

>SlCycA1;1

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>SlCycA2;1

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>SlCycA2;2

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>SiCycD6;2

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DQSEHRDNEGLIFNLQSVHRMENLILTTLNWRLRSITPFAFLQFFNSLFELSNNQSLQSLE
DRASDVIFISHYEIKLVEYKPSVLASSAFLCAAHELIPQQVPFFLDAINSKCGHIYKEEVLNC
WNVMMREVSVDVVCSSTLPNSVLDQESTSENADRNNSSKKRRLNDFRNNQTFYISQVQH
Y

>SiCycD6;3

MEFDLQNPLTGFEYHFDVSAFANESDHMPDFKSTDVRFSLRRHSVFLISQVQSSY
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NFDARSIRHMESLILDTLKWRLRSITPFSFLHFFNSFLKLGDSSTLRSKQRASDIIFSTQYEI
KLFVYKPSIIAASALLRAAEELIPLQYSSCLAAISENQYSNKEELMNALAVMREMPIEGREY
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>SiCycD7;1

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ALSAKMNDNIDDVPSLSEYPLGPYEINVNHITRMERLVLDQFSWNMNCVTPFDGFFVS
RFCRDVTRLHITRITTARIIMIALRDLCLMNRHPSVIAAAATLLAVNRDFTIQELVMEINDLP
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>SiCycH1;1

MADFVTSTHRTKWIFTPQDIKHKYK VANHRAKQALEKYGTTRMEVDIDGSFSYAESQND
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HHPKDIMLTCIYAACKAEENHVSAAELGKGIGQDHHVILNNEMLVFQSLGFDLIAYAPYRA
LEGFISNLEEFCAQDNDQLLALKGALDTARIEADKIMRSDGPLLFPQGQLALTALHRADA
AHGIFDFERYLRSVLSHHDQPGHAISELTDSINVIDSLVGKLLTPTSKDVKHIDRKLKSCLD
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>SiCycH1;2

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LEGFISDLEEFCAKDEDQLVALKGS�DTARIEADKIMRSDGPLLFPQGQLALTALHRANA
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>SiCycL1;1

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SSKKYVDLKAIDLIRTERHLLKEMGFICHVEHPHKFISNYLATLETPAELRQEAWNLANDSL
RTTLCVRFKSEVVACGVVYAAARRFQVPLPENPPWWKAFDADKAGIDEVCRVLAHLVSL
PKAKYIPVCKEGGSFATSNRSRDSPLSPVSKEGLLDAPPVNEDTEEVAKEADTKAVLDK
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RRHH

>SiCycU1;1

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SLLVSLNVHRLLVTSVMVASKMLDDAHYNNAFYARVGGVSNAELNKLELELLFLLDFGV
NMSARVFESYCQYLEREMLSNGPTLKIEKSAINSITTSTVDDATEISVEDIDHTSSPSQLLD

>SiCycU2;1

MANISPRKLRDDLYCYSYEQDSNIPLVISVLASLIERNLARNERIAKNCTWALSKNVTRTVF
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ASKYVEDMNYRNSYFARVGGGLTTNEMNKLEMEFVFMGMGFKFHVNVSVFESYCCHLERE
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>SiCycU3;1

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CTSAHLTSLNVHRLLLITSVMVAAKFIDDAFYNNAYYARVGGVTTKELNKLEMKFLFGLDF
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>SiCycU3;2

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ITSIMLAVKFVDDDCYNNAYYAKVGGITTRELNKLEVKFLAALDFRLHVSVESFDKYCLQ
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>SiCycU4;1

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PKLQYICFNEDESSSQQLVV

>SiCycU4;2

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ESSTHELDVSQII

>SiCycT1;1

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GIGDGEMNDRQTHEREQVSYQGSTAQVQSRQSRYSSESNTEDDQEGNTRKSEPRHKGESK
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>SiCycT1;2

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SKRDRSWLEDTEEDNIFMSKEEIEKCSPSRKDGIDAMHEAHLRYSYCAFLQNLGIRLDLPQ
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LMELF

>SlCycT1;3

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SAPLRIRQKDIYDKEKESVVAGERMLLVTVAFDLSIQHPYKALVAAMKRLEISNNEMVKVA
WNFVNDWLR TTLCLQYKPHYVAAGSMFLAAKFLKVKLPAKKGNPWWMQFDVAPKKLE
EVIQKMLQLLEQNQKQVTPSTSSKLTESKPVAEKATSSIAESCISSVSVVAQESRNIGLVKTR
GLSTSVTSKFIEKASCNSINTVKEETEHWETSECGSANS AVENGVCQPVKNEGGKEICHAV
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>SDS

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DPVDISRDIVSKRNRNAKVIEGTEDSEVISRFLKASGGFCGESSKSGEDAVARSRNAAKIIH
EDVVSFNSVLQSPSESKCGNLSVQSIKCSENRAAEVESEVSRVCPEVELSAVEQAHEKLV
EAELDLECSENFSIVDVSDDYSSAYSELQSEIFPESSDIDISDYSPSYWYDSGSQFSEKSNAD
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YLHDYAE EYCSTTDYGDLIVQQRLQMVHWILEQATR KDLQKETMFLSVNLFDRFLSKGY
FKTKRCLQIAGIACLTAVRIEENQPFNSIRQKTFSVAGTTYSCSEVVAMEWL VQEVLNFQC
FLPTIYNFLWFY LKAATATEYMEKTAKYLAVLALLGHEHLCYRPSTVASALVILALSAANL
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>J18

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GMKSIPFRFFEDLLLKFSEVARV GKQLSLEACMDIMDLVYEKGKISSFNCSSHHLAASIVVA
AYAITVPLQQSEFPILLWVKFVTSCKEEDIVVTVKNILMYVFEA