## QUERY

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20
保 VRLPFSGFRLQKVLRESARDKIIFLHGKVNDSEDAVVILE VFTLSGFRVEEVLRESAREKTLFLHGKVGEGKDAVVIL VRDPFSGFRVKKVLRESARDKDIFLHGKVNDGEDAVVILE RDNFAGFKLIKKLNENTRNKCVFVHGSFEGDDDAVVLME

GFRLSAVLRE
EDDLRA
FQLTRIRDNPREKSTFIHGKVKD EFKMSKVLSESAREKTMFIHGKLV
SPVS KFKFEKLLNQDQAGRRTS

RSTLS EFKFERQLNK K VESLIPKKKFERQLNQ VPCDFTK FQDFRKRILN FQDFEMTKILNDHPKNKSVFVHGKFK FKLDRVLN KFKFERL N

VE SLI KQDAGRRCALYGA! ESLLI P K SKFEVSKVLNQ | D |
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FFQFERLKN QFKLERLLQNNCVRKSMCI EGTFED
FKOFKLERLLODQAGRRIVLLGSIQ SKPALLLA
NFKGFKLKRILNQNLSSKSVFLLGNLSGNDQAVVLLE SSLLAGFTFERLLREDPRTRFIALLGAFKEGAPAVLLAE ELNLSAFEVTRVLQNNCARKLICLEGKFEDREPAVVLLE KFKLDRVLNQDQAGRRTSLLGTVDSTPALLILE
$\square R$
LPKFKLEKVLNQDQAGRRVSLYGTI DNEPAVMVVE VRFPFSGFRVKKVLRESARDKRLELHGKVNDGEDAIVTAE SLLT DFKFTRLLNTTSQGRRIILLGTIS SQPALLIAE R $\square$ D
 QALE! EFRFSRVLNSNPQTKQISLLGTIK QLELK QF OLKNNNTNKTACVVGKFKDKSVALI LFE DLS $\stackrel{L}{\text { LS }}$
LK K FLKRILNNTVRKSI KHFQFVRVLDENPATHYAALETAD LHRFELERVLSHDTFTQTIALLGSFPAPDPAVVRI E ESTP RF QFRFTRVKSDSOTKT NLAKFKFTRVLKSDSQTKTISILGKIN

ETL

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## QUERY

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RAAFATDSAEVLRAFHSSVRTDNDI KNAYRESDVPSYFSSDLKVRTDFVNN KNAFPVDSWSAMLTGSTVLKPIMKNDI KLHFGSQELPNLLKELQWISTDNNDI RAPFPTSYAEGVAVGLDSVNNLGAND, KTPLNASDASKFLGLVRRIQLEESTDI KTPFQEESDPDLLKKHTKLQLQMSNDI KTHFHCENEYSCINGWEDLKEELTSNDI KTHFHRENQYSCLTGLEGVQEISSNDI
KNAYQESDVSSYSAELKVDTEFINNN KNAYQESDVPSYFSAELKVDTEFNNND KTHFNCENEYSCDNGUQELKEITSND KRAAQSEQGRTFFTTSSRLHKEFVNDVYGNFLCTVDPELN KTALPAFSDD LVGSTKLIESTDIYSWLFGWSSEERP EVPKGDDTLKKMFLGAPEVQLQNNNSV
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 $K S H F Q S N D T V D L N Q T V Q E L Q L I Q N N D V$
$K T A F D L N T L D Q L A T L Q D T K L E N N D V$


 RAPFPSSYLGRLPKSLARLQNLGANDV KSHF VVDHKSLASLVQDSETINQND KTHFHVENEYSCIGG|VEELKLLSNDI KSHFDAENLTNGLDKVEDAESINSND
 KSVFNVDSUNSLSHCLDTVSTINSNDVYNWGKALLNQDLQ KTHFDEAYLKQFASPDPEVKQLENNDVYHWNLATLAQDVN KSHFLPDTQSEDIKSLTAIEQLFNNDCYFTGVTQLSQKLQ KTPYASDEAPSLPSAWSKLQTVTSNDIYSTSLGWPGRATA KYAFDEASVLNVCNAQTKLKKNLDNDVYKSFSCVPDENTP RSHFPVDNHFDVQEWVEKLELLENNDVYYFWSSALVKQSLP KQNFSDDELKELENKDTVLHKLYNNNIYGNYECYPLKKYN KTHFQAQTLQAMEAGDRDLKEVCKNDVYFWGVSELEQDHP
 RAAFPTALLSQLPSTLINLTSLGVNEIYHWYLANSPESTA RKPFDPTSIEECTR QTKTKETLHNDI
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| UniRef90_K8EMC5 |  |
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P
N Q P T|IV|K|V|E
S S L L D G F K L A R V L D Q D P K L K K S A N L L G E F S N T E Q A
 VQELFRRLFKFEKVI NEDPSHHTVNLLGTI KDGQQAIVII N

L D S F R L E R I L D Q D P K QK S AN L L GT L E H GQQQA L I L E

IE TLL K R F Q Y R R V L N S N P Q V K V L S L L G S I D R K D A I L T F E


LD GF K F E I R R I L L N S Q N P Q N K L I I T V L L G K
 $\square P A L L$ K H F K L D E IL L D E P E K C RITLLLGTLP D GPPAMV RVVE FKFVRVLDSNPQTKVLSLLGK\| DDKDAVLTAE LT GFEP ERILSESSTMTGSTFILGTLABGQQAIFHVQ


 $\square$ NME TF K F K E I L A E D G S R K T A F V L L D V N D S N Q A I L I V E $\square S K F E L$ QRVL Y E D P R S K L CN AL G H I D DNEPALLL L ME
 $F S G$
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## $\square \mathrm{KE}$ L

 FR GLLKD D K V I NND P LI I K K A A I S G L L $L$ N D A K F Q $V R D D L S T F Q L E D I T T K M D S E M C V V G T F P G N D K V L L V L Q$

LA KAKYEEVLGADSSHKTEFLLLSHPNGEQGILDAN
D D AVLLLA
LD GFK LD R ILDDQDPKSKSVNLLGTLSSRERAILLVVE
L Q Q F R P S QV I N WD D NTK RL I L L G S L N D


$S N T Q \| L R V L R N D T R N K M D C L H V K F K D D E D G \square I I L Q$
LK D F VFEKVLSNDVTNKQIVGLGHMKNGDKALLKLE
 $R$ L N F E G F K L N Q I L S D N I Q S K S V F F E G S F E A T D D A V V L L E VRRLP F S G F R V V Q K V L


 P A $1 . V$ R V
 ENMHLISVLQNDLRNKCLVLHTREGDEKEAIVGLQ
 R A F K P V K V L S H D P VA H S IV L L G T HA H G D Q A L L L
 $E N M H L M S V L Q N D L R N K S Q V L H T R F D D S K E A D V T L Q$


LH NAKVVEVLGSDSSHKTVCLLLE SNGVKGGLVAN L R G F N L V K K T D T E L A L L G T F N N K N E A V L V D IK Q F KVDAVT LE S E N V VY T GT DK N E P AIICI FP GKKAALL: FP
$R L P L R G F K T S R V L N D G K E S G V V A L L G T F P G C D E S V V K L S$




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560E77




80 KTPLIDEDTLSELFSG SRLTLDMKNDIYSTYRLQPPPHLN KTHFSESFYEQLEGPSSTLSDLGQNDIYTWILAWPDPTDP KTHFDKDALQSCTQNDRGLKEVCKNDVYFWGVSELEQDHP KTHFTNQQLERTNSQI QRITTIGANDVYQWSLAWLNPQVP KTHFSEASFSC|GSLLESVANLGQNDIYTWLLAWLSPSSA
 TAAMDVGALGQLLTGMS LHKMLTNDIYSTFQGDVSRDFK KTPLDDPDLHAVFRSVSLHEVLNNDIYSTYGGDVSRDFK KQNFADDQLKEFFSKDTVFQKLYSNTIYGNYECFPAKKYS K LPFHRADRNHWLQNDTLCQNL QNDVYTSYRASDSQK KLHFQSDDENS SSLSSSVKQLFHNDVYFNGVTGQGDGFN KTHFSERFFSRLDDGAERVSDLGQNDVYTWLLAWLQRSAS KTAFAAECGKDFFSLVRK円EVMESTDIYTWLAGWGSSPLS TTPFQEEKDEELLRKHTKLELQMANDIYSTYRLYPPPELS KTHFHCENEYSCVNGTEEVTEIASNDIYHWGLVLDKQDIP KTVVVGKDAQEAINTLENIKLLLENAPYYSAHAWTKPEPP KT TLELKDAMLFIEENGLFELIEKNDPYHLFLGGDPKKSP KSPFSEEDITATVKSAELTEILKNDIFGNYDIIIPSNLN KESFLSESISKI FQEENEIKINFITNDIYYGYQYFPKSDFN KSAFQVNDDFSVSQSUKNVKQLLHNDVYYTGLUSYDLQDNP
 KTHYTPETLRQLASTFPRIEVIGQNDIYVWLFGWEQEEAC KKPFDVEDVKRMLNAGAEYKLDFQNAEYCVYNLQVPPALDN KTHESEDDDSNAMGSCRRDTDLGQNDDYTWLLAWSRDSMA RTPLSADFPRHLSSGLVQD KTHFSESFFEHLTATFSTLSDLGQNDTYTWL LA WQSETDA KSAFHESNFNQFDHLTDNHVMIQQNDIYHVLTSSSTPTPN KLTFDDENVDQQQQTSTSHKLIQQNDIYHVLLSDSNDKLD KSPFSEEDIEKLLATAQLQEMSRNDIFGSYNIEVDPKLN QYPGEEDELVMFLLSDR AT KTHFHCDDEFSCVNNDECTREVASNDVYHWGLVVLKQHDP KQKFAAGDACQLLANLSTLKQ1HANDVYSNHUGVLSGDND KTPFSEQQIEELGKS AKLKELSKNDIFGSYEDEDDKHGN KTAFTEDSVENMLSGESTFKVNIQNDIYNTLDLYPSALLN KKPFDAGDLDGAGPGLELVQLDGQNDIVTWLLGFGSRSVD
KKPF EKEVQQLAFDQAVQYGHNDI KTHFSEESQASKLSKATHLSSDGQNDIYTWLLAWSARSDA KKPFQSNECQEMLSECODLDLDLKNDRYHTFDGTDAAKYQ KTAFDANVVQEMLTEKTELVETLKNDVYSTYKTFPDKTLN KSSFPSEVSALCAAKTGKVEEVVNNDVYHRSSVMAGESVN KLAFEEKDLQKLQEELIEKDEHYFQNDIYNKMFIWMNKDIS KLPVPKK SVVEKLLNN ADLALNLK SKSVANYACYPKVDRG KTPFQTS KKPFLDEYKNNLNK【TNQNKY FENDDFTRYNDLQNDEWN RTSLDTKRAPQLVKGLEENLDVFDDNRPYFSAHALLPDSLP KRPFD SKXCIPNSNI IQAQSNDVYFWGTGSRSTEDS RRAFDEGRLPFLLTSATHLDIQFHNDVYAKYTGTDDPSDG KMSFNESII AQLGQDDSNLTLDFQNDIYSKFMLNMTPEDS KQSFPGDALAHLLSLPME NTQSNDVYHSYRVLSPTH TAVMDSDTLDRLLAGMSLLE ILVNDNYSTFQEDVARDVK RTPLTEDLLVDDUMSPATDYSIEMKNDUYTSLRLFPKVSAG $K T A F A A D C A K D F F S L V R K D E V M E S T D I Y T W L A G W I S S S A G S$
$N Q P V P Q D T F S Q F L G P M E S T Q S N V Y Y Y K V L G N M$ NQPVPQDTFSQFLSLPME KRGQSNDVYYTYKVLGNM KTPFPTDPVHKLLTSSDEAKSULSNDVYHRLLVTAGESLN KVAFNEEY KVAFSOEF AWEKLESLESNDIYTQQRLKLVRETP KTAFDPFDISFFTGNGLHBNLIGKNDVYHLGSATDAEHSK KSSFPSDPVNKLLSLHTQAKSHMTNDIYHRFFVTNGLELN K $\square P F Q E E N \square T E L L K K H T G L Q L Q V S N D I Y S T Y H L F L P P E D N$
KTPFP EAVDQLLGLVPK
AAEMDTGDLDRMLAGMS $\square H E T M K N D T Y S T F Q G D V S R D V$ KKSVLQDEFLDVIKEDVELKL HNDIYYAYTVKAKVDL KQLVLPDTFSQVLSLDPK NTQSNDI YYSFKASVPMN



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[^0]:    lupas_21
    lupas_14
    lupas_28

