

>sp|Q9BXJ3|C1QT4_HUMAN

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>sp|Q8WTP9|XAGE3_HUMAN

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>sp|Q6IMI6|ST1C3_HUMAN

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>sp|Q6IMN6|CAPR2_HUMAN

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>sp|Q9ULW0|TPX2_HUMAN

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>sp|Q8WWW8|GAB3_HUMAN

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>sp|P13761|2B17_HUMAN

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>sp|Q2QD12|RPEL1_HUMAN

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>sp|Q15172|2A5A_HUMAN

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>sp|Q9NXE8|CWC25_HUMAN

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>sp|Q9COD6|FHDC1_HUMAN

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>sp|Q9H0C2|ADT4_HUMAN

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>sp|Q16762|THTR_HUMAN

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>sp|E9PQR5|NP1B8_HUMAN

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>sp|Q5VY80|RET1L_HUMAN

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>sp|P10451|OSTP_HUMAN

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>sp|Q5VWJ9|SNX30_HUMAN

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>sp|Q7Z4B0|CRO20_HUMAN

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>sp|Q9HAA7|YGO46_HUMAN

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>sp|Q6P582|MZT2A_HUMAN

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>sp|Q3V5L5|MGT5B_HUMAN

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ENC SARVQH C I C S V A S D H S V G L L S L R E K K C I M L A S R H L F P I Q V I K W R P S D D Y L V V G C S D G S V Y V W Q M D T G A L D R C V M G I T A V E I L N A
C D E A V P A A V D S L S H P A V N L K Q A M T R R S L A A L K N M A H H K L Q T L A T N L L A S E A S D K G N L P K Y S H N S L M V Q A I K T N L T D P D I H V L F F D V E
A L I I Q L L T E E A S R P N T A L I S P E N L Q K A S G S S D K G G S F L T G K R A A V L F Q Q V K E T I K E N I K E H L L D D E E E D E E I M R Q R R E E S D P E Y R S S
K S K P L T L L E Y N L T M D A K L F M S C L H A W G L N E V L D E V C L D R L G M L K P H C T V S F G L L S R G G H M S L M L P G Y N Q P A C K L S H G K T E V G R K L P
A S E G V G K G T Y G V S R A V T T Q H L L S I I S L A N T L M S M T N A T F I G D H M K K G P T R P P R P S T P D L S K A R G S P P T S S N I V Q G Q I K Q V A A P V V S A
R S D A D H S G S D P P S A P A L H T C F L V N E G W S Q L A A M H C V M L P D L L G L D K F R P P L L E M L A R R W Q D R C L E V R E A A Q A L L L A E L R R I E Q A G R K
E A I D A W A P Y L P Q Y I D H V I S P G V T S E A A Q T I T T A P D A S G P E A K V Q E E E H D L V D D D I T T G C L S S V P Q M K K I S T S Y E E R R K Q A T A I V L L G
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S A V L M G L L E L C A D A E K Q L A N I T M G L P L S P A A D S A R S A R H A L S L I A T A R P P A F I T T I A K E V H R H T A L A A N T Q S Q Q N M H T T T L A R A K G E
I L R V I E I L I E K M P T D V V D L L V E V M D I I M Y C L E G S L V K K K G L Q E C F P A I C R F Y M V S Y Y E R N H R I A V G A R H G S V A L Y D I R T G K C Q T I H G
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D Y D E E L Y V A G N M V I W S K G S K S Q A L A V Y K A F T V D S P V Q Q A L W C D F I I S Q D K S E K A Y S S N E V E K C I C I L Q S S C I N M H S I E G K D Y I A S L P
F Q V A N V W P T K Y G L L F E R S A S S H E V P P G S P R E P L P T M F S M L H P L D E I T P L V C K S G S L F G S S R V Q Y V D H A M K I V F L N T D P S I V M T Y D A
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L C G Q K F L C F L V E S Q L Q L R C V K F Q E S N D K T Q L I F G S V T N I P A K D A A P V E K I D T M L V L E G S G N L V L Y T G V V R V G K V F I P G L P A P S L T M S
N T M P R P S T P L D G V S T P K P L S K L L G S L D E V V L L S P V P E L R D S S K L H D S L Y N E D C T F Q Q L G T Y I H S I R D P V H N R V T L E L S N G S M V R I T I
P E I A T S E L V Q T C L Q A I K F I L P K E I A V Q M L V K W Y N V H S A P G G S Y H S E W N L F V T C L M N M G Y N T D R L A W T R N F D F E G S L S P V I A P K K A
R P S E T G S D D D W E Y L L N S D Y H Q N V E S H L L N R S L C L S P S E A S Q M K D E D F S Q N L S L D S S T L L F T H I P A I F F V L H L V Y E E L K N T L M G E G I
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R L V L S I A L Y I L G D E S L V S D E S S Q Y L T R I T I A P Q K L Q V E Q E E N R F S F R H S T S V S S L A E R L V V W M T N V G F T L R D L E T L P F G I A L P I R D
A I Y H C R E Q P A S D W P E A V C L L I G R Q D L S K Q A C E G N L P K G K S V L S S D V P S G T E T E E E D D G M N D M N H E V M S L I W S E D L R V Q D V R R L L Q S A
H P V R V N V V Q Y P E L S D H E F I E E K E N R L L Q L C Q R T M A L P V G R G M F T L F S Y H P V P T E P L P I P K L N L T G R A P P R N T T V D L N S G N I D V P P N M
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>sp|Q9BVA6|FICD_HUMAN

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>sp|P09488|GSTM1_HUMAN

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>sp|Q03112|EVI1_HUMAN

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>sp|043502|RA51C_HUMAN

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>sp|Q6PI97|CK088_HUMAN

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>sp|Q9UGK3|STAP2_HUMAN

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>sp|Q8IWX7|UN45B_HUMAN

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>sp|A6NCL1|GEMC1_HUMAN

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>sp|P34130|NTF4_HUMAN

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>sp|Q9UL12|SARDH_HUMAN

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>sp|Q8WWQ2|HPSE2_HUMAN

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>sp|Q9H1C7|CYTM1_HUMAN

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>sp|POCG34|TB15A_HUMAN

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>sp|Q9C0B5|ZDHC5_HUMAN

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>sp|Q71RG6|YH006_HUMAN

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>sp|Q9NQT6|FSCN3_HUMAN

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>sp|P02766|TTHY_HUMAN

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>sp|O95081|AGFG2_HUMAN

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>sp|Q8IW92|GLBL2_HUMAN

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>sp|Q2WGJ6|KLH38_HUMAN

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>sp|Q8NGY1|O10Z1_HUMAN

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>sp|Q8IZ21|PHAR4_HUMAN

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>sp|P25963|IKBA_HUMAN

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>sp|Q96P16|RPR1A_HUMAN

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>sp|P61550|ENVT1_HUMAN

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>sp|P35227|PCGF2_HUMAN

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>sp|Q9H208|O10A2_HUMAN

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>sp|Q9UHD1|CHRD1_HUMAN

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>sp|Q14773|ICAM4_HUMAN

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>sp|Q6ZP80|TM182_HUMAN

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>sp|AOFGR9|ESYT3_HUMAN

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>sp|O76082|S22A5_HUMAN

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>sp|P07203|GPX1_HUMAN

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>sp|Q9BSD7|NTPCR_HUMAN

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>sp|O95007|OR6B1_HUMAN

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>sp|Q8IVF2|AHNK2_HUMAN

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>sp|Q6ZP29|LAAT1_HUMAN

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>sp|Q9BQE4|SELS_HUMAN

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>sp|PODMV9|HS71B_HUMAN

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>sp|A6NGB7|TM221_HUMAN

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>sp|Q96G61|NUD11_HUMAN

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>sp|Q8TD07|N2DL4_HUMAN

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>sp|Q9HB19|PKHA2_HUMAN

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>sp|Q6UWH4|F198B_HUMAN

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>sp|Q8N5G0|SMI20_HUMAN

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>sp|Q96S42|NODAL_HUMAN

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>sp|Q9NPA2|MMP25_HUMAN

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>sp|Q9NW82|WDR70_HUMAN

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>sp|Q9BY41|HDAC8_HUMAN

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>sp|Q9Y6D0|SELK_HUMAN

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>sp|Q7Z6K5|ARPIN_HUMAN

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>sp|Q6UWV2|MPZL3_HUMAN

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>sp | Q9BVL4 | SELO_HUMAN

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>sp|Q96RU3|FBNP1_HUMAN

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>sp|P01037|CYTN_HUMAN

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>sp|P27540|ARNT_HUMAN

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>sp|Q15053|K0040_HUMAN

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>sp|P51795|CLCN5_HUMAN

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>sp|Q5T7W0|ZN618_HUMAN

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>sp|Q92845|KIFA3_HUMAN

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>sp|A2A3N6|PIPSL_HUMAN

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>sp|P01160|ANF_HUMAN

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>sp|Q9P2S2|NRX2A_HUMAN

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S G G Q S S L F I H A G A P T V S S E N F I L P K G A A V N G P V S H S S L T K T S N M N K G S V S L T T G Q P V D Q P T T E S C S T L K V A A D L Q L S T P Q K A S Q H Q V
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R Y Y C C E E C N F M A V T E N E L E C H R G I A H G A V V K C P M V T S D I A Q R K T Q K K T F M K D S V V G S S K S A T Y I C K M C P F T T S A K S V L K K H T E Y L H
S S S C V D S F G S P L G L D K R K N D I L E E P V D S D S T K T L T K Q Q S T T F P K N S A L K Q D V K R T F G S T S Q S S S F S K I H K R P H R I Q K A R K S I A Q S G V
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T Q A I E D E T Y S D I N Q E H V N L F P L F K S K V E G Q E P G E N A T L S Y D Q N D G F Y F E Y Y E D T G S N N F L H E I H D P Q H L E T A D A S L S K H S S V F H W T D
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>sp|P56703|WNT3_HUMAN

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>sp|Q96M02|CJ090_HUMAN

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>sp|P12643|BMP2_HUMAN

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>sp|POCW23|AKAI1_HUMAN

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>sp|Q7Z3Z0|K1C25_HUMAN

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>sp|O14964|HGS_HUMAN

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>sp|Q9GZY8|MFF_HUMAN

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>sp|Q9GZW5|SCND2_HUMAN

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>sp|Q9UJK0|TSR3_HUMAN

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>sp|Q8NCM2|KCNH5_HUMAN

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>sp|A6NCF6|MA13P_HUMAN

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>sp|O43390|HNRPR_HUMAN

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>sp|Q9BVG9|PTSS2_HUMAN

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>sp|Q7L592|NDUF7_HUMAN

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>sp|A6NC98|CC88B_HUMAN

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>sp|Q9BSE5|SPEB_HUMAN

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>sp|P07947|YES_HUMAN

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>sp|Q9NVX0|HAUS2_HUMAN

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>sp|Q6IPR1|ETFR1_HUMAN

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>sp|P04004|VTNC_HUMAN

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>sp|P59990|KR121_HUMAN

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>sp|Q9BU23|LMF2_HUMAN

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>sp|Q8N584|TT39C_HUMAN

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>sp|Q92823|NRCAM_HUMAN

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>sp|075528|TADA3_HUMAN

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>sp|Q01831|XPC_HUMAN

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>sp|Q9Y6R1|S4A4_HUMAN

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>sp|Q8NDW4|ZN248_HUMAN

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>sp|Q13571|LAPM5_HUMAN

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>sp|O95182|NDUA7_HUMAN

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>sp|Q8NGQ3|ORIS2_HUMAN

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>sp|Q8IZ57|NRSN1_HUMAN

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>sp|Q8WTS1|ABHD5_HUMAN

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>sp|Q9P2Y5|UVRAG_HUMAN

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>sp|043610|SPY3_HUMAN

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>sp|P10253|LYAG_HUMAN

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>sp|POC6P0|BCL8_HUMAN

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>sp|Q15029|U5S1_HUMAN

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>sp|Q8NGG2|OR5T2_HUMAN

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>sp|Q53T59|H1BP3_HUMAN

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>sp|Q9HC44|GPBL1_HUMAN

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>sp|Q99719|SEPT5_HUMAN

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>sp|P49238|CX3C1_HUMAN

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>sp|Q5VUJ9|EFCB2_HUMAN

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G P L Q S G A S P T P S I S A S A S T L Q L S P P R T K N I T A K N P A K S N T S K P N T V K S N A S K P N T S K P N G S K S K Y K P K I S N M Q K K Q S T L A S S N K K S K
V N T A L R N L R Y R R G I H K C I E C C S E I K D F A N H F P T Y V H C S F C R Y N T S C S K A Y V N H M M S F H S N R P S K R F C I F K K H S E N L R G I T L V C L N C D
F L S D V S G L D N M A T H L S Q H K T H T C Q V V M Q K V S V C I P T S E H L S E L K K E A P A K E Q E P V S K E I A R P N M A E R E T E T S N S E S K Q D K A A S S K E K
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S S E A R F S S K N I K D L R L A S D N V S I D Q F L R K R H E P E S V S S D V S E Q G S I H L E P L T P S E V L E Y E A T E I L Q K G S G D P S A K T D E V V S D Q T D D I
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D H I F L L S L V D L N K N F K K I Y W P A A K E R V E L C K L A G K D A N T E C A N F I R V L Q P Y N K T H I Y V C G T G A F H P I C G Y I D L G V Y K E D I I F K L D T H
N L E S G R L K C P F D P Q Q P F A S V M T D E Y L Y S G T A S D F L G K D T A F T R S L G P T H D H H Y I R T D I S E H Y W L N G A K F I G T F F I P D T Y N P D D D K I Y
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S S I F K G S A V C V Y S M A D I R A V F N G P Y A H K E S A D H R W V Q Y D G R I P Y P R P G T C P S K T Y D P L I K S T R D F P D D V I S F I K R H S V M Y K S V Y P V A
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K G E R G E D S S E E K H L E E P G E T Q N A F L N E R K Q A S A I K K E E L V A R S E T H A A G H S Q E K T H S R E K S S Q E S G E E T G S Q E N H P Q E S K G Q P R S Q E
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>sp|P52907|CAZA1_HUMAN

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>sp|Q9NUJ3|T11L1_HUMAN

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>sp|Q9UBW7|ZMYM2_HUMAN

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>sp|Q96NU1|SAM11_HUMAN

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>sp|Q6ZQN5|FOX12_HUMAN

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>sp|Q8NEM7|SP20H_HUMAN

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>sp|Q8NBZO|IN80E_HUMAN

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>sp|Q9H1J7|WNT5B_HUMAN

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>sp|Q8NDF8|PAPD5_HUMAN

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>sp|Q5TKA1|LIN9_HUMAN

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>sp|Q9NQ89|CLO04_HUMAN

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>sp|P04180|LCAT_HUMAN

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>sp|Q8NH05|OR4Q3_HUMAN

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>sp|Q9P243|ZFAT_HUMAN

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>sp|P07919|QCR6_HUMAN

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>sp|O95136|S1PR2_HUMAN

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>sp|Q1HG43|DOXA1_HUMAN

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>sp|POCT01|SPDE6_HUMAN

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>sp|Q9NRW7|VPS45_HUMAN

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>sp|Q8WXU2|DYXC1_HUMAN

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>sp|O95677|EYA4_HUMAN

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>sp|Q9H4I2|ZHX3_HUMAN

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>sp|O15079|SNPH_HUMAN

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>sp|Q13882|PTK6_HUMAN

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>sp|O75439|MPPB_HUMAN

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>sp|Q96BD0|S04A1_HUMAN

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>sp|PODN77|OPSG2_HUMAN

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>sp|Q9H4Y5|GSTO2_HUMAN

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>sp|POCG31|Z286B_HUMAN

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>sp|Q9ULJ1|ODF2L_HUMAN

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>sp|Q14D04|MELT_HUMAN

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>sp|Q9BXT6|M10L1_HUMAN

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MQQ

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>sp|P01210|PENK_HUMAN

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>sp|Q9NWW9|HRSL2_HUMAN

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>sp|O75054|IGSF3_HUMAN

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>sp|Q96LT9|RBM40_HUMAN

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>sp|Q8NBK3|SUMF1_HUMAN

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>sp|Q9H244|P2Y12_HUMAN

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>sp|Q96LA8|ANM6_HUMAN

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>sp|Q9HC62|SENP2_HUMAN

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>sp|Q5H9T9|FSCB_HUMAN

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>sp|Q8N859|ZN713_HUMAN

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>sp|P42694|HELZ_HUMAN

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>sp|O14815|CAN9_HUMAN

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>sp|Q9Y252|RNF6_HUMAN

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>sp|Q8TBE9|NANP_HUMAN

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>sp|POC623|OR4Q2_HUMAN

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>sp|A5D6W6|FITM1_HUMAN

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>sp|Q9BZW7|TSG10_HUMAN

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>sp|Q5T8A7|PPR26_HUMAN

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>sp|Q9NZ32|ARP10_HUMAN

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>sp|P11021|GRP78_HUMAN

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>sp|P33763|S10A5_HUMAN

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>sp|Q8TAX0|OSR1_HUMAN

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>sp|Q9H5L6|THAP9_HUMAN

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>sp|POC7N5|OR8U9_HUMAN

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>sp|Q8IZP0|ABI1_HUMAN

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>sp|Q9UKI9|PO2F3_HUMAN

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>sp|P17213|BPI_HUMAN

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>sp|Q15286|RAB35_HUMAN

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>sp|P07998|RNAS1_HUMAN

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>sp|Q9UMN6|KMT2B_HUMAN

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>sp|Q9Y483|MTF2_HUMAN

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>sp|Q14563|SEM3A_HUMAN

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>sp|P00403|COX2_HUMAN

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>sp|O60516|4EBP3_HUMAN

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>sp|P13591|NCAM1_HUMAN

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>sp|Q8N1D0|BWR1B_HUMAN

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>sp|Q16718|NDUA5_HUMAN

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>sp|Q8NI51|CTCFL_HUMAN

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>sp|P00488|F13A_HUMAN

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>sp|Q9HCJ6|VAT1L_HUMAN

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>sp|Q8WZ04|TOMT_HUMAN

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>sp|Q96RD0|OR8B2_HUMAN

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>sp|Q9NX61|T161A_HUMAN

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>sp|Q15723|ELF2_HUMAN

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>sp|Q9NUL3|STAU2_HUMAN

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>sp|E9PKD4|NP1A5_HUMAN

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>sp|Q13685|AAMP_HUMAN

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>sp|Q9Y4P8|WIPI2_HUMAN

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>sp|Q9H6U6|BCAS3_HUMAN

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>sp|Q8WWP7|GIMA1_HUMAN

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>sp|Q9BZW4|TM6S2_HUMAN

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>sp|Q16572|VACHT_HUMAN

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>sp|Q9BXE9|VN1R3_HUMAN

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>sp|A6NDS4|TBC3B_HUMAN

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>sp|C9JI98|TM238_HUMAN

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>sp|Q7KZI7|MARK2_HUMAN

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>sp|Q9BQ83|SLX1_HUMAN

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>sp|O15067|PUR4_HUMAN

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>sp|Q9NYR9|KBRS2_HUMAN

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>sp|Q9BY10|TSCOT_HUMAN

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>sp|O60547|GMDS_HUMAN

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>sp|Q9NXN4|GDAP2_HUMAN

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>sp|Q15849|UT2_HUMAN

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>sp|Q96HP4|OXND1_HUMAN

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>sp|Q8TDV5|GP119_HUMAN

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>sp|094979|SC31A_HUMAN

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>sp|P09871|C1S_HUMAN

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>sp|Q969S8|HDA10_HUMAN

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>sp|Q14145|KEAP1_HUMAN

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>sp|P29460|IL12B_HUMAN

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>sp|Q86TB3|ALPK2_HUMAN

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>sp|Q9BQ15|SOSB1_HUMAN

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>sp|Q9POU3|SENP1_HUMAN

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>sp|Q6UXA7|CF015_HUMAN

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>sp|Q9Y237|PIN4_HUMAN

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>sp|Q6UXU4|GSG1L_HUMAN

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>sp|Q99550|MPP9_HUMAN

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>sp|Q5GH70|XKR9_HUMAN

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>sp|P51160|PDE6C_HUMAN

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>sp|Q9H707|ZN552_HUMAN

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>sp|Q5T751|LCE1C_HUMAN

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>sp|Q9HCX4|TRPC7_HUMAN

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>sp|Q96P67|GPR82_HUMAN

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>sp|Q96A26|F162A_HUMAN

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>sp|O15084|ANR28_HUMAN

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>sp|P25116|PAR1_HUMAN

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>sp|A6NHL2|TBAL3_HUMAN

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>sp|P30483|1B45_HUMAN

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>sp|Q8IZQ1|WDFY3_HUMAN

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>sp|Q9Y216|MTMR7_HUMAN

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>sp|A2RU48|SMC03_HUMAN

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>sp|Q8N8G2|VGLL2_HUMAN

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>sp|Q6ZRP5|YD019_HUMAN

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>sp|014836|TR13B_HUMAN

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>sp|Q6ZSI9|CAN12_HUMAN

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>sp|Q3KR37|GRM1B_HUMAN

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>sp|P28340|DPOD1_HUMAN

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>sp|P33527|MRP1_HUMAN

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>sp|Q8NH85|OR5R1_HUMAN

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>sp|Q8IY50|S35F3_HUMAN

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>sp|Q9H1E1|RNAS7_HUMAN

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TEETENİRİSSVİNSİREİKEEİGNİLKSSHİSGVLEİENSVDİLİSRMDİLEERİDİSLEDQİEEFSKDTMQMTKQİİSKERQRDİEER
SRSCNİRİLİGİPEKESYENRAEDİİKEİİDİNFaelkkGSİEİVSACRVPİSKİDEKRLTPRHİLVKFWNSSDKEKİİRASRERREİ
TYQGTİRİRLTADİSLDİTLDARSKWSNVFKVİLEKGFNPRİLYPAKMAFDİFRGKTİKVFLSİEEFRDİVİLHMPTİLRELLGNİP

>sp|Q8NGN5|O10G8_HUMAN

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HYİCDAPİİKLACADİTSAİETVİFVTGİVASGCFVİLİVSİVİVCSİLİRTİSİEGKHRAFQTCASHCİVİLCFFİGPGFLİYİLRP
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>sp|Q9Y5M1|F215A_HUMAN

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>sp|P49005|DPOD2_HUMAN

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VSDİFRİYSİMEDHİİLEWİTLRVRHİSPTAPDİİLGİPYFYKİDİPİFİPECPHİVYFCGNTİPSFGİSKİİRGPEİDQİVİİVTVPDİFSATQ
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>sp|P59646|FXD4_HUMAN

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TC

>sp|Q9NRF2|SH2B1_HUMAN

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EGPİSEYİMETVDAQHVKAWVSDİİQECLSPGİPCPATİSPRİMTİLPAGTİSFLİTRENTİDİSİELİSCLNİHİSİELPSQDİİİİGPİSİENDRİS

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>sp|Q7RTY8|TMPS7_HUMAN

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>sp|Q15572|TAF1C_HUMAN

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>sp|Q9Y6U3|ADSV_HUMAN

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>sp|P31749|AKT1_HUMAN

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