---------- ----------
C.IN.93.93IN101.AB023804        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.IN.93.93IN904.AF067157        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.IN.93.93IN999.AF067154        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GAGAAAG TAAGA----- ---------- ---------- ----------
C.IN.94.94IN11246.AF067159      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.IN.94.94IN476.AF286223        ---------- ---------- ---------- ---------- ------TTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.IN.95.95IN21068.AF067155      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.IN.98.98IN012.AF286231        ---------- ---------- ---------- ---------- ------CTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.IN.98.98IN022.AF286232        ---------- ---------- ---------- ---------- ------GCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.IN.99.01IN565_10.AY049708     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.IN.x.VB39.EF694033            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.IN.x.VB49.EF694036            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.MM.99.mIDU101_3.AB097871      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.TZ.98.98TZ013.AF286234        ---------- ---------- ---------- ---------- -------CGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.TZ.98.98TZ017.AF286235        ---------- ---------- ---------- ---------- ------TTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1069MB.AY838567         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.00.1119MB.AY463229         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.00.1134MB.AY463217         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1157M3M.AY585266        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1162MB.AY463224         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1165MB.AY463230         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1168MB.AY463231         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1170MB.AY463225         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1171MB.AY463232         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
C.ZA.00.1176MB.AY463218         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1178MB.AY463233         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1184MB.AY838566         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1189MB.AY838565         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1192M3M.AY463219        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1195MB.AY463220         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1197MB.AY463234         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAGG TAAGA----- ---------- ---------- ----------
C.ZA.00.1210MB.AY463221         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1214MB.AY463236         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1217MB.AY463226         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.1225MB.AY463227         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
C.ZA.00.1228MB.AY463222         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.00.J112MA.AY838568         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.00.J38MA.AY463228          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.01.01ZATM45.AY228557       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.01.2134MB.AY463237         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTAA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.01.J54Ma.AY463223          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.02.02ZAPS001MB1.DQ275648   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGG----- ---------- ---------- ----------
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C.ZA.02.02ZAPS013MB1.DQ351222   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
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C.ZA.02.02ZAPS015MB1.DQ369995   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS017MB1.DQ351224   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS020MB1.DQ275653   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
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C.ZA.03.03ZAPS024MB1.DQ396367   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS025MB1.DQ351226   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
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C.ZA.03.03ZAPS030MB1.DQ369996   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS032MB1.DQ445633   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
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C.ZA.03.03ZAPS042MB1.DQ369977   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS043MB1.DQ351227   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS044MB1.DQ396384   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
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C.ZA.03.03ZAPS048MB1.DQ396364   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.ZA.03.03ZAPS049MB1.DQ369986   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS050MB1.DQ369980   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS051MB1.DQ396385   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS052MB1.DQ369987   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS054MB2.DQ369988   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS055MB1.DQ396373   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS056MB1.DQ396374   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS057MB2.DQ369989   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS059MB2.DQ445634   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS063MB1.DQ396388   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS066MB2.DQ396375   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS067MB2.DQ396389   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TACGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS071MB1.DQ396376   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS073MB1.DQ275649   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
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C.ZA.03.03ZAPS079B1.DQ093592    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS081MB1.DQ351219   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS083MB1.DQ351229   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS086MB1.DQ275654   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS088MB1.DQ275651   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS089MB1.DQ351216   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.ZA.03.03ZAPS091MB1.DQ275645   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
C.ZA.03.03ZAPS094MB1.DQ396377   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS095MB1.DQ275652   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS097MB1.DQ351230   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTAA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS099MB1.DQ275655   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS103MB2.DQ275656   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS104MB1.DQ369990   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS105MB2.DQ445632   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS108MB1.DQ396378   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
C.ZA.03.03ZAPS112MB2.DQ396386   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.ZA.03.03ZAPS113MB2.DQ396365   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS116MB1.DQ445635   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS118MB1.DQ396368   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS122MB1.DQ396370   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGG----- ---------- ---------- ----------
C.ZA.03.03ZAPS123MB1.DQ396369   AG-TGGCACC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS124MB1.DQ369976   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS125MB1.DQ396390   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
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C.ZA.03.03ZAPS126MB1.DQ275657    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS128MB1.DQ275643    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS130MB1.DQ275658    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS131MB1.DQ396380    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS133MB1.DQ275646    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GGCTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS136MB1.DQ351231    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS140MB1.DQ369981    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS143MB1.DQ396391    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-ACGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS151MB1.DQ396392    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS152MB1.DQ396399    AG-TGGCGCC TGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZAPS155MB1.DQ396371    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK005B2.DQ011175     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK006B2.AY878056     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK010B2.DQ164104     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK011B2.AY901965     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK013B2.DQ275660     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK016MB2.DQ351233    AG-TGGCGCC CGAGCAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK019B2.AY878063     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACACGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK020B2.AY878064     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK026B2.DQ011165     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK034B1.AY878065     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGG----- ---------- ---------- ----------
C.ZA.03.03ZASK036B1.AY901966     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK039B2.AY878068     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK058B2.AY901967     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK061B1.AY901968     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.03.03ZASK062B1.DQ164113     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK066B1.AY901969     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK067B1.DQ275642     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK072B1.DQ093593     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAGG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK073B1.AY901970     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK076B1.AY901975     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.ZA.03.03ZASK078B1.AY901971     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK084B1.AY901981     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK092B1.AY878057     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK094B1.AY878070     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK097B1.AY878060     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK098B1.AY878061     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.03.03ZASK103B1.DQ164106     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK104B1.DQ396395     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK107B1.DQ056410     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK110B1.DQ056411     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.03.03ZASK111B1.DQ056404     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK113B1.DQ351237     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.ZA.03.03ZASK117B1.DQ056408     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK118B1.DQ011169     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK120B1.DQ011176     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK211B1.DQ093601     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
C.ZA.03.03ZASK212B.DQ978981      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK213B1.DQ093607     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK215M6W.DQ275661    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK223B1.DQ093597     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK224MB1.DQ275664    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGA AA-GCGAAAG CAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK226B1.DQ164108     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK232B1.DQ093589     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.03ZASK233B1.DQ351234     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.04ZAPS194MB1.DQ275650    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GTGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.03.04ZASK082B1.AY901972     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAGG TAAAA----- ---------- ---------- ----------
C.ZA.03.SK023B2.AY772690         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GGCTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.SK029B2.AY772691         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.SK033B2.AY772692         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.03.SK040B1.AY703908         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.03.SK041B1.AY772693         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.SK043B1.AY772700         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.SK065B1.AY772694         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.03.SK112B1.AY772695         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.03.SK116B1.AY772696         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS157MB1.DQ351232    AG-TGGTGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
C.ZA.04.04ZAPS160B1.DQ164107     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS161B1.DQ164115     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS165MB1.DQ351221    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS168B1.DQ164121     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS169MB1.DQ396393    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS172MB1.DQ275659    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS177MB1.DQ396394    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS188B1.DQ164122     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS189B1.DQ164117     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS190B1.DQ093602     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS195B1.DQ164118     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS197MB1.DQ369997    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS198MB1.DQ445637    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS199B1.DQ164110     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS202B1.DQ093598     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS205B1.DQ093599     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-TCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS206B1.DQ164126     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS208B1.DQ093603     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS214B1.DQ093588     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS216B1.DQ093600     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZAPS217B1.DQ164119     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK031B2.AY901974     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.ZA.04.04ZASK083B2.DQ093585     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK127B1.DQ011172     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK128B1.DQ093586     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK131B1.DQ056412     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGG----- ---------- ---------- ----------
C.ZA.04.04ZASK132B1.DQ093594     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK135B1.DQ011166     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK136B1.DQ011177     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.04.04ZASK139B1.AY878072     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK142B1.DQ011167     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTAGA AA-GCGAAAG TCAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK145B1.AY901976     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG CAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK146.AY772699       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCGGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
C.ZA.04.04ZASK147B1.DQ164114     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK148B1.DQ093590     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK150B1.DQ011178     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK151B1.AY901977     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK154B1.AY878071     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
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C.ZA.04.04ZASK155B1.AY901978     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK156B1.DQ011171     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK159B1.DQ011179     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK160B1.DQ011173     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
C.ZA.04.04ZASK161B1.DQ011170     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK163B1.AY901979     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK164B1.DQ056405     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK165B1.DQ396387     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.04.04ZASK167B1.DQ164127     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK168B1.AY878058     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
C.ZA.04.04ZASK169B1.DQ396381     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK170B1.DQ093595     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK171B1.DQ351217     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK172B1.DQ369998     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.04.04ZASK173B1.DQ093604     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK174B1.AY901980     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.ZA.04.04ZASK175B1.DQ164129     AG-TGGAAAA TCTCTAGCAG TGGCGCCCGA AC-------- AGGGACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK176B1.DQ056416     AG-TGGCGCC CGACCAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK178B1.DQ093587     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK180B1.AY878059     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK181B1.AY878062     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK182B1.AY878054     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK183B1.AY878055     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK184B1.DQ056418     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK185B1.DQ011174     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK190B1.DQ056409     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK191B1.DQ369993     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK192B1.DQ396382     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK193B1.DQ396396     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-ACGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK196B1.DQ056413     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK200B1.DQ396383     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK201B1.DQ396397     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK202B1.DQ011180     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK204B1.DQ056414     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK206B1.DQ056415     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
C.ZA.04.04ZASK208B1.DQ056406     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK217B1.DQ056417     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG CAAGA----- ---------- ---------- ----------
C.ZA.04.04ZASK234B1.DQ093605     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.04.CAP30_5w_F4.GQ999973     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.CAP61_8w_F3.GQ999975     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.SK133B1.AY772698         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.SK134B1.AY703909         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.SK140B1.AY901973         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-ACGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.04.SK143B1.AY703910         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.04.SK144B1.AY703911         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGG----- ---------- ---------- ----------
C.ZA.05.05ZAPSK240B1.DQ369991    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.05ZASK243B1.DQ396372     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.05ZASK244B1.DQ369992     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.05ZASK245B1.DQ369982     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.05ZASK246B1.DQ369983     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TCAGA----- ---------- ---------- ----------
C.ZA.05.05ZASK247B1.DQ369994     AG-TGGCGCC TGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.ZA.05.CAP174_4w.GQ999981       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.CAP206_8w_F1.GQ999982    AG-TGGCGCC CGAACAGA-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.CAP210_5w.GQ999983       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.CAP228_8w_F2.GQ999984    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.CAP229_7w.GQ999985       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.CAP239_5w_F1.GQ999991    AG-CGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.CAP244_8w_F1.GQ999986    AG-TGGAACC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.CAP248_9w.GQ999987       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.CAP255_8w_F1.GQ999988    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG CAAGA----- ---------- ---------- ----------
C.ZA.05.CAP256_6w.GQ999989       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.CAP257_7w_F1.GQ999990    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.CAP45_5w_F1.GQ999974     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.CAP63_5w_F4.GQ999976     AG-TGGCGCC CGAGCAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.05.CAP65_6w.GQ999977        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.ZA.05.CAP84_3w_F2.GQ999978     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.CAP85_5w_F1.GQ999979     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.CAP88_5w_F2.GQ999980     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.05.CAP8_3w_F2.GQ999972      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
C.ZA.07.DEMC07ZA011.JX140664     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.08.DEMC08ZA011.JX140666     -G-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.09.DEMC09ZA008.JX140667     -G-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.09.DEMC09ZA009.JX140668     -G-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.10.DEMC10ZA001.JX140669     ------CGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.90.pZAC_R3714.JN188292      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTAGA AA-ACGAAAG TAAGG----- ---------- ---------- ----------
C.ZA.97.97ZA003.AY118165         ---------- ---------- ---------- ---------- ------TTGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
C.ZA.97.97ZA009.AY118166         ---------- ---------- ---------- ---------- ------CCGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.97.97ZA012.AF286227         ---------- ---------- ---------- ---------- ------TTGA AA-ACGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.98.98ZA445.AY158533         ---------- ---------- ---------- ---------- -------TGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.98.98ZA502.AY158534         ---------- ---------- ---------- ---------- ------TTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.ZA.98.98ZA528.AY158535         ---------- ---------- ---------- ---------- -------AGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.98.TV001.AY162223           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGA AA-GTGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.98.TV002.AY162224           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
C.ZA.98.TV012.AY162225           AG-TGGCGCC TGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.99.99ZACM9.AF411967         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.99.99ZALT21.EU293446        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.99.99ZALT39.EU293447        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.99.99ZALT42.EU293448        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.ZA.99.99ZALT45.EU293449        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.99.99ZALT46.EU293450        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.99.99ZALT4.EU293444         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.99.99ZALT5.EU293445         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGGA----- ---------- ---------- ----------
C.ZA.99.99ZATM10.AY228556        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZA.99.ZASW7.AF411966           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZM.02.02ZM108.AB254141         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.ZM.02.02ZM110.AB254142         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TACGA----- ---------- ---------- ----------
C.ZM.02.02ZM114.AB254146         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
C.ZM.02.02ZM115.AB254148         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZM.02.02ZMBC.AB254149          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
C.ZM.02.02ZMDB.AB254153          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
C.ZM.02.02ZMJC.AB254155          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZM.03.ZM246F_flA10.FJ496186    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
C.ZM.03.ZM247F_flA1.FJ496195     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZM.03.ZM249M_flC1.FJ496209     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
C.ZM.89.ZAM18.AB485645           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
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C.ZM.96.96ZM651.AF286224          ---------- ---------- ---------- ---------- ------CTTG AA-GCGAAAG TAGA------ ---------- ---------- ----------
C.ZM.96.96ZM751.AF286225          ---------- ---------- ---------- ---------- ------CCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
C.ZM.x.HIV1084i.AY805330          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
D.CD.83.ELI.K03454                AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
D.CD.83.NDK.M27323                AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
D.CD.84.84ZR085.U88822            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
D.CD.85.Z2Z6_Z2_CDC_Z34.M22639    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
D.CM.10.DEMD10CM009.JX140670      -G-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
D.KE.93.MB2059.AF133821           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
D.KE.97.ML415_2.AY322189          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
D.KR.04.04KBH8.DQ054367           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
D.SN.90.SE365.AB485648            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
D.UG.05.p190049.JX236668          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAGAA----- ---------- ---------- ----------
D.UG.07.p191647.JX236670          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
D.UG.07.p191882.JX236673          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
D.UG.07.pSC191727.JX236679        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
D.UG.08.p191859.JX236672          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
D.UG.91.UG270.AB485650            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
D.UG.92.92UG001.AJ320484          AG-TGGCGCC CGAACAGGGA CC-------- ---------- ------TGAA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
D.UG.94.94UG114.U88824            ---------- ---------- ---------- ---------- ------CTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
D.ZA.84.R2.AY773338               ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
D.ZA.85.R286.AY773340             ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
D.ZA.86.R482.AY773341             ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
D.ZA.90.R1.EF633445               ---------- ---------- ---------- ---------- --------TC GG-CTGGAAC GAGTT----- ---------- ---------- ----------
F1.AR.02.ARE933.DQ189088          ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
F1.BE.93.VI850.AF077336           ---------- ---------- ---------- ---------- ---GGACTTG AA-GCGAAAG TAGAA----- ---------- ---------- ----------
F1.BR.02.02BR082.FJ771006         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAGAA----- ---------- ---------- ----------
F1.BR.02.02BR170.FJ771007         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
F1.BR.06.06BR564.FJ771008         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
F1.BR.06.06BR579.FJ771009         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
F1.BR.07.07BR844.FJ771010         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
F1.BR.89.BZ126.AY173957           AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
F1.BR.93.93BR020_1.AF005494       ---------- ---------- ---------- ---------- ------CTGA AA-GCGAAAG TAAA------ ---------- ---------- ----------
F1.DE.x.MVP_30846.EU446022        ---------- ---------- ---------- ---------- ---------- --AAGCGAAA GCGAA----- ---------- ---------- ----------
F1.ES.02.ES_X845_4.FJ670516       AG-TGGCGCC CGAACAGG-- ---------- ---------- --GACGCGAA AA-GCGAAAA TAAAA----- ---------- ---------- ----------
F1.ES.11.DEMF110ES001.JX140671    ---------- ---------- ---------- ---------- --GGACCTGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
F1.ES.x.P1146.DQ979023            ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
F1.ES.x.X1093_2.DQ979025          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
F1.ES.x.X1670.DQ979024            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACYYGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
F1.FI.93.FIN9363.AF075703         ---------- ---------- ---------- ---------- ---------A AA-GCGAAAG TAGAA----- ---------- ---------- ----------
F1.RO.96.BCI_R07.AB485658         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
F1.RU.08.D88_845.GQ290462         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
F2.CM.10.DEMF210CM007.JX140673    -G-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
G.BE.96.DRCBL.AF084936            AG-TGGCGCC CGAACAGGGA CTTGAAA-AC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.CM.01.A1786.FJ389367            AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.CM.04.178_15.FJ389363           AG-TGGCGCC CGAACAGGGA CTTGAAA-AC GAAAGTTAAC AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
G.CM.04.314_40.FJ389364           AG-TGGCGCC CGAACAGGGA CTTGAAA-AC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.CM.04.515_28.FJ389365           AG-TGGCGCC CGAACAGGGA CKYGAAA-GC GAAAGTTAAM AGGGACTCGA AA-GCGRAAG TT-------- ---------- ---------- ----------
G.CM.04.944_5.FJ389366            AG-TGGCGCC CGAACAGGGA CTTGAAA-AC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.CM.10.DEMG10CM008.JX140676      ---TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.CM.96.96CMABB55.AY772535        AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAA AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.CN.06.sh52.HM067749             AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TTAATATGGG AATGGAAAGG GAAAGTT--- ----------
G.CU.99.Cu74.AY586547             AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAC AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.CU.99.Cu85.AY586548             AG-TGGCGCC CGAACAGGGA CTTGAAA-GY GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.CU.99.Cu87.AY586549             AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.ES.05.P962.EU786670             AG-TGGCGCC CGAACAGGGA CTTGAAA-AC GAAAGTTAAC AGGGACTTGT AA-GCGGAAG TTAACGGGGA CTCGAAAGCG AAAGTT---- ----------
G.ES.08.P1981_2.FJ670530          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GAGAAAG TTAAAAGGGA CTCGAAAGCG AAAGTT---- ----------
G.ES.09.X2634_2.GU362882          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TTAACAGGGA CTCGAAAGCG AAAGTT---- ----------
G.ES.99.X138.AF450098             AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTMAC AGGSACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.GH.03.03GH175G.AB287004         AG-TGGCGCC CGAACAGGGA CCCGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.KE.93.HH8793_12_1.AF061641      AG-TGGCGCC CGAACAGGGA CTTGAAA-G- TGAAGTTAAC AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.NG.92.92NG083_JV10832.U88826    ---------- ---------- -ATGAAA-GC GAAAGTTAAT AGGGACTCGA AA-ACGAAAG TT-------- ---------- ---------- ----------
G.PT.x.PT2695.AY612637            AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAA CGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.PT.x.PT3037.FR846408            AG-TGGCGCC CGAACAGGGA CCTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.PT.x.PT3306.FR846409            AG-TGGCGCC CGAACAGGGA CTTGGAA-GC GAAAGTTAAC AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.PT.x.PT988.FR846410             AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
G.SE.93.SE6165_G6165.AF061642     AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAC AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
H.BE.93.VI991.AF190127            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG CAAAA----- ---------- ---------- ----------
H.BE.93.VI997.AF190128            ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
H.CF.90.056.AF005496              ---------- ---------- ---------- ---------- ------TTGA TA-GTGAAAG TAAAA----- ---------- ---------- ----------
H.GB.00.00GBAC4001.FJ711703       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
J.CM.04.04CMU11421.GU237072       AG-TGGCGCC CGAACAGG-- ---------- ---------- --GACCTTGA AA-GTGAAAG TAGGA----- ---------- ---------- ----------
J.SE.93.SE9280_7887.AF082394      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
J.SE.94.SE9173_7022.AF082395      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.CF.90.90CF11697.AF197340    AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.CF.90.90CF4071.AF197341     AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAC AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.CF.90.90CR402_CAR_E_4002    AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.CN.05.FJ051.DQ859178        AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
01_AE.CN.05.FJ053.DQ859179        AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.CN.05.Fj052.EF036528        AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.CN.05.Fj055.EF036527        AGGTGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.CN.05.Fj056.EF036529        AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.CN.05.Fj057.EF036530        AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.CN.05.Fj065.EF036534        AG--GGCGCC CGAACAGGGA CCTTGAA-GC GGAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.CN.05.Fj066.EF036535        AG-TGGCGCC CGAACAGGGA CTTGA---GT GACAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.CN.06.FJ054.DQ859180        AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
01_AE.CN.06.Fj062.EF036531        AG-TGGTGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.CN.06.Fj063.EF036532        AG-TGGCGCC CGAACAGGGA CTCGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.CN.06.Fj064.EF036533        AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.HK.04.HK001.DQ234790        ---------- CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.JP.x.DR0492.AB253423        AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.JP.x.DR2594.AB253660        AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.JP.x.DR6824.AB253426        AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.JP.x.JRC77AE.AB565504       AG-TGGCGCC CGAACAGGGA CTTGAAA-AC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.04.AA027a_wg4.JX44702    ---------- -------GGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.04.AA075a_WG7.JX44754    AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.04.BKM.DQ314732          AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.05.AA004a_wg4a.JX4467    AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.05.AA033a_wg6a.JX4470    AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.05.AA049a_WG13.JX4472    ---------- ---------- ---------- ---------- ---------- -----GAAAG TT-------- ---------- ---------- ----------
01_AE.TH.05.AA064a_WG2.JX44741    ---------- ---------- ---------- ---------- ---------- ----CGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.05.AA079a_WG4.JX44759    ---TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
01_AE.TH.05.AA101a_WG1.JX44793    -G-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.05.AA107a_wg4.JX44802    ---------- --------GA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.06.AA002a_WG1.JX44666    AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTAAA- AGGGACTCGA AA-GCGAAAG GTT------- ---------- ---------- ----------
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01_AE.TH.06.AA017a_wg1.JX44689   ---------C CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.06.AA034a_wg2.JX44708   AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGAAAG GTT------- ---------- ---------- ----------
01_AE.TH.06.AA038a_WG3.JX44713   ---------- -GAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.06.AA055a_WG4.JX44731   ---------- --------GA CTTGAAA-GT GAAAGATAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.06.AA056a_WG5.JX44731   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.06.AA059a_WG5.JX44735   AG-TGGCGCC CGAACAGGGA CCTGAAA-AC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.06.AA063a_WG37.JX4474   AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.06.AA068a_14.JX447465   ---------- ---------- -TTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.06.AA082a_WG9.JX44764   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.06.AA085a_wg2.JX44768   AG-TGGCGCC CGAACAGGGA CTTTAAA-GT GAAAGTAAA- AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
01_AE.TH.06.AA088a_wg14.JX4477   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.06.AA099a_WG9.JX44789   AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGCAAG TT-------- ---------- ---------- ----------
01_AE.TH.07.AA015a_WG4.JX44687   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.07.AA019a_WG11.JX4469   AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-ACGAAAG GTT------- ---------- ---------- ----------
01_AE.TH.07.AA028a_wg3.JX44702   AG-TGGCGCC CGAACAGGGA CTTGAAA-AC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.07.AA050a_WG7.JX44728   ---------- --------GA CTTGAAA-AC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.08.AA007a_WG10.JX4467   ---------- --------GA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
01_AE.TH.08.AA037a_WG6.JX44712   AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
01_AE.TH.08.AA060a_WG1.JX44735   ---------- ---------- ---------- ---------- --------GA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.08.AA067a_WG12.JX4474   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.08.AA108a_WG6.JX44802   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.09.AA090a_WG11.JX4477   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.09.AA111a_WG11.JX4480   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.90.CM240.U54771         AG-TGGCGCC CGAACAGGGC ACTCGAAAGC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.93.93TH253.U51189       AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.93.93TH9021.AF164485    AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGCTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.TH.95.95TNIH022.AB032740   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
01_AE.TH.95.95TNIH047.AB032741   AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
01_AE.TH.x.NP03.AB485654         AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01_AE.VN.97.97VNAG204.FJ185247   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNAG206.FJ185248   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNAG207.FJ185249   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNAG210.FJ185251   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNAG212.FJ185252   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNAG214.FJ185253   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNAG218.FJ185255   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNHCM314.FJ18524   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNHCM319.FJ18524   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNHCM343.FJ18524   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.97.97VNHCM345.FJ18524   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01_AE.VN.98.98VNND15.FJ185235    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
02_AG.CM.08.DE00208CM001.JX140   AG-TGGCGCC CGAACAGGGA CCGG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.CM.08.DE00208CM004.JX140   -------GCC CGAACAGGGA CTTG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.CM.99.pBD6_15.AY271690     AG-TGGCGCC CGAACAGGGA CTTG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.ES.06.P1261.EU786671       AG-TGGCGCC CGAACAGGGA CCGG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.ES.06.P1423.EU884501       AG-TGGCGCC CGAACAGGGA CCGG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.FR.91.DJ263.AF063223       ---------- -----GGACC GGAA------ ----GCTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.GH.03.03GH181AG.AB286855   AG-TGGCGCC CGAACAGGGA CCTG------ --AAATTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.GH.03.03GH182AG.AB286857   AG-TGGCGCC CGAACAGGGA CCGG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.GH.03.03GH189AG.AB286862   AG-TGGCGCC CGAACAGGGA CCGG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.GH.03.03GH197AG.AB286863   AG-TGGCGCC CGAACAGGGA CTTG------ --GAGTTAAT AGGGACTCGA AA-GCGAAAG TG-------- ---------- ---------- ----------
02_AG.GH.03.GHNJ185.AB231895     AG-TGGCGCC CGAACAGGGA CTTGAAG--- -----TTAAT AGGGACTCGG AAAGCGAAAG TACCAGAGAA GTCTCTCCGA CGCAGTCGAA AGCGAAAGTT
02_AG.GH.03.GHNJ188.AB231896     AG-TGGCGCC CGAACAGGGA CCGG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.GH.03.GHNJ196.AB231898     AG-TGGCGCC CGAACAGGGA CCGG------ --AAGTTAAT AGGGACGCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.GH.97.97GH_AG1.AB049811    AG-TGGCGCC CGAACAGGGA CTTG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.GH.x.I_2496.AB485633       AG-TGGCGCC CGAACAGGGA CCGG------ --AAATTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.GW.04.CC_0030.FJ694791     ---------- --AACAGGGA C--------C GGAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.GW.05.CC_0048.FJ694792     ---------- ---------- ---------- -GAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.KR.07.07MHI10.JQ316136     AG-TGGCGCC CGAACAGGGA CTTG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.LR.x.POC44951.AB485636     AG-TGGCGCC CGAACAGGGA CTTG------ --AAGATAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.NG.x.IBNG.L39106           AG-TGGCGCC CGAACAGGAC TTGAC----- ----GGTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02_AG.SE.94.SE7812.AF107770      AG-TGGCGCC CGAACAGGGA CTTG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
03_AB.BY.00.98BY10443.AF414006   AG-TGGCGCC CGAACAGGGA CTTGAAA--- ---------- ---------- ---GCGAAAG TT-------- ---------- ---------- ----------
03_AB.RU.98.RU98001_98RU001.AF   AG-TGGCGCC CGAACAGGGA CTTGAAA--- ---------- ---------- ---GCGAAAG TT-------- ---------- ---------- ----------
04_cpx.CY.94.94CY032_3.AF04933   ---------- -----TTGAA AGTG------ -AAAGTTAAT AG-GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
04_cpx.GR.00.DE00400GR002.JX14   ---------- -GAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
04_cpx.GR.91.GR11_97PVCH.AF119   AG-TGCCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
04_cpx.GR.97.GR84_97PVMY.AF119   AG-TGGCGCC CGAACAGGGA CCCGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
05_DF.BE.93.VI961.AF076998       ---------- ---------- ---------- ---------- -GGACCTTGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
05_DF.BE.x.VI1310.AF193253       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAGAA----- ---------- ---------- ----------
05_DF.ES.99.X492.AY227107        ---------- ---------- ---------- ---------- -GGGACTTGA AA-ACGAAAG TAAA------ ---------- ---------- ----------
06_cpx.AU.96.BFP90.AF064699      AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
06_cpx.CD.x.BCF_Dioum.AB485660   AG-TGGCGCC CGAACAGGGA CTTTAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
06_cpx.EE.01.EE0359.AY535659     AG-TGGCGCC TGAACAGGGA CCAGAAA-AC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
06_cpx.GH.03.03GH173_06.AB2868   AG-TGGCGCC CGAACAGGGA CCCGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
06_cpx.ML.95.95ML127.AJ288982    AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
06_cpx.ML.95.95ML84.AJ245481     AG-TGGCGCC CGAACAGGGA CCTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
06_cpx.SN.97.97SE1078.AJ288981   AG-TGGCGCC CGAACAGGGA CCCGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
07_BC.CN.05.XJDC6431_2.EF36837   ---------- ---------- ---------- ---------- ------GCGA AA-GCGAAAG TAGAG----- ---------- ---------- ----------
07_BC.CN.05.XJDC6441.EF368370    ---------- ---------- ---------- ---------- ------TTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
07_BC.CN.06.Sichuan_2006_SC006   ---------- ---------- ---------- ---------- ---------- ---------G TAAGA----- ---------- ---------- ----------
07_BC.CN.06.Sichuan_2006_SC008   ---------- ---------- ---------- ---------- ---------- ---------G TAAGA----- ---------- ---------- ----------
07_BC.CN.06.Xinjiang_2006_709.   ---------- ---------- ---------- ---------- ---------- -----GAAAG TAAAG----- ---------- ---------- ----------
07_BC.CN.07.CNGZD.JQ423923       ---------- ---------- ---------- ---------- ------TTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
07_BC.CN.98.98CN009.AF286230     ---------- ---------- ---------- ---------- ------TTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
08_BC.CN.00.p00CH_HH090_08_BC3   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGT-AAG A--------- ---------- ---------- ---------- ---------- ----------
08_BC.CN.00.p00CH_WS035_08_BC5   AG-TGGCGCC CGAACAGGGA CCTGAAA-GC GAAAGT-GAG A--------- ---------- ---------- ---------- ---------- ----------
08_BC.CN.01.p01CH_DL001_08_BC0   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGT-AAG A--------- ---------- ---------- ---------- ---------- ----------
08_BC.CN.06.nx2.HM067748         AG-TGGCGCC CGAACAGGGG CAAGAAA-AG GAAAAT-GAG A--------- ---------- ---------- ---------- ---------- ----------
08_BC.CN.98.98CN006.AF286229     ---------- ---------- -TTGAAA-GC GAAAGT-AAG A--------- ---------- ---------- ---------- ---------- ----------
09_cpx.CI.00.00IC_10092.AJ8665   AG-TGGCGCC CGAACAGGGA CTTTAAA-GT GAAAGT-AAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
10_CD.TZ.96.96TZ_BF061.AF28954   AG-TGGCGCC CGAACAGG-- ---------- ---------- --GTACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
10_CD.TZ.96.96TZ_BF071.AF28954   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
10_CD.TZ.96.96TZ_BF110.AF28955   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
11_cpx.CM.95.95CM_1816.AF49262   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
11_cpx.CM.96.96CM_4496.AF49262   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
11_cpx.CM.97.MP818.AJ291718      AG-GGGGGCC CCAACAGG-- ---------- ---------- ---GACCCGA AA-GCCAAAG TTAGA----- ---------- ---------- ----------
11_cpx.FR.99.MP1298.AJ291719     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
11_cpx.FR.99.MP1307.AJ291720     GAAGGGCGCC CGAACCTG-- ---------- ---------- --GGACCCCG AA-GCGAAAG TAAGA----- ---------- ---------- ----------
11_cpx.GR.x.GR17.AF179368        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
12_BF.AR.97.A32879.AF408629      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
12_BF.AR.97.A32989.AF408630      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
12_BF.AR.99.ARMA159.AF385936     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
12_BF.UY.99.URTR23.AF385934      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTAGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
12_BF.UY.99.URTR35.AF385935      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
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13_cpx.CM.02.02CM_A1394.DQ8453   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGG GA-AGCGAAA GTGAA----- ---------- ---------- ----------
13_cpx.CM.04.04CM_173_9.DQ8453   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG CTAA------ ---------- ---------- ----------
13_cpx.CM.04.04CM_632_28.DQ845   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AG-CGAAAGC GAAAG----- ---------- ---------- ----------
13_cpx.CM.96.96CM_1849.AF46097   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGTGA AA-GCGAAAG CTAA------ ---------- ---------- ----------
13_cpx.CM.96.96CM_4164.AF46097   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTAA AA-GCGAAAG CTAA------ ---------- ---------- ----------
14_BG.ES.00.X605.AF450096        AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAC AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
14_BG.ES.00.X623.AF450097        AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
14_BG.ES.05.X1870.FJ670522       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TTAACAGGGA CTCGAAAGCG AAAGTT---- ----------
14_BG.ES.05.X772_8.FJ670528      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TTAACAGGGA CTCGAAAGCG AAAGTT---- ----------
15_01B.TH.99.99TH_MU2079.AF516   AG-TGGCGCC CGAACAGGGA CCTTAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
15_01B.TH.99.99TH_R2399.AF5305   ---------- ---------A CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
16_A2D.KR.97.97KR004.AF286239    ---------- -----CGTAA AGTG------ -AAAG-TAAT AGGGACTTGA AA-GCGAAAG TT-------- ---------- ---------- ----------
18_cpx.CU.99.CU14.AY586541       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
18_cpx.CU.99.CU68.AY894993       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
18_cpx.CU.99.CU76.AY586540       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
19_cpx.CU.99.CU29.AY588971       AG-TGGCGCC CGAACAGGGA CTTAAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
20_BG.CU.03.CB134.DQ020274       AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAA AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
20_BG.CU.99.Cu103.AY586545       AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAA AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
20_BG.ES.99.R77.AY586544         AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAA AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
22_01A1.CM.02.02CAMLT72.EU7439   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
22_01A1.CM.06.LPH27MF.JN864049   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
22_01A1.CM.10.LB054.JN864059     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
23_BG.CU.03.CB118.AY900571       AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAA AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
23_BG.CU.03.CB347.AY900572       AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAA AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
24_BG.CU.03.CB378.AY900574       AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
24_BG.CU.03.CB471.AY900575       AG-TGGCGCC CGAACAGGGA CCTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
24_BG.ES.08.X2456_2.FJ670526     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TTAATAGGGA CTCGAAAGCG AAAGTT---- ----------
25_cpx.CM.06.06CM_BA_040.EU693   AG-TGGCGCC CGAACAGGGA CCTGAAA-AA GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
25_cpx.SA.03.J11233.EU697906     AG-TGGCGCC CGAACAGGGA CCCGAAA-GY GAAAGTTAAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
25_cpx.SA.03.J11451.EU697908     AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
26_AU.CD.02.02CD_KS069.FM87778   AG-GGGCGCC CAACCAGGGC CTTTAAA-GC GAAAGT-AAA AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
26_AU.CD.02.02CD_LBTB084.FM877   AG-TGGCGCT CGAACAGGGA CCTTAAA-GC GAAAGT-AAA AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
26_AU.CD.02.02CD_MBTB047.FM877   AG-TGGCGCC CGAACAGGGA CCTTAAA-GC GAAAGT-AAA AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
26_AU.CD.97.97CD_KTB119.FM8777   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAG-TAAA AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
27_cpx.FR.04.04CD_FR_KZS.AM851   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GA-CTTG AA-AGCGAAA GTAA------ ---------- ---------- ----------
28_BF.BR.05.0679SV.JF804812      ---------- ---------- ---------- -------AGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
28_BF.BR.99.BREPM12313.DQ08587   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
28_BF.BR.99.BREPM12609.DQ08587   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
28_BF.BR.99.BREPM12817.DQ08587   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACC-GA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
29_BF.BR.01.BREPM16704.DQ08587   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGT GA-GCGAAAG TTGAA----- ---------- ---------- ----------
29_BF.BR.05.0264RI.JF804807      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
29_BF.BR.99.BREPM11948.DQ08587   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
31_BC.BR.02.110PA.EF091932       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
31_BC.BR.04.04BR142.AY727527     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
32_06A1.EE.01.EE0369.AY535660    AG-TGGCGCC CGAACAGGGA CCTGAAA-AC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
32_06A1.EE.02.EST2002_1169.DQ1   AG-TGGCGCC CGAACAGGGA CTTGAAA-AC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
33_01B.ID.07.JKT189_C.AB547463   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
33_01B.MY.05.05MYKL007_1.DQ366   AG-TGGCGCC CGAACAGGGA CTTGAAA-AC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
33_01B.MY.05.05MYKL015_2.DQ366   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
33_01B.MY.05.05MYKL031_1.DQ366   AG-TGGCGCC CGAACAGGGA CCTTAAA-GA GGAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
33_01B.MY.05.05MYKL045_1.DQ366   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
38_BF1.UY.03.UY03_3389.FJ21378   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
38_BF1.UY.04.UY04_3987.FJ21378   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
38_BF1.UY.04.UY04_4022.FJ21378   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
38_BF1.UY.05.UY05_4752.FJ21378   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
39_BF.BR.03.03BRRJ103.EU735534   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
39_BF.BR.03.03BRRJ327.EU735536   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GTGAAAG TAGAA----- ---------- ---------- ----------
39_BF.BR.04.04BRRJ179.EU735535   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
40_BF.BR.04.04BRRJ115.EU735538   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
40_BF.BR.04.04BRSQ46.EU735540    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
40_BF.BR.05.05BRRJ055.EU735537   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
40_BF.BR.05.05BRRJ200.EU735539   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.03.luBF_01_03.EU17015   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACYTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.03.luBF_02_03.EU17015   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.03.luBF_04_03.EU17014   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.03.luBF_05_03.EU17015   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACYTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.03.luBF_06_03.EU17014   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.03.luBF_07_03.EU17014   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCYGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.03.luBF_08_03.EU17014   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.03.luBF_09_03.EU17015   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACYTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.04.luBF_11_04.EU17014   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACYTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.04.luBF_12_04.EU17014   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.05.luBF_13_05.EU17013   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACYTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.05.luBF_14_05.EU17014   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.05.luBF_16_05.EU17015   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.05.luBF_17_05.EU17014   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACYTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.06.luBF_18_06.EU17013   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
42_BF.LU.06.luBF_19_06.EU17013   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
43_02G.SA.03.J11223.EU697904     AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGRAAG TT-------- ---------- ---------- ----------
43_02G.SA.03.J11232.EU697905     AG-TGGCGCC CGAACAGGGA CTTGAAA-AG CGAAAGTAAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
43_02G.SA.03.J11243.EU697907     AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGCAAG TT-------- ---------- ---------- ----------
43_02G.SA.03.J11456.EU697909     AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
44_BF.CL.00.CH80.FJ358521        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
44_BF.CL.01.CH12.AY536235        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-RCGAAAG TAAAA----- ---------- ---------- ----------
45_cpx.CD.97.97CD_MBFE185.FN39   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
45_cpx.CD.97.97CD_MBS30.FN3928   AG-TGGCGCC CGAACAGGGA CTTGAAA-AC GAAAG----- ---------- ------TAAG TT-------- ---------- ---------- ----------
45_cpx.CM.97.97CM_MP814.FN3928   AG-TGGCGCC CGAACAGGGA CTTG------ ---------- ---------A AA-GCGAAAG TAAAAGTT-- ---------- ---------- ----------
45_cpx.FR.04.04FR_AUK.EU448295   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
45_cpx.GA.97.97GA_TB45.FN39287   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TA-------- ---------- ---------- ----------
46_BF.BR.01.01BR087.DQ358801     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
46_BF.BR.01.01BR125.DQ358802     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
46_BF.BR.07.07BR_FPS625.HM0264   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
46_BF.BR.07.07BR_FPS742.HM0264   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
46_BF.BR.07.07BR_FPS783.HM0264   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
46_BF.BR.07.07BR_FPS810.HM0264   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
46_BF.BR.07.07BR_FPS812.HM0264   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
47_BF.ES.08.P1942.GQ372987       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
47_BF.ES.08.X2457_2.FJ670529     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
49_cpx.GM.02.N18380.HQ385477     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
49_cpx.GM.03.N26677.HQ385479     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-ACGAAAG TAAGG----- ---------- ---------- ----------
49_cpx.GM.97.N28353.HQ385478     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGG----- ---------- ---------- ----------
52_01B.MY.03.03MYKL018_1.DQ366   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
53_01B.MY.04.04MYKL016_1.DQ366   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
53_01B.MY.10.10MYKJ067.JX39061   ---------- ---------- ---------- --------AT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
53_01B.MY.10.10MYKJ079.JX39061   ---------- ---------- ---------- --------AT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
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53_01B.MY.11.11FIR164.JX390610   ---------- ---------- ---------- ---------- -GGGACTCGA AA-GCGRAAG TT-------- ---------- ---------- ----------
54_01B.MY.07.07MYKLD49.EU03191   AG-TGGCGCC CGAACAGGGA CTTGAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
54_01B.MY.08.08MYKL044.JX39097   ---------- ---------- ---------- ---------- ----ACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
54_01B.MY.09.09MYSB023.JX39097   AG-TGGCGCC CGAACAGGGA CTTGAAA-AT AAAAGTTATT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
55_01B.CN.10.HNCS102056.JX5746   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
55_01B.CN.11.GDDG318.JX574662    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01A1.MM.99.mCSW105.AB097872      AG-TGGCGCC CGAACAGGGA CTCGAAA-GC GGAAGTTAAT AGGGACTCGA AA-GCG-AAG TT-------- ---------- ---------- ----------
01B.CN.08.08CYM047.JF340054      ---------A GTTATAGG-- ---------- ---------- ---GACTCG- ---AAAGCGA AAGTT----- ---------- ---------- ----------
01B.JP.x.pHIV_1_Y271B01AE64.AB   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTGGA AAAGCGAAAG TAAAA----- ---------- ---------- ----------
01B.MM.00.mIDU502.AB097865       AG-TGGCGCC CGAACAGG-- ---------- ---------- --GCACGCGA AA-ACGAAAG TGAGA----- ---------- ---------- ----------
01B.MM.99.mCSW104.AB097867       AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01B.MY.04.04MYKL019_1.DQ366665   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01B.MY.05.05MYKL043_1.DQ366666   AG-TGGCGCC CGAACAGGGA CTTGAAG-GC GGAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01B.MY.06.06MMYKLD46.EF495062    AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01B.MY.07.07MYKLD47.EU031913     ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01B.MY.07.07MYKLD48.EU031914     AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
01B.TH.05.05TH140456.JN631793    AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01B.TH.05.AA095a_WG21.JX447830   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01B.TH.06.AA020a_wg2.JX446927    ---------- --------GA CTTTAAA-GT GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01B.TH.06.AA025a_WG13.JX447000   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01B.TH.06.AA084a_WG10.JX447668   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAA AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
01B.TH.91.CM237.AY167123         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
01BC.MM.00.mCSW503.AB097866      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
01BC.MM.99.mIDU107.AB097868      AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGT-AAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
01F2G.CM.02.LT31.JN864056        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
0206.DZ.10.DEURF10DZ001.JX1406   ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
0206.GH.03.03GH195AG_06.AB2868   AG-TGGCGCC CGAACAGGGA CTTGGAG--- -----TTAAT AGGGACTCGA AA-GCGAAAG TG-------- ---------- ---------- ----------
0206.NE.00.NE36.AJ508597         AG-TGGCGCC CGAACAGGGA CTTG------ -AAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
0206.NE.00.NE95.AJ508596         AG-TGGCGCC CGAACAGGGA CTCGAAA-GC GAAAGTTAAT AGGGACTCGA AA-ACGAAAG TT-------- ---------- ---------- ----------
0206.NE.97.NE03.AJ508595         AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
0209.CI.01.01IC_17395.AJ866554   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
0209.CI.01.01IC_PCI118.AJ86655   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
0209.CI.01.01IC_PCI127.AJ86655   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
0209.CI.97.97IC_PCI3.AJ866555    AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
0222.CM.02.02CAMLT04.EU743964    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
0222.CM.08.BDSH129.JN864052      ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
0222.CM.10.LB045.JN864053        -------CCG CAACCTCA-- ---------- ---------- ---ACCTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02A1.ES.05.X230_10.FJ670515      AG-TGGCGCC CGAACAGGGA C--------C GGAAGTTAAT AGGGACTCGA AA-ACGAAAG TT-------- ---------- ---------- ----------
02A1.ES.07.ES_P1751.GQ372986     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02A1.GH.03.GHNJ176.AB231894      AG-TGGCGCC CGAACAGGGA CCGG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02A1.GH.97.97GH_AG2.AB052867     AG-TGGCGCC CGAACAGGGA CCGG------ --AAGTTAAT AGGGGCTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02A1.RU.10.10RU6637.JN230353     AG-TGGCGCC CGAACAGGGA CCGG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02B.ES.99.99SP_11339.DQ926899    AG-TGGCGCC CGAACAGGGA CCGGAAA-TA TAGG------ ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02B.FR.02.URF4.JN882652          ---------- ---------- ---------- ---------- -GGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02B.FR.09.URF2.JN882654          ---------- ---------- ---------- ---------- ---------- -------AAG AGAAA----- ---------- ---------- ----------
02BD.FR.08.URF1.JN882653         AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02BG.FR.10.URF5.JN882655         ---------- ---------- ---------- ------TAAA AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02C.BE.93.VI1035.AJ276595        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
02D.GH.03.GHNJ193.AB231897       AG-TGGCGCC CGAACAGGGA CTTG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02D.GH.03.p03GH194AG09.AB48004   AG-TGGCGCC CGAACAGGGA CTTG------ -AAAGT-AAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
02F2G.CM.02.LT66.JN864057        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
02GK.CI.01.01IC_PCI123.AJ86655   AG-TGGCGCC CGAACAGGGA CTCGAAA-GC GAAAGTTAAT AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
06A1.BJ.x.B76.AJ293865           ---------- ---------- ---------- --------AT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
1819.CU.99.CU64.AY894995         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGTT----- ---------- ---------- ----------
26C.CD.02.02CD_LBTB032.FM87777   AG-TGGCGCC CGAACAGGGA CCTGAAA-AG CGAAAGTAAA AGGGACTCGA AA-GCGGCGG TT-------- ---------- ---------- ----------
26C.CD.97.97CD_KFE267.FM877778   AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTAAA- AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
26C.CD.97.MBFE250.FM877783       AG-TGGCGCC CGAACAGGGA CTTGAAAAGC GAAAGT-AAA AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
A1B.EE.02.EST2002_394.DQ167216   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1BD.GB.03.34567.JN417238        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
A1C.CA.04.04CA7750.EU220698      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TGTT------ ---------- ---------- ----------
A1C.IN.01.1579A.DQ083238         AG-TGGCGCC CGAACAGGGA CTTGAAAAGC GAAAG-TAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1C.IN.95.95IN21301.AF067156     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
A1C.RW.92.92RW009_06.U88823      ---------- ---------- ---------- ---------- ------ATGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
A1CD.KE.02.ML2000.EU110093       ---------- ---------- ---------- ---------- ---------- ---------- -T-------- ---------- ---------- ----------
A1CD.SE.95.SE8603.AF075702       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1CDGKU.ZA.99.CM4_99ZACM4.AF41   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAAGA----- ---------- ---------- ----------
A1CG.KE.02.ML1979PCR.EU110096    ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1D.DK.96.FSA.DQ912822           ---------- ---------- ---------- ---------- --GGACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
A1D.DK.98.MA.DQ912823            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
A1D.GB.00.8179.JN417236          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
A1D.GB.03.33365.JN417239         AG-GGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
A1D.GB.10.11762.JN417241         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
A1D.GB.10.12792.JN417240         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
A1D.KE.02.ML1974.EU110090        ---------- ---------- ---------- ---------- ---------- ---------- -A-------- ---------- ---------- ----------
A1D.SE.93.SE6954.AF075701        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1D.TZ.96.TZBFL0011.AF442569     -G-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA GA-GCGAAAG TAGAA----- ---------- ---------- ----------
A1D.TZ.96.TZBFL0088.AF442570     -G-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1D.TZ.97.TZBFL0086.AF442566     -G-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
A1D.UG.07.p191947.JX236674       AG-TGGCGCC CGAACAGGGA CCTGAAA-GC GAAAGTTAAC AGGGACTCGA AA-ACGAAAG TT-------- ---------- ---------- ----------
A1D.UG.08.p191982.JX236675       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
A1D.UG.90.UG266.AY352657         -G-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1D.UG.92.UG035.AY352656         -G-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1DHK.NO.97.97NOGIL3.AJ237565    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TGGAA----- ---------- ---------- ----------
A1DK.CD.85.MAL_MALCG.X04415      AG-TGGCGCC CGAACAGGGA CTTTAAA-GT GAAAGT-AAC AGGGACTCGA AA-GCGGAAG TT-------- ---------- ---------- ----------
A1G.BE.94.VI1197.AJ276596        ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
A1G.NG.92.92NG003.U88825         ---------- -----TTGAA AGCG------ -AAAGTTAAC AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1G.RU.11.11RU6900.JX500697      AG-TGGCGCC CGAACAGGGA CCGG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1G.RU.11.11RU6939.JX500706      AG-TGGCGCC CGAACAGGGA CCTG------ --AAGTTAAT AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1GHU.GA.x.VI354.AF076474        ---------- -------GGA CTTGAAA-GC GAAAG-TAAC AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
A1GJ.BW.98.BW2117.AF192135       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
A2C.ZA.98.DU178_98ZADU178.AF41   AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
A2C.ZM.90.ZAM184.U86780          AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
A2CD.KE.01.ML1956.EU110089       ---------- ---------- ---------- ---------- ---------- ---------- -A-------- ---------- ---------- ----------
AC.KE.97.ML672.AY322191          AG-TGGCGCC CGAACAGGGA CTTGAAA-GC GAAAGTAAAT AGGGACTTGA AA-GCGAAAG TT-------- ---------- ---------- ----------
AC.ZA.04.04ZAPS204B1.DQ093606    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
AGKU.CD.x.BCF_Kita.AB485665      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTCGA AA-GCGAAAG TAA------- ---------- ---------- ----------
BC.AR.01.ARE195FL.AY968312       ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
BC.BR.01.01_BR_RGS45.GQ365651    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
BC.BR.01.01_BR_RGS69.GQ365652    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAAGA----- ---------- ---------- ----------
BC.BR.02.02_BR_HF31.GQ365649     AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
BC.BR.05.05_BR_NSP24.GQ365650    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
BC.BR.06.06BR1114.JN692478       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
BC.BR.92.92BR023.HM100716        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
BC.CN.07.309.HM776938            AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
BC.CN.07.341.HM776939            -G-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
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BC.IN.02.INDNARI_0218440.EU000   AG-TGGCGCC CGAACAGG-- ---------- ---------- --GACCCGAA AA-GTGAAAG TAAGA----- ---------- ---------- ----------
BC.IN.02.NARI7_3.EU000511        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
BC.IN.02.NARI9_3.EU000508        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
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BC.MM.99.mIDU103.AB097873        AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
BCU.FR.06.06FR_CRN.EU448296      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAAGA----- ---------- ---------- ----------
BF.BR.03.BREPM1026.EF637055      ---------- ---------- ---------- ---------- -GGAACTTGA AA-GCGAAAG TGAGA----- ---------- ---------- ----------
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BF1.BR.00.BREPM13853.DQ085875    AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-ACGAAAG TAAAA----- ---------- ---------- ----------
BF1.BR.01.01BR042.DQ358799       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
BF1.BR.01.01BR047.DQ358800       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
BF1.BR.01.01BR226.DQ358803       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
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BF1.BR.02.02BR006.DQ358807       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
BF1.BR.02.02BR033.DQ358811       ---------- ---------- ---------- ---------- -------TGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
BF1.BR.02.02BR034.DQ358812       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
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BF1.BR.03.03BR2018.JN692449      AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCAGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
BF1.BR.03.03BR2019.JN692448      ---------- ---------- ---------- ---------- ---------- ---------- ----T----- ---------- ---------- ----------
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BF1.ES.09.DEMBF09ES006.JX14066   ---------- ---ACAGG-- ---------- ---------- ---GACTTGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
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BF1.IT.02.30638.GU595148         ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ---------- ----------
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BF1.JP.x.DR0769.AB253430         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAGAA----- ---------- ---------- ----------
BFG.MO.05.MO108.GU207884         ---------C CGAACAGGGA CTTGAAA-GC GAAAGTTAAC AGGGACTCGA AA-GCGAAAG TT-------- ---------- ---------- ----------
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CU.JP.04.DR5782.AB286849         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACGCGA AA-GCGAAAG TAAAA----- ---------- ---------- ----------
DF1G.ES.04.X963_4.FJ670527       AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACTTGA AA-GTGAAAG TTAATAGGGA CTCGAAAGCG GAAGTT---- -----C----
DO.FR.08.RBF208.GQ351296         AG-TGGCGCC CGAACAGG-- ---------- ---------- ---GACCGGA AA-GAGAAAG TGAAA----- ---------- ---------- ----------
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                                                   ---SD/3WJ----                                                                                        ---------SD-----------
                                                                                                                                                                     ------SDa----
                                    ....|....| ....|....| ....|....| ....|....| ....|....| ....|....| ....|....| ....|....| ....|....| ....|....|
                                        205        215        225        235        245        255        265        275        285        295
B.FR.83.HXB2_LAI_IIIB_BRU.K034      CCAGAGGAG- --CTCTCTCG ACGCAGG--A CTCGGCTTGC TGAA------ GCGCGCACGG CAAGAGGCGA GGGG----CG GCGA---CTG --GTGAGTAC
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A1.RU.00.RU00051.EF545108           ---------- ---------- ---------- CTCGGCTTGC TGAG------ GTGC-CACAG CAAGAGGCGA GAG-----CG GCGA---CYG --GTGAGTAC
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B.BR.02.02BR1013.JN692432           CCAGAGGAG- --ATCTCTCG ACGCAGG--A CTCGGCTTGC TGAA------ GCGCGCACGG CAAGAGGCGA GGGG----CG GCGA---CTG --GTGAGTAC
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C.ZA.03.SK065B1.AY772694         CCAGAGGAG- --ATCTCTCG ACGCAGG--A CTCGGCTTGC TGAA------ GTGCACTCGG CAGGAGGCGA GGG-----GG GCGA---CTG --GTGAGTAC
C.ZA.03.SK112B1.AY772695         CCAGAGGAG- --ATCTCTCG ACGCAGG--A CTCGGCTTGC TGAA------ GTGCACACGG CAAGAGGCGA GAG-----CG GCGG---CTG --GTGAGTAC
C.ZA.03.SK116B1.AY772696         CCAGAGAAG- --ATCTCTCG ACGCAGG--A CTCGGCTTGC TGAA------ GTGCACTCGG CAAGAGGCGA GAG-----CG GCGG---CTG --GTGAGTAC
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                                 SD-
                                 SDa-                                                                  -----U5/AUG-----
                                 ....|....| ....|....| ....|....| ....|....| ....|....| ....|....| ....|....| ....|....| ..
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B.US.04.USPI83747EI6y04121pcWG   GCCAAAT--- ---------- ---------- --TTTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.05.04013396_0_flE6.FJ4960   GCCATAT--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.05.502_0223_wg1.JF320059    GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.05.502_1400_FL02.JF320043   GCCG------ ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.05.502_2136_FL02.JF320185   GCCAAT---- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.05.502_2495_wg02.JF320054   GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.05.CR0307R.FJ469717         GCCAATG--- ---------- ---------- ----TTTTTT TCAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.05.USPI38417EI33y05051pcW   GCCAAAA--- ---------- ---------- ---ACTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGCGCGAGAG CG
B.US.05.USPI88403EI14y05121pcW   GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.06.502_0341_FL05.JF320003   GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.06.502_1046_FL04.JF320564   GCCATAT--- ---------- ---------- ----TTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.06.502_1174_FL09.JF320053   GCCAAAA--- ---------- ---------- -ATTTTGACT AGCGGAGGCT AGAAGG-AGA GAGAG--ATG GGTGCGAGAG CG
B.US.06.502_1211_FL01.JF320151   GCCATAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.06.502_1619_FL06.JF320126   GCCGAAA--- ---------- ---------- --AATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.06.502_2667_FL03.JF320145   GCCAAAA--- ---------- ---------- -AATTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.06.CH106_TF1.JN944897       GCCAAAA--- ---------- ---------- ---ACTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.06.CH40E_flA1.FJ495818      GCCAAAA--- ---------- ---------- ----TATTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.06.CH58E_flA9.FJ495941      GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.06.CH77E_flA1.FJ496000      GCCAATA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.07.502_0364_wg2.JF320563    GCCAAAA--- ---------- ---------- -AATTTTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.07.502_1115_wg1.JF320045    GCCGAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.07.502_1478_wg4.JF320150    GCCT------ ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.07.502_1897_wg6.JF320182    GCCAAAA--- ---------- --------AT AAAGTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.07.502_2289_05.JF320197     GCCAA----- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.09.C1P.GU733713             GCCAAA---- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.09.DEMB09US002.JX140657     GCCATAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.10.CP1.JN397365             GCCAAAG--- ---------- ---------- --ATTCTTTG ACTAACGGAG GCTAGA-AAG AGAGG--ATG GGTGCGAGAG CG
B.US.10.VC1.JN397364             GCCAAAA--- ---------- ---------- -GTTTTTTGA CTAGCGAAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.11.ES38.JN397362            GCCA------ ---------- ---------- --TATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.83.5018_83.AY835777         GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.83.5157_83.AY835781         GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.83.RF_HAT3.M17451           GCCGAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.83.SF2_LAV2_ARV2.K02007     GCCAA----- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAGAGATG GGTGCGAGAG CG
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B.US.84.5019_84.AY835779          GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.84.MNCG_MN.M17449            GCCAAAA--- ---------- ---------- --ATTCTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.84.NY5CG.M38431              GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.84.SF33.AY352275             GCCAAAA--- ---------- ---------- --AATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.85.5077_85.AY835769          GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.85.Ba_L.AB221005             GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.86.5084_86.AY835775          GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.86.5096_86.AY835749          GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.86.5127_86.AY835774          GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.86.AD87_ADA.AF004394         GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.86.JRFL_JR_FL.U63632         GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.86.YU_2.M93258               GCCAAAA--- ---------- ---------- -AATTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.87.5113_87.AY835758          GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.87.BC_BCSG3.L02317           GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.88.5160_88.AY835763          GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.88.WR27.AF286365             GCCATAA--- ---------- ---------- ----TTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.89.P896_89_6.U39362          GCCAAAT--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.90.US1.AY173952              GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.90.US2.AY173953              GCCAAAA--- ---------- ---------- --ATTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.90.US3.AY173954              GCCAAAA--- ---------- ---------- ----TTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.90.US4.AY173955              GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.90.WCIPR.U69591              GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.90.WEAU160_GHOSH.U21135      GCCGAAA--- ---------- ---------- ----TTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.91.5048_91.AY835761          GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.91.DH12_3.AF069140           GCCAAAA--- ---------- ---------- ----TTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.91.SUMA_flC5.FJ496145        GCCGAAA--- ---------- ---------- --ATTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.93.WCD32P0793.DQ487188       GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.93.WCM32P0793.DQ487190       GCCAAAA--- ---------- ---------- ----TTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.94.5082_94.AY835773          GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.95.5073_95.AY835768          GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.95.USPI90770EI72y95091pcW    GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.96.5155_96.AY835753          GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.96.USPI55751EI32y96071pcW    GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.97.ARES2.AB078005            GCCAAAC--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.98.15384_1.DQ853463          GCCATAA--- ---------- ---------- ----TTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.98.98USHVTN1925c1.AY56010    GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.98.98USHVTN3605c9.AY56010    GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.98.98USHVTN8229c6.AY56010    GCCAAAA--- ---------- ---------- -ATTTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.98.98USHVTN941c1.AY560110    GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.98.WC3_0498_4.EF175212       GCTGAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.x.CR0059T.FJ469694           GCCAAAA--- ---------- ---------- -CTTTTTTGA CTGGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.US.x.sample_C_BID_D617.JX503    GCCA------ ---------- ---------- --TTTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.VE.10.DEMB10VE001.JX140659      GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
B.ZA.03.03ZAPS045MB2.DQ396398     GCCAAA---- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BR.02.02BR2022.JN692434         GCCA------ ---------- ---------T TTTCATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BR.04.04BR013.AY727522          GCCA------ ---------- ---------T TTTATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BR.04.04BR021.AY727523          GCCA------ ---------- ---------T TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BR.04.04BR038.AY727524          GCCA------ ---------- ---------T TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BR.04.04BR073.AY727525          GCCA------ ---------- ---------T TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BR.07.DEMC07BR003.JX140663      GCCAATT--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BR.92.BR025_d.U52953            GCCA------ ---------- ---------T TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BR.98.98BR004.AF286228          GCCA------ ---------- ---------T TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW07621.AF443088        GCCAAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW076820.AF443089       GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW087421.AF443090       GCCA------ ---------- --------AT TTTATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW147127.AF443091       GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW16162.AF443092        GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW1686.AF443093         GCCAAAC--- ---------- ---------- TTTAATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW17593.AF443094        GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW17732.AF443095        GCCAAAT--- ---------- ---------- TTTATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGCG CG
C.BW.00.00BW17835.AF443096        GCCAAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW17956.AF443097        GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW18113.AF443098        GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW18595.AF443099        GCCAAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW18802.AF443100        GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW192113.AF443101       GCCAAAT--- ---------- ---------- TTTATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW20361.AF443102        GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW20636.AF443103        GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW20872.AF443104        GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW2127214.AF443105      GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW22767.AF443107        GCCAAAT--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW38193.AF443108        GCCGATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW38428.AF443109        GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW38713.AF443110        GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW3876_9.AF443111       GCCAAAA--- ---------- --------AA ATTTTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW3886_8.AF443112       GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW3891_6.AF443113       GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW3970_2.AF443114       GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.00.00BW5031_1.AF443115       GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.96.96BW01B03.AF110959        GCCAAAT--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAAAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.96.96BW0402.AF110962         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.96.96BW0502.AF110967         GCCA------ ---------- ---------T TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.96.96BW06.AF290028           GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.96.96BW1104.AF110969         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.96.96BW1210.AF110972         GCCA------ ---------- --------AT TTTATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.96.96BW15C02.AF110974        GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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C.BW.96.96BW17.AF110980           GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.96.96BWM032.AF443075         GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.96.96BWMO1_5.AF443074        GCCA------ ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.98.98BWMC122.AF443076        GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.98.98BWMC134.AF443077        GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.98.98BWMC14A3.AF443078       GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.98.98BWMO1410.AF443079       GCCAATT--- ---------- ---------T TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.98.98BWMO18D5.AF443080       GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.98.98BWMO36A5.AF443081       GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGTGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.98.98BWMO37D5.AF443082       GCCAAAA--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.99.99BW393212.AF443083       GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.99.99BW46424.AF443084        GCCAAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.99.99BW4745.AF443085         GCCTATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.99.99BW47547.AF443086        GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.BW.99.99BWMC168.AF443087        GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.DJ.x.DJ259.AB485643             GCCATAT--- ---------- ---------- -TTTATTTGA CTAGCAAGGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.DK.01.CTL_015.EF514713          GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ES.06.X1936.EU786673            GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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C.ES.07.X2118_2.EU884500        GCCAAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ES.08.X2363_2.EU786681        ---------- ---------- ---CGCCAAA TTTTAATTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ET.86.ETH2220.U46016          GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.IL.98.98IS002.AF286233        GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.IN.03.D24.EF469243            GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.IN.93.93IN101.AB023804        GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.IN.93.93IN904.AF067157        GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.IN.93.93IN999.AF067154        GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.IN.94.94IN11246.AF067159      GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.IN.94.94IN476.AF286223        GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.IN.95.95IN21068.AF067155      GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.IN.98.98IN012.AF286231        GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.IN.98.98IN022.AF286232        GCCGATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.IN.99.01IN565_10.AY049708     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.IN.x.VB39.EF694033            GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.IN.x.VB49.EF694036            GCCTAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGCG CG
C.MM.99.mIDU101_3.AB097871      GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.TZ.98.98TZ013.AF286234        GCCAATT--- ---------- ---------- --TTATTTGA CTGGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.TZ.98.98TZ017.AF286235        GCCA------ ---------- --------TT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1069MB.AY838567         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1119MB.AY463229         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1134MB.AY463217         GCCGAAT--- ---------- ---------- -TTTACTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1157M3M.AY585266        GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1162MB.AY463224         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1165MB.AY463230         GCCA------ ---------- --------AT TTTATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1168MB.AY463231         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1170MB.AY463225         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--GTG GGTGCGAGAG CG
C.ZA.00.1171MB.AY463232         GCCAAAT--- ---------- ---------- TTTATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1176MB.AY463218         GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGTGG CTAGAA-GGA GAGAGAGATG GGTGCGAGAG CG
C.ZA.00.1178MB.AY463233         GCCAAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1184MB.AY838566         GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1189MB.AY838565         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1192M3M.AY463219        GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-AGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1195MB.AY463220         GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1197MB.AY463234         CCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1210MB.AY463221         GCCGAAT--- ---------- ---------- -TTTATTGGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGTG CG
C.ZA.00.1214MB.AY463236         GCCAAAA--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1217MB.AY463226         GCCAAAT--- ---------- ---------- TTTATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1225MB.AY463227         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.1228MB.AY463222         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.J112MA.AY838568         GCCA------ ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.00.J38MA.AY463228          GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.01.01ZATM45.AY228557       GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.01.2134MB.AY463237         GCCA------ ---------- --------TT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.01.J54Ma.AY463223          GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.02.02ZAPS001MB1.DQ275648   GCCAAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.02.02ZAPS005MB1.DQ351235   GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.02.02ZAPS006MB1.DQ351220   GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.02.02ZAPS008MB1.DQ275647   GCCT------ ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGGG--ATG GGTGCGAGAG CG
C.ZA.02.02ZAPS013MB1.DQ351222   GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG TTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.02.02ZAPS014MB1.DQ351218   GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.02.02ZAPS015MB1.DQ369995   GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS017MB1.DQ351224   GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS020MB1.DQ275653   GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS021MB1.DQ369978   GCCGTTT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS023MB1.DQ351225   GCCTTTT--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS024MB1.DQ396367   GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS025MB1.DQ351226   GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS026MB1.DQ369985   GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGTG CG
C.ZA.03.03ZAPS027MB1.DQ351223   GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS030MB1.DQ369996   GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS032MB1.DQ445633   GCCGAAT--- ---------- ---------- TTTATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS034MB1.DQ369979   GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS042MB1.DQ369977   GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS043MB1.DQ351227   GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS044MB1.DQ396384   GCCGAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS046MB1.DQ369984   GCCGAAT--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS048MB1.DQ396364   GCCAAGT--- ---------- ---------- -TTATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS049MB1.DQ369986   GCCA------ ---------- --------AT TTTTATTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS050MB1.DQ369980   GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS051MB1.DQ396385   GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS052MB1.DQ369987   GCCATTT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS054MB2.DQ369988   GCCAAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS055MB1.DQ396373   GCCAAAT--- ---------- ---------- -TTTCTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS056MB1.DQ396374   GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS057MB2.DQ369989   GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS059MB2.DQ445634   GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS063MB1.DQ396388   GCCA------ ---------- --------AT TTTTATTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS066MB2.DQ396375   GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS067MB2.DQ396389   GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS071MB1.DQ396376   GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS073MB1.DQ275649   GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS074MB2.DQ351228   GCCAAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS077B1.DQ093591    GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGTG CG
C.ZA.03.03ZAPS079B1.DQ093592    GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGTGAGAG CG
C.ZA.03.03ZAPS081MB1.DQ351219   GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS083MB1.DQ351229   GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS086MB1.DQ275654   GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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C.ZA.03.03ZAPS089MB1.DQ351216   GCCAAAT--- ---------- ---------- --TTATTGGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS091MB1.DQ275645   GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS094MB1.DQ396377   GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS095MB1.DQ275652   GCCAAAT--- ---------- ---------- -TTCATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS097MB1.DQ351230   GCCGATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS099MB1.DQ275655   GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS103MB2.DQ275656   GCCAAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS104MB1.DQ369990   GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS105MB2.DQ445632   GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS108MB1.DQ396378   GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS112MB2.DQ396386   GCCA------ ---------- ---------- ATTTTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS113MB2.DQ396365   GCCAAAT--- ---------- ---------- TTTATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS116MB1.DQ445635   GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS118MB1.DQ396368   GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS122MB1.DQ396370   GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS123MB1.DQ396369   GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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C.ZA.03.03ZAPS124MB1.DQ369976    GCCA------ ---------- --------AT TTTTATTTGA CCAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS125MB1.DQ396390    GCCAAAT--- ---------- ---------- TTTATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS126MB1.DQ275657    GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS128MB1.DQ275643    GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGTGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS130MB1.DQ275658    GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS131MB1.DQ396380    GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS133MB1.DQ275646    GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS136MB1.DQ351231    GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS140MB1.DQ369981    GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS143MB1.DQ396391    GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS151MB1.DQ396392    GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS152MB1.DQ396399    GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZAPS155MB1.DQ396371    GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK005B2.DQ011175     GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK006B2.AY878056     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK010B2.DQ164104     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK011B2.AY901965     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK013B2.DQ275660     GCCG------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGTG CG
C.ZA.03.03ZASK016MB2.DQ351233    GCCAAAT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK019B2.AY878063     GCCA------ ---------- --------AT TTTATTTTGA CTAGCGGAGG CTAGAG-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK020B2.AY878064     GCCA------ ---------- --------TT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK026B2.DQ011165     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK034B1.AY878065     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK036B1.AY901966     GCCT------ ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK039B2.AY878068     GCCAATT--- ---------- ---------- --TTATTTGA CTAGTGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
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C.ZA.03.03ZASK061B1.AY901968     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK062B1.DQ164113     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK066B1.AY901969     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK067B1.DQ275642     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK072B1.DQ093593     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK073B1.AY901970     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK076B1.AY901975     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK078B1.AY901971     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK084B1.AY901981     GCCG------ ---------- --------AT TTTATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK092B1.AY878057     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK094B1.AY878070     GCCG------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK097B1.AY878060     GCCA------ ---------- --------AT TTTTATTTGA CCAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK098B1.AY878061     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK103B1.DQ164106     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK104B1.DQ396395     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK107B1.DQ056410     GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK110B1.DQ056411     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGTG CG
C.ZA.03.03ZASK111B1.DQ056404     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK113B1.DQ351237     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK117B1.DQ056408     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK118B1.DQ011169     GCCA------ ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK120B1.DQ011176     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGTG CG
C.ZA.03.03ZASK211B1.DQ093601     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK212B.DQ978981      GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK213B1.DQ093607     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK215M6W.DQ275661    GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK223B1.DQ093597     GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK224MB1.DQ275664    GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK226B1.DQ164108     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK232B1.DQ093589     GCCAAAC--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.03ZASK233B1.DQ351234     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.04ZAPS194MB1.DQ275650    GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.04ZASK082B1.AY901972     GCCT------ ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.SK023B2.AY772690         GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.SK029B2.AY772691         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.SK033B2.AY772692         GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.SK040B1.AY703908         GCCA------ ---------- ---------- TTTTATTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.SK041B1.AY772693         GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.SK043B1.AY772700         GCCAAAT--- ---------- ---------- -TTTACTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.SK065B1.AY772694         GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.SK112B1.AY772695         GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.03.SK116B1.AY772696         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS157MB1.DQ351232    GCCATTT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS160B1.DQ164107     GCCTAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS161B1.DQ164115     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS165MB1.DQ351221    GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS168B1.DQ164121     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS169MB1.DQ396393    GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS172MB1.DQ275659    GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS177MB1.DQ396394    GCCAAAA--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS188B1.DQ164122     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS189B1.DQ164117     GCCGAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS190B1.DQ093602     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS195B1.DQ164118     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS197MB1.DQ369997    GCCA------ ---------- --------AT TTTTAGTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS198MB1.DQ445637    GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS199B1.DQ164110     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS202B1.DQ093598     GCCAAAT--- ---------- ---------- TTTATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS205B1.DQ093599     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS206B1.DQ164126     GCCT------ ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS208B1.DQ093603     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS214B1.DQ093588     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS216B1.DQ093600     GCCGAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZAPS217B1.DQ164119     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK031B2.AY901974     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGTG CG
C.ZA.04.04ZASK083B2.DQ093585     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK127B1.DQ011172     GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--GTG GGTGCGAGAG CG
C.ZA.04.04ZASK128B1.DQ093586     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK131B1.DQ056412     GCCAATT--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGTG CG
C.ZA.04.04ZASK132B1.DQ093594     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK135B1.DQ011166     GCCA------ ---------- --------AT TTTATCTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK136B1.DQ011177     GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGCGCGAGAG CG
C.ZA.04.04ZASK139B1.AY878072     GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK142B1.DQ011167     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK145B1.AY901976     GCCATAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK146.AY772699       GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK147B1.DQ164114     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK148B1.DQ093590     GCCAACT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK150B1.DQ011178     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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C.ZA.04.04ZASK151B1.AY901977     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGTG CG
C.ZA.04.04ZASK154B1.AY878071     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK155B1.AY901978     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK156B1.DQ011171     GCCAAAA--- ---------- ---------T TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK159B1.DQ011179     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK160B1.DQ011173     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK161B1.DQ011170     GCCAAAT--- ---------- ---------- -TTCATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK163B1.AY901979     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGGGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK164B1.DQ056405     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK165B1.DQ396387     GCCAAAT--- ---------- ---------- -TTTTCTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK167B1.DQ164127     GCCAAAT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-AGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK168B1.AY878058     GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK169B1.DQ396381     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK170B1.DQ093595     GCCAAAT--- ---------- ---------- TTTCTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK171B1.DQ351217     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGCG CG
C.ZA.04.04ZASK172B1.DQ369998     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK173B1.DQ093604     GCCA------ ---------- --------AT TTCTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK174B1.AY901980     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK175B1.DQ164129     GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK176B1.DQ056416     GCCGAAT--- ---------- ---------- --TTTATTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK178B1.DQ093587     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGCG CG
C.ZA.04.04ZASK180B1.AY878059     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK181B1.AY878062     GCCTAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK182B1.AY878054     GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK183B1.AY878055     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK184B1.DQ056418     GCCATTT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK185B1.DQ011174     GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK190B1.DQ056409     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ACG GGTGCGAGAG CG
C.ZA.04.04ZASK191B1.DQ369993     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK192B1.DQ396382     GCCA------ ---------- --------AT TTTATTTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK193B1.DQ396396     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK196B1.DQ056413     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK200B1.DQ396383     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK201B1.DQ396397     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK202B1.DQ011180     GCCTAAT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK204B1.DQ056414     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK206B1.DQ056415     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK208B1.DQ056406     GCCAAAT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK217B1.DQ056417     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.04ZASK234B1.DQ093605     GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.CAP30_5w_F4.GQ999973     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGGG-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.CAP61_8w_F3.GQ999975     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.SK133B1.AY772698         GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.SK134B1.AY703909         GCCATTT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.SK140B1.AY901973         GCCGAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.SK143B1.AY703910         GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.04.SK144B1.AY703911         GCCAAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.05ZAPSK240B1.DQ369991    GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.05ZASK243B1.DQ396372     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.05ZASK244B1.DQ369992     GCCTATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.05ZASK245B1.DQ369982     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.05ZASK246B1.DQ369983     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.05ZASK247B1.DQ369994     GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP174_4w.GQ999981       GCCAATT--- ---------- ---------- --TTCTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP206_8w_F1.GQ999982    GCCAAAT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GGGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP210_5w.GQ999983       GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP228_8w_F2.GQ999984    GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP229_7w.GQ999985       GCCAAAT--- ---------- ---------- -TTTATTTGA CTGGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAA CG
C.ZA.05.CAP239_5w_F1.GQ999991    GCCGATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP244_8w_F1.GQ999986    GCCATTT--- ---------- ---------- --TATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP248_9w.GQ999987       GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP255_8w_F1.GQ999988    GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP256_6w.GQ999989       ACCAATT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP257_7w_F1.GQ999990    GCCAAAA--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP45_5w_F1.GQ999974     GCCAAAT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP63_5w_F4.GQ999976     GCCAATT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP65_6w.GQ999977        GCCAAAT--- ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP84_3w_F2.GQ999978     GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGGGG CTAGAA-GGA GAGAG--ATG GGTGCGAGTG CG
C.ZA.05.CAP85_5w_F1.GQ999979     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP88_5w_F2.GQ999980     GCCGATT--- ---------- ---------- -TGTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.05.CAP8_3w_F2.GQ999972      GCCAATT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.07.DEMC07ZA011.JX140664     GCCAAAT--- ---------- ---------- --TTTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.08.DEMC08ZA011.JX140666     GCCTATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.09.DEMC09ZA008.JX140667     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.09.DEMC09ZA009.JX140668     GCCAATT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.10.DEMC10ZA001.JX140669     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.90.pZAC_R3714.JN188292      GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.97.97ZA003.AY118165         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.97.97ZA009.AY118166         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.97.97ZA012.AF286227         GCTG------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.98.98ZA445.AY158533         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.98.98ZA502.AY158534         GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.98.98ZA528.AY158535         GCCAATT--- ---------- ---------- -TTTATTTGA CTAGCGGTGG CTAGAA-GGA GAGAAG-ATG GGTGCGAGAG CG
C.ZA.98.TV001.AY162223           GCCAATT--- ---------- ---------- --TTACTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.98.TV002.AY162224           GCCAAAA--- ---------- ---------- TTTTTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.98.TV012.AY162225           GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.99.99ZACM9.AF411967         GCCGAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.99.99ZALT21.EU293446        GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.99.99ZALT39.EU293447        GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.99.99ZALT42.EU293448        GCCA------ ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.99.99ZALT45.EU293449        GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.99.99ZALT46.EU293450        GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.99.99ZALT4.EU293444         GCCAATT--- ---------- ---------- --TTATTAGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.99.99ZALT5.EU293445         GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.99.99ZATM10.AY228556        GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZA.99.ZASW7.AF411966           GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZM.02.02ZM108.AB254141         GCCA------ ---------- ---------- TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGTG--ATG GGTGCGAGAG CG
C.ZM.02.02ZM110.AB254142         GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGGGAG CG
C.ZM.02.02ZM114.AB254146         GCCCAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZM.02.02ZM115.AB254148         GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZM.02.02ZMBC.AB254149          GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZM.02.02ZMDB.AB254153          GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZM.02.02ZMJC.AB254155          GCCATTT--- ---------- ---------- --TATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZM.03.ZM246F_flA10.FJ496186    GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZM.03.ZM247F_flA1.FJ496195     GCCGAAT--- ---------- ---------- TTTTGTCTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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C.ZM.03.ZM249M_flC1.FJ496209     GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZM.89.ZAM18.AB485645           GCCA------ ---------- --------AT TTTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZM.96.96ZM651.AF286224         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZM.96.96ZM751.AF286225         GCCAATT--- ---------- ---------- --TTATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
C.ZM.x.HIV1084i.AY805330         GCCAAAT--- ---------- ---------- -TTTATTTGA CTAGCGGAGG CTAGGA-GGA GAGAG--ATG GGTGCGAGAG CG
D.CD.83.ELI.K03454               GCTAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
D.CD.83.NDK.M27323               GCTGAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
D.CD.84.84ZR085.U88822           GCTGAAAA-- ---------- ---------- --TTTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
D.CD.85.Z2Z6_Z2_CDC_Z34.M22639   GCTAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
D.CM.10.DEMD10CM009.JX140670     GCTAAAA--- ---------- ---------- --ATTTTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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D.KE.97.ML415_2.AY322189         GCTGAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
D.KR.04.04KBH8.DQ054367          GCTAAAAA-- ---------- ---------- --TTTTTTGA CTAGCAGAGG CTAGAA-GGG GAAAG--ATG GGTGCAAGAG CG
D.SN.90.SE365.AB485648           GCTGAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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D.UG.07.p191647.JX236670         GCAAAAA--- ---------- ---------- --TTTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
D.UG.07.p191882.JX236673         GCTGAAT--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
D.UG.07.pSC191727.JX236679       GCTGAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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D.UG.94.94UG114.U88824           GCTAAAAA-- ---------- ---------- --ATTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
D.ZA.84.R2.AY773338              GCTAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
D.ZA.85.R286.AY773340            GCTAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
D.ZA.86.R482.AY773341            GCTAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
D.ZA.90.R1.EF633445              GCTAAAA--- ---------- ---------- -TAATTTTGA CTAGCGGAGG CTAAAA-GGA GAGAG--ATG GGTGCGAGAG CG
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F1.BR.02.02BR170.FJ771007        GCCAA----- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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F1.BR.89.BZ126.AY173957          GCCAA----- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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01_AE.VN.97.97VNAG218.FJ185255   GCCAA----- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
01_AE.VN.97.97VNHCM314.FJ18524   GCCAA----- ---------- ---------- ---ATTTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
01_AE.VN.97.97VNHCM319.FJ18524   GCCAA----- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGGG--ATG GGTGCGAGAG CG
01_AE.VN.97.97VNHCM343.FJ18524   GCCAA----- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGCA-GGA GAGAG--ATG GGTGCGAGAG CG
01_AE.VN.97.97VNHCM345.FJ18524   GCCAA----- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
01_AE.VN.98.98VNND15.FJ185235    GCCAA----- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
02_AG.CM.08.DE00208CM001.JX140   GCCAA----- ---------- ---------- ---ATTTTGA CTAGTGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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02_AG.ES.06.P1261.EU786671       GCCAA----- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
02_AG.ES.06.P1423.EU884501       GCCAA----- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
02_AG.FR.91.DJ263.AF063223       GCCAA----- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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02_AG.GH.03.03GH182AG.AB286857   GCCA------ ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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12_BF.UY.99.URTR23.AF385934      GCCAACA--- ---------- ---------- -AATTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
12_BF.UY.99.URTR35.AF385935      GCCGAAA--- ---------- ---------- -AATTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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18_cpx.CU.99.CU68.AY894993       GCCGAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
18_cpx.CU.99.CU76.AY586540       GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
19_cpx.CU.99.CU29.AY588971       GCCA------ ---------- ---------- --TTTTTTAA CTAGCGGAGG CTAGAA-AGA GAGAG--ATG GGTGCGAGAG CG
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20_BG.CU.99.Cu103.AY586545       GCCAAAC--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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23_BG.CU.03.CB347.AY900572       GCCAAAC--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
24_BG.CU.03.CB378.AY900574       GCCAAAC--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
24_BG.CU.03.CB471.AY900575       GCCAAAC--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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43_02G.SA.03.J11232.EU697905     GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
43_02G.SA.03.J11243.EU697907     GCCA------ ---------- ---------- --TTTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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45_cpx.CD.97.97CD_MBS30.FN3928   GCCAAGA--- ---------- ---------- ---ATTTTGA CTAGCGAAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
45_cpx.CM.97.97CM_MP814.FN3928   GCCA------ ---------- ---------- --TTTTTTGA CTAGCGRAGG CTARAA-GGA RARAG--ATG GGGGCGAGAG CG
45_cpx.FR.04.04FR_AUK.EU448295   GCCA------ ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
45_cpx.GA.97.97GA_TB45.FN39287   GCCAAAG--- ---------- ---------- AATTTTTTGA CTAGCGGTGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
46_BF.BR.01.01BR087.DQ358801     GCCAA----- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
46_BF.BR.01.01BR125.DQ358802     GCCAA----- ---------- ---------- ---ATTTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
46_BF.BR.07.07BR_FPS625.HM0264   GCCAA----- ---------- ---------- ---ATTTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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46_BF.BR.07.07BR_FPS810.HM0264   GCCATAT--- ---------- ---------- ----TTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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53_01B.MY.10.10MYKJ079.JX39061   GCCAA----- ---------- ---------- ---ATTTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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54_01B.MY.07.07MYKLD49.EU03191   GCCAA----- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
54_01B.MY.08.08MYKL044.JX39097   GCCAA----- ---------- ---------- ---AATTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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01B.MY.07.07MYKLD48.EU031914     GCCAA----- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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0206.NE.97.NE03.AJ508595         GCCAT----- ---------- ---------- ---ATTTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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BF1.BR.04.04BR1067.JN692456      GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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BF1.IT.06.89072.GU595153         GCCAAAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
BF1.JP.04.DR6082.AB480298        GCCAA----- ---------- ---------- ---TTTTTGA CTAGTGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
BF1.JP.04.DR6190.AB480300        GCCAAT---- ---------- ---------- ---TTTTTGA CTAGCAGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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BFG.MO.05.MO108.GU207884         GCCAA----- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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BG.DE.01.9196_01.AY882421        GCCATAA--- ---------- ---------- ---ATTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
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CU.JP.04.DR5782.AB286849         GCCAAAA--- ---------- ---------- ---TTTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
DF1G.ES.04.X963_4.FJ670527       GCTGAAA--- ---------- ---------- ----TTTTGA CTAGCGGAGG CTAGAA-GGA GAGAG--ATG GGTGCGAGAG CG
DO.FR.08.RBF208.GQ351296         ---------W TTTGCTGGCG GTGGCCAGAC CTAGGGGAAG -GGCGAAGTC CCTAGG-GGA GGAAG--ATG GGTGCGAGTG CG
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Figure S1. Alignment of all HIV-1 group M sequences described in the Los Alamos HIV sequence database (Premade web alignment of complete sequences 2012; www.hiv.lanl.gov) and used in the phylogenetic analysis (Figure 2). The region encompassing the nucleotides involved in the SD, SDa, SD-3WJ and U5-AUG interactions (positions 554-804 in the proviral HXB2 reference sequence) is shown. Sequences with ambiguous nts, nt insertions and deletions and nt variations in the analyzed region that are observed only in a single virus isolate, were excluded from the phylogenetic analysis, as these may represent reverse transcription, PCR or sequencing errors.

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