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TGTGHDFNE
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lupas 21
lupas_2
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QUERY UniRef90 Q8KWG5 UniRef90-T6MQH1 UniRef90-P46006 UniRef90_Q9X4L4 UniRef90 G2B334 UniRef90-UPI0003EE1207 UniRef90-D8EF36<br>UniRef90 B3V22<br>UniRef90 Q9XDG6<br>UniRef90-Q9XDG7<br>UniRef90_G4C9M1 UniRef90 A0A028K61s UniRef90 ${ }^{-}$A0A034T3D UniRef90_UPI00041592B5 UniRef90 UPI000478D208<br>UniRef90 C4UJG7<br>UniRef90-UPI0002E0A4F-<br>UniRef90-D4BUN4 UniRef90 LOMML6 UniRef90_L3G4Z0<br>UniRef90_E0LZA3<br>UniRef90 U4W52<br>UniRef90 Q48BA6 UniRef90 B2Q5R1 UniRef90- F915A8 UniRef90 U4W0F8 UniRef90-B5R1L8 UniRef90-B3IA45<br>UniRef90 S5T0Z8<br>UniRef90 X2MX62<br>UniRef90-G9UCR8<br>UniRef90_G5NPB4 UniRef90_A0A028K442

ERQVDGRAE ${ }^{50}$ Y VVQSKDGR ${ }^{60}$ E LDRVRTGGDGWSPVKGEGGK ${ }^{80}$ ARQDGGRPEHYIVQNSKGQHELRVRIGGNGWIISSFGEAQR DRQEGGHPRHYVIQNKGNQGELRVRIGGNSWFSSLDEEWQ ASQNNGKVGHYIVQNRETKHELKVKI GGGGWSSSLDEGQR $A R Q V E G R P G H Y V Q G K D T Q S K L W V R L G R E G W T S P T G G G Q Q$ ARQQESRPGHYI I QGRDSRHELRVRIAGEGWSPSVTGGQE AQKAGYQPEHYI IRGQNSQNEIRVRLTGDRWSPSVDKGQN AQKAGERSEY I KGRTKNQHEI RVRLGGDGWSPSVLKEQY ARQEGGGAGKYIVQSTEGQHNLRIRIGGNGWSSFVEKGDQ GREDEGRPGHLLI QGKDSRNEIRARVEGEGWSAIT EGEH SRQMGNDPGHYII LGRDSHNEMRVRLDGAGWLPSVS DGQ AERNGESPDRYVLTGKSNQHKLYVII GQNGWVP DNKGGE EQKAGDSPGVLLLSGQDTHHHLQVRLEGEEWQP DTASGH AERNGDAPNRYVLTGKNSQHKLYVII GQEGWVP DTKDGL APRSSEAPNRYILMGKNNQHKLFVIIGQAGWQP DSQGGL VQTSSFPTGHFVLSGKNDAHKLRIRLVPRGLTAEGSPGSS ANKSGAQPNSYVLLGKNNSNKLRVRIEKKDWQP DHEGGK AEAEPGHPGRCTLAGRDPGHRLRVRLSGPGWQAGAAGDGT GTQVQGQVHRYQLMSARG HPLNIRLAGKDWQP DAQSGK DKPVNGQSNVYHLQGRNPSNELTVRIEGNGWSPSD KVGN QAVSGQQGGRYVLTGQNPAHQLRVRLVPQQQVVTEMADMQ AYAAGGSSARCVLTGQRHANTLRVRLSGEGWLPGPPPEG
 APLNNGRSQRYRL FSSGKQ LKVKI G G
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GETWNTFNTNTPVNTVTIGDQDTPPDTYPMTVDVVGYQP GETWSTFDTNTPVNIITIGDQDIPPDTYPLTL QWF VYN $\square$ NQR LDVKVAGDRNIPADSYPLNINDAAYQ

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jnet
conf
sol25
sol5
sol0
jhmm
jpssm

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[^0]:    lupas 21
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    lupas_28
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    jpssm

