


## QUERY

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PVTTNQASA GATDTVTYQGCADAGFCYPPET 1100
KTVPLSETVV 120
 $P V \operatorname{T} T \mathrm{TA}$ GKEASLKVVYQGCAEAGFCYPPETKVVPLREVA PLTVTQAAKGGTLTVVYQGCADAGFCYPPESKTVPLSEVA PITVVAA NKGSSVSVTYQGCAEAGFCYPPETRVVPLSAI A

 N I P L T A A Q Q GA L V R V T | $P$ | $V$ | $T$ | $L$ | $S$ | $E$ | $A$ | $E$ | $Q$ | $G$ | $A$ | $T$ | $L$ |
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| $P$ | $V$ | $T$ | $V$ | $V$ | $Q$ | $A$ | $L$ | $T$ | $G$ | $A$ | $T$ | $L$ |
| $T$ | $V$ | $T$ |  |  |  |  |  |  |  |  |  |  |


 TLPITEAAQGATLRVTYQGCAEAGFCYPPETKVVPLT PVVLKQAGPDATLSVTYQGCAAAGFCYPPETRSVPVSEDA


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$P L T L Q Q A G D N A S L S V T Y Q G C A E A G F C Y$ C

$N$ L R QNA D QNA T L L,

 P V A I K



 P L T L L Q A D G D A T L R V TYY Q G C A A A G F C Y P P E S R D I P L S K V A

 $D L P R T D D Q R P F T L V V T Y Q G C A D K G L C Y P P E T E R L S I A G$



 DVPLNEYQTGARLIVQYQGCAKAGFCYPPETRVIDLSP



 TYPI LQSQDDSVVKI RYQGCADAGLCYPPTTQVVYLNAV



 EI PRPNE QRPFTVLVVGYQGCADKGLCYPPETARLDSVDGTA ELPRND SRSFTLAVTYQGCADKGLCYPPETERLSIEGNA $R M P V S G A V D R F T V V A V S Q G C A D K G L C Y P P M T L K L T M S P Q A$




 $R V T$ APAAQA LT $\quad$ T WQ GCADA GLLCYPP

 CYAPETAHVDVSA
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 S E ASQDEFPVDQA QQDFVAVDEA
N
SSEPV EANAKPKFLQVDEAFQDESEQRGDQLLLTLHI ADDYYLYR ASNAPTFVPVDQAFPFNFTQQGNRVYLDWQVTPGYYLYE KPTFLTSQQAFQFDAKQTGNQLALNWQ ADGYYLYQ ATQTDFLPVEKAFRFTWERDDGQVQLRWQUAPGYYLYQ SSGADLLPADEAFQLQVTPNNNQLELDFRVADGYHLYK ATEDDFLPPEQAFRFAASQDDKTI EVVFDVAPGYYMYR SAARADDFLDPSAAFRFSASEQPGEVDVKFKIADGYYGYR ASNNTFVPVDQAFPFRFEQEGNRVYLDWQVKPDYYLYQ PLAQEQAFGFEADAHGNQLLLRFTPARGYYLYR NNNGFVPQDQAFPFNYYQQDGKVLLRWQVKEGYYLYQ SNSRFVTVDEAFPFSSFQQGHQI YLDWQVKDGYYLYQ SNNNTFVPVDEAFPFNHYQQDANLLLDWQVKSDYYLYQ PLPADQAFNFEAVVDGNSLLLRFTPAPGYYLYR APKQDTFLPVDEAFKFDFDQQGNDLF GWD AEPGYY YQ APPQQTFLQVDQAFVFDFDQQDNTLFLGWDAPEYYLYK ARAADDFLDPSVAFRFSAAEQPGQVVVTYKIADGYYMYR AEEFLDPADAFKPTAQADGQTVQVRFEAKGYYGYR AYGGDEFLPPEQAFRFSAMQDGQTVEVKFGUADGYYMYR SLAGNADFLPVHEAFKLSTQADAHAIKLRLVATDGYYLYR NSADFLPVREAFRLNVDSSPEQVKLRFVAAEGYYLYR
NSNDFLPVRDAFRLNIDSTPETIKLRFVPTEGYYLYR
NSKDFLPVREAFRLSVDSSPTSIKLRFANAEGYYLYK NAPAANTFVPVDEAFPMNFYQQGNKVFMDWQVKEDYYLYQ

GMPQLLPPDEAF VKKAMDAQTVKLDYAPTANTYLYR AQAEEEFLDPEVAFVMSATQTPDTLDVHFKDAPKYYMYR NQDDLLPPEQAFAAQVSRSGDQLQLTLEVAPGYYLYR QNEFMPSQRAFAFDFKQQDNQLVLHWQI QPGYYLYR ATASEEFRPPDQAFAREQTSQGDQTQLHWDUAEGYYLYR NSADFLPVREAFKLSDDSSGDSTKLRFVPAEGYYLYK SQDSSFVTVNEAFPFNFMQQGDRVYLDWQVKPDYYLYQ

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NQNDFVSVEQAFSFNFYQQNQQLMLDWQVKEGYYLYQ N $\square F S Q P K F L K S E E A F V F S T Q N E N E T L Q L N W D I A D N Y Y L Y K$ ASAAEGFLPVEDAFKLTSEQRNGEVVPRWE APGYYLYR AASQALPLPSEKAFGFDA VDGNTVLLRFSPAPGYYLYR AASQALPLPSEKAFGFDALVDGNTVLLRFSPAPGYYLYR ASAKPAFLKVDEAFALESEQQGNRLLLRLREADDYY YR
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 AVDGRDLLPPERAFPVAARQSGGVVSLVWDVDGYYLYR EDGLLKPDQAYRPMVESDGKTLNVHWEIADGTYLYE AHAGDGFLDPSVAFEFSASERPGEVDVRYKIADGYYMYR SATKPAFLPVDKAFTLSSQRDGDQLVVTWKI ADGYYLYQ SRSNNPEFLPVEQAFVLSVNKRGEHLI ATWDVQDGYYLYK ADEDDFLPPEQAFVFSAAADPATLVLNYRIA ABEYMYR ALADDGFLEPDQAFKFSARVDGHTVSVSFLIADGYYMYR SAEDEFLSPDAAFGVSDKQDGQNIRANFTVAPNYYLYK

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N S A
NSADFLPREAFRLSEVTEQQLERFNAEGYYLYR
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EDDYLAPEVAFKFSARADAKTAEVTYAI ADGYYMYR AVKKNEFLPVGQAFAFSSEKESGQTRLYWQIADNYYLYQ QEFLDPLVAFKPEARADDKTVEVRYSUAKGYYLYR DAAGRAQPLPPEQAFGFEA\ADGGRLLLRFTPARGYYLYR DAAGRAQPLPPEQAFGFEAIADGGRLLLRFTPARGYYLYR
 S S E D E L L T A E E A F R F Q A E A A P D R I Q L S S E DEEL LTAE EAFRF QA EAA P D R I Q L T W WD I A P G T Y L Y R

 A DEFLDDPAVVAFKPSVRADGQTI EVVSYEIAKGYYLYR N P F A Q S D F L P VN KA F VV S S ER E T G QM R L H WQ I T D G Y Y L Y Q A A GDGAF L P PE QAFRFTARQDA QADEVHFDVA Q GY H L Y R
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 ERFAFKS GDATLGAP QLPPGKDKFDPTBGDVETYRGKVVV


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 $\frac{L}{P} V A \operatorname{L} T$ A GAS G Q V K V S WQ G CA DA G L C Y P P Q S Q T V D L G G T A R L P L T R VGG D A A S L T L S Y Y Q G C T E $\quad$ G L C Y P P T D K R I A V

 $K V P L A Q Q \quad G Q Q L K V R F R G C A D A G L C Y P P T C R T R L P$

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 $K I P \quad A G A S G Q I K V G F Q G C A D A G L C Y P P Q T Q V V D L G P D A$ TLPASAKGTLRMGWQGCADAGLCYPPQTSMVDLGGAA TLPVTRTATTAKLTARFQGCQNDGICYPPMTRTVDVELPA TLPVTRTATTAKLTARFQGCQNDGI CYPPMTRTVDVELPA QLP DADLASLEDGWQGCADAGLCYPPQSRTVELGGSS


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 A D D D L P A E EAFLLFSAEVAGORLQQTWDDVAEGTYLYS DSNDDGELLPADKAFAFTSHQDNGNLILNFKVTQGYHLYK DSNDDGELLPADKAFAFTSHQDNGNLILNFKVTQGYHLYK AVDASQLLPPEQAFVPQVKADKQGVTVEFKIADGYYLYR S SQQDFLPVEEAFRFELMDDVEPAVLVWQDAPEHYLYR SSSNQSDFLPVLEAFQPTAWHDGNAVYI GIDDIADDYYLYR AADFLEPEQAFKLTATQDARTVRLAWKI APDYHLYR ADDEFLDPQVAFKFSARQDPSTVAVSYQUADGYYMYR
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NSSDFLPVEQAFKVQAENSDGSVAI RFSNAQGYYLYR AVNPN EFLDPPAVAFQPSVRADGQS|EVRFEMAKNYYLYR ALNSEDLLPPDQVFKLTVTQSRQQLEALEWQUEDGYYLYQ ALINSED ELPPDQVFKLTVTQSRQQLELEWQUEPGYHLYR SRLSEPEFLTVDQAFQLSVSAKDGKLMATWQIHDGYYLYQ A DAGQLLPAEQAFRPTVAAGEQDVAVQFQUADGYYQ ASQEVLEPEQAFRHSARLDERTLALRFEIADGYYMYR
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A VD $\stackrel{1}{\mathrm{E}} \stackrel{1}{\mathrm{Q}}$ DTSS Q AVDASKLLPPEQAFVPQVEVSDKGVDVQFAVADGYYLYQ STN EDFLDPADAFVLTAAANPTELQVHFRIAPDYYMYR STNNDFLPVEKALPFNYTTDSGSLVLNWDVAPGYYLYK DDFLDPQQAFVFSAQADAKHI AVHYAI ADGYHLYR ASDELLEVDDAFTLQPTMKDQNITLSWKIADDYHLYR AVDPA QLLPPEEAFVPKVYAEEGGVSVEFKIADGYYLYR GQRDLLPVEQAFPLSVERSEDRIVATWSVQD

## ELLGP

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## SNPSFLPVE

QKKYLKPEEAY

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## Q

K N I

## N S

SKMTVTS GTATVLGNPSFPP
H M L E A L T L






 E Y L E AK S QA GEALDEVRSGP GK I K D DP GF GTMEVYY GTATA


 D Q FK V Till N N A P V K L N L P S G H E K D D P T F G R T F V V Y N N Q V Q S

 DRI AV GAPGARAGEPCLWPAATPFNDAHFGDVQVYFDQVEV

 SK】AATAPEGVLGEPVFSK

D DK FRFAA E PVQLGKA IL P P
SK $\square$ TAETAPAELADPEMSEGKEKEDEFFGKQTVYYHQAEG DKTK\ES D G\QQ GAF QMPEGAVEHDAVFGDMSVYRDRDLVG
 E QFA LSSGDAAKLHFSPFPA GEIK Q DPYYGEVVTVYREQLTLL EK $⿴ 囗 ⿱ 一 一 厶 儿 V T E P G L L G T A K F S Q G K E K E D E F F G K Q T V H Y H Q A V V$
 DKLSLKLQGVQLTPLQLP DKLS



 ERFKFEAVGAKLGTPVYPAGKVKFDDTFQKVETFRKTLTI D K L S V V GE DK T ELDP LT L V S GEVIDWD DYFK ETEVYR WQL EVV




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 $P V A L E A N A T G Q V V V R Y Q G C T \quad K G F C Y P P E E V T L D L Q A A T$

 A L S L P A A M T E G QL R RVT S Q G CAD A GL C YP PRTTQQFALLT GNLL S

 $T$ L P L L R GP TEEVTLLI A K Y Q G C A E I G I C Y P P Q T Q Q V T









## R V

$P L P V L K A P A$ T F K MLLV TYQ G C A E


$$
\begin{aligned}
& A \\
& E L V T P G B P V A S G S \\
& L
\end{aligned}
$$


$T$ I SVK P $\qquad$ EVVPLVRRSAQATVVARYQGCAERGVCYP PQTRRRLPLALPT
 $P L P L L R S A A P A T V T A T F Q G C Q T G G I C Y P P M T R T V A L L R L P P$ TQRLTNPPTNVTLNDQFQGCADTGVCYPPMNKTVSLS Q L P L S A P T R P M T L K A T S Q G C T R L G GVCY VP Y T H R F R I S P
 NWP YKKAAPTYKLTLTYQGCAEAGVCYPPVDTSFEIN

 $R L P V E R N S S P L I L K V T S Q G C A D A G I C Y$
$K W A Y P A A A Q N Y T L T L H Y Q G C A D V G V C Y P P V E T Q F D D K G A$

 PI Y YDAA GTQVNAVLSY Q GCADKGLCYAPQKMPI QFTVAA $K L P F Q A A P A T Y R L T L T Y Q G C A D V G V C Y P P V T K T L E D$ VV P LS DALTNSKL QV T Y Q G C S E K G F C Y P P M K QVI S L D G V



E P L A A A S T E TAE LTAARS Q G C A D L G V C Y P P H F Q T V S

$T \cap P V E A A G A$ FTLDKATGQGCSDKGLCYAPQDATA QL

L WPYK Q GVQ N NYKL K L A Y Q GCA DVGGVCYP PV EN T FE $R M P V T E G T T E A T F R V T F Q G C A K A G L C Y P P Q T R D V$


S YPFRNPA G GYK L TVVMY Q GCA D A G V C Y P P P S E T V F D D T A P
 SLPMLLNADAPFTLNVSYQGCAEDGLCFPPITKRFRVNP




$$
\begin{aligned}
& \text { LI SLDD ASAPLRRLHISFQGCNDTGVCYPPMEQTLLVPMAE }
\end{aligned}
$$

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## EGA

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ASA APAFLPVDDAFQFSAAKSENQVTVNWS AEGYYLYQ


FAAQDFLPPEKAFRAEATWNSNQLELEFLPAKGYYDYQ SADSPPEVDEAFDFEVVVDPKT LAGWQUMEGNYLYQ GREDPLPRDEAFKLKVVAKGNSVVAEFTPAKGHYLYK SAGQSKFLPVDQAFQVSTDSTGTRLSINFDITPEHYVYK NNHELPTPEQAGHVDANVDAEAIDLDFEDTDQVYLYQ SVAAREAPLPPAQAFAMTYVRQGDAVQVRYMIPAGYYLYR NGAGQSKFLPVDEAFQVSATAKGTQLSTIFNITPGHYVYK QSESKFLDASQVLTLSVTEVDGH:EEVGN:AKDYYVYR AQAADSLLRFNQAYRLSARAETGAVHLDWSIAPGYYLYR QAQDQLLSAEDAFSFTVESNEHEAQLSWHMPERYYLYQ SLFGDQDRLPVDEAVVLDADWDNQQVLLDFELYDNLYLYQ PKFLEASQVLTLAVSHVDGRFEVVGQI AKDYYVYR SAGQSKFLSVDKAFQVTSSSKGTQLSINFDITPEHYVYK PAQAETLRAREAF IPVARVAPGSLMVHYEVTPGYYLYK ANPLPAAEVFHVNVSVDPNTFAl NFQUKPNYFLYS
AFPDPA
A A Q
AKPLPASEVFKVSVKKDPNTFAIGWNGPKYFLYS
KFLPVKEAFQARAWRSDGRLFVGFKNAPGYYLHR GNDQPPTVDEADQVDADWDKQTVLVGFELLDDVYLYR NSAQQSKFRPVDKAFQVTATTKGTQLSAFDITPGHYVYK

SSSSFLAPAKAFQFRAHMAPNELVLQWVAAKHYHLYR NGAEKSKFLPVDKAFQVKATSKATRLSINFDITPGHYVYK
AESTEEFLTVDEAYKLSDRASDTSIIAEWVIEEKYFLYG NSASSSQLLAPSNAFRFKAQMAPNNLILRWLVAPRYHLYR


HAQQALLSAEEAFPVQVEVTAQEVQLHWDTPEHYYLYQ ANPLPASEVFHVNVTVDPNTFAINFQIKPDYFLYS
 ATQTDFLPVEKAFRFTWERDDGQVQLRWQLAPGYYLYQ
QNPTFLHHEQAFILSYDLTDTDVELTWDVAPGYYYYQ QNQA $\square \mathrm{PLH} \square R E A F P Y H V S V D P G K Y L V T W N L A P G H Y L Y G$
NSNFKNK LLPSQKAF LFNWYKKDNS DF DWWEDKEGYYLFQ SAAAPKFLPVGEAFQVTSSTQGTQLSINFDITPEHYVYK SDILPADQAFRVFPTQKGNVISVRVEVQDGYYAYQ RSPFLAPAKAFRFKARMAPNTLVLQWIAAPHYHLYR
 ES GVLPPDKAFNI ETVISDEAVLI SWDVKEGYYLYF AFVLDSSVPGNVLRFHWKIHPGYYLYK
 FEVSVHQQPEDLHVHWDI APGLQLYK
GYANPDPSAHVFQVNATKDPNTFANVFQVKPNYFLYS S S VD D GF L K VY D A Y QV QP Q T R Q T D L N L E WK A A D G Y Y L Y Q N N K IL L P A N EA F K L E T S L M Q D T L K I K WII E D E Y Y L Y L
 S QARANDFLLPPEEAF QFT LSVVQDPADDVRAKVAPEYYLYR


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UniRef90 B1XST0 UniRef90-11XF15 UniRef90-B5EKA6 UniRef90 ${ }^{-}$F5STH5 UniRef90-UPIO002881216 UniRef90-K2H2U1 UniRef90-V4NRX8 UniRef90-E2T5N0 UniRef90-U4T9B1
UniRef90 R7WU32

UniRef90-A6X401 | UniRef90-TOZU98 |
| :--- |
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UniRet90 U5T1M2 UniRef90-Z2D9B9 UniRef90_UPIo0041F91B9 UniRet90-E0XPM1 UniRet90-S6FH00 UniRef90-S3DID4 UniRef90-UPIO00410C99C UniRef90-L2FBF9 UniRef90_UPIO004650BFB UniRef90-C8PXH4 UniRef90_UPI000413143F UniRef90-UP100037A0838 UniRef90-E1VN67 UniRef90-B6BVM9 UniRef90-D7A883 UniRef90-U4T952 UniRef90-G9END2 UniRef90-Q21MK3 UniRef90-Q1MZT9 UniRef90_Q6Q931 UniRef90-A6GPI7 UniRef90-UP1000377AC4E UniRef90-K6VFC8 UniRef90-G0JM07 UniRef90-A0AOO9ZZG67 UniRef90-J5KCB1 UniRef90-C6RCZ9 UniRef90-A1WZ19 UniRef90-A4GJU5
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UniRef90 L6YKX6 UniRef90-L6YKX6
UniRef9O-A6QBE6 UniRef90-E7C112
UniRef
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 NSLRFQTGKLSNDRPSTPSGVEKFDETFQKLQVYKDPFLV DKLRFEDDSVQLGEFTLPGGDMKNDAYFGLSEVYTSDVQU NKTFFASSGVFVKEVRLPPGEVKNDAFFGRMEIYKNPFQA DK SLKPEGVTATPLKFSSSQNIDDPSFGKVAVFKQPQVT $K Q L K L T P A G V T A A P F T F S Q P Y S M D D P T F G K V L V F D Q A N M L L$
$H A F S F R D T T D F S D F T V P D G K L K H D E L F G E V Q V Y Y G Q L A L L$ EHDTASDAQGRTVKIRTPNGTMKADPTFGPVEVYYGTARA DRFSFKTDNVTVAGVELPPPLEKFDKAMGQNMRYYERVAV DQLT DRPKGLKASPFKFSQPVSDNDPTFGQVPVFTQRNMV HSLRAEDVSGQPLDLTLSDGAKHTDEFFGDTEIYRDALRI DHLEAKAATGEPIPLSTGPGVVTEDANFGPSEVYYLEAVA DRTHFKAAGVTMGKPAFPPGVVENDPYLGRLVVFYKHMDA HKFEVQQGTQKLA LKKPPATDLYDENYGHTQVYFQNVQF
$E R F S L Q D G P A G L A P Q L P A G R L V F D S N F G D M A L W E G T V H V$ ERFSLQDGPAGLPAPQLPAGRLVFDSNFGDMALWEGTVHV SQFSFTPATVPDSKDELPKGRTQQNSVRGVYQAYTGDVKM HSLRAEDASGSPVALTLSAGSTHTDEFFGETEMYTDDLRM NQLKLTPAGVTAAPFTFSQPVLDNDPTFGKVPVFDQKNMV

| D | $\mathrm{R} \mid$ |
| :--- | :--- |
| D | Q |

HQFALESRDPSVGEGEVPPGEPVDHPQLGKIMVFYDKVVL
QRFEFITRDIDFSAFQMP
DKLGLKPEGVSASAMKFS
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EQFRFFDNGDKV DATRP
DKLTLRPKGVSATPFTFPPGKPMHDPGLGTFAVYEGDDTV DKLTLRPKGVSATPFTFQSPVSDDDPTFGKVPVFTQRTTL ESLHFKPNKLEAVKAPLPPGLEKFDETFQKMQVYKQAFLL HKFEVR QGQHVLPLKLP
KRTKLASKSTHLGPLRLPPTVKKTDKQGHVYAVYRNQVSI DQVRLEVDSGSVTPPVLPEGEQALDPLTGEAAVYHDSLRM DSVT Q GADGSALALAFPEPQUNVDEFFGRQPIYKRPLDL KRLRFDGLDPALQ PQLPPG



 NR|QVSPRSAHLAPYTLPPGKPMDDPGVGTLAVYEGDTTD NQLKVRPKGLSATPFTYS ESIKIENTNETFKF
SKFKFSSSSNTDVQFNFP
DSLEASRDGAPVA LERPT
DRIKKTPQADHLGSIRFPATTKKVDKQGHVYTVYRNHVSUM
DKMKLSFADKTSHFGFEESPVVIEDPAFGKVAVFYEQATL NSUKVMANKEPMKHKDLSKVLDHEDDFFGKTKILKDKLVI

 HKFEVK QGLHRLPGDLPKAVEQFDDNFGQSQVYYQQVQF DS K IQNKKEPIRYKILESSVETEDEYFGKTKYYKDYL
DSLAASLGDKDLSHLNFPKGEIYDER KSMELKNNNELISYNIFSKDELHEDEFFGETQLUKKKFNV KQISTTPTKADI AAVQLPAGVWHEDEFYGKSEIYRKRL

NSVQVKNANIIPYKVIYSEQSTYSDEFFGTTQUCRKEFK KDLHFK\|IGPKSFELDVKKPEPVDYEG QKVYYGTLN


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K Y F K $\quad$ L $\quad$ K $\quad$ L





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UniRef90 B5EKA6

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UniRef90-Q5ZXP5
UniRef90-X0QLE9
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TFKIVINHPPपKAEVVEIIAYQGCT
110
R
 $N L P L D R T Q S D S T L T A H Y Q G C S E T G U C Y P P T R K T F D L E L D P A$ EIVLDRKSTTKRLTLGYQGCNEKGVCYPPQQKSFDL

 DIRHKGA I TLSYRGCQADGLCYDPVRVVIP
RVRDAGGNQAVKLVATAQGCAAVGVCYAPITKAFSVPA TTNNSKGAKNAPIV 1 GWQGCAKAGLCYPPEK $\mathbb{R} K T T V N D A A$
 RL GTEAPGQVAVTYQGCKKDSICYPPDTDTV TLPFSAPAGLQHLAVTYQGCHAVADCFPPHTEALSDT $\square G G$ $\stackrel{1}{N}$ ATEASKTYQVTWQGCAQDRICYPPQTDEFQ RQA IPATVREGTLQLSYQGCAEQGLCYPPQHGQLHFSRGA ALPLQSVPVDTRLEVRYQGCDEDTLCYPPTKKIVEFASP LIPLNHRSSGLNLNVNYQGCSHAGFCYTPDKKIVNLSEVK HLDPAT AT GPVTLHWQGCADAGDCYPPQTMTVALPAGM ATTTDS DNDNAAVVI GWQGCAKAGLCCYPPEKDKTTVVDVA TLVPVASPGKPSSLTVQSQGCSEKRVCFPPEQHIFRQUKA

 SAP AASESGP LTTMVVTFQGCSDEGLCYPQAQTELQ ATTMMTAAKNVPVVI GWQGCAQAGLCYPPEKVKTTVNDAV ATATLTNTTNDDVIVTWQGCADAGLCYPPQKEILS
 TTNNGKGAKNAPI IIGGWQGCAKAGLCYPPEK CKTTVVDDAV
 ITSNGKGAKNVPVIIGWQGCATAGLCYPPEKVKTKINDAA TLDKKAVGKPLYLELELQGCAEAGLCYPPMTLKFLLAGP
 NVPA DKPGETYVLHYQGCSDDGFCYAPETKADRDS TVPFTATGSPFVLRVWHQGCADEGICYPPAATLFRIDP EVEDSNSDKNNNIVTYQGCAEGKYCYPK\|IKSL P $I$ K GDD GLAASTITIT FQGCSESGVCYPPYQVT

HFAKSADKQTTFSVSYRGCAEAGLCYPPVTVEDNL SLSLPTVPRPAQLVIVYQGCADWGFCYPPQS PIPINSDTPVQFSVTYQGCT KGFCYPPETKRVE E IPLKNSYQGQVLTVMYKGCAKSGFCYPDT IVTVSVNE ASTTLTSTNGGEVTIGWQGCAKAGLCYPPEKDKTTVS


 TVVPVLATPDGFEVRVQYQGCAEKGLCYMPQSHTVTDPTNA


 $\frac{L}{S} D R S K L P A G T E L T M K Y Q G C D E A G L C Y P P Q K A Q L D I T P A S$
 QI RYNNPVDMAKAVMISQGCADAGLLCYPPMTTPV GID D $\square$ N S SLNLSEVLIVYQGCSEAGFCYPDQKNQL
 RIPLQFPPKADTVTSSYQGCNTSVCYPQVTTRLPFTAVV

 HLTAEPVGEGAQLYAEFQGCNEESLCYPPSSLEAELASGA KADLNYILNDSDVFDYYQGCSESGFCYPLQKNKL
$\square L P L G S D R T L K V R M Q G C L D T G L C F E Q Q T R I V Q \square$
A VP LAA A SDDFVLNGDF $\square$ GCTKS GFCYQPQQF GFSFKRVV ElNNSDLLDLSNILKYQGCSDSGFCYPMQTKQ
S S DEEIETNLI EI AYQGCSDRGFCYPVQNKNIK
K LK S R SNKTLPLVI AYQGCSESGDCFAPDQQKFNDD


 PV HF
PV H F
VI Q IQ Q PDELDTPFSLHAQGCQENVICYPPETWKDDPQALA $V D V I M T E G R P V D L V V T D Q G C A D V G V C F P P L E R R I H L G$ PVTVNQAAAGATLTITYQGCA
NI PVKEDTGDYTLEI EVVGCSDKGDCYQPFRKSFTFKGAK EINKSNIDGHNTFEISYQGCSEGTYCYPVVKKEIKD

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A NVLS AK DAFKLD 20
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 $K$ D Q RVK F SVVERD DESI ELLTWII A P K S F L Y Y R




A VE K P K L LTP E E A F NVT A THSNQ GVMI NVTLGEK I Y L Y D


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 A E PL K V E D A F K V K VT A D K $\square$ GTN F N F D



 F A E P K F L M P E E A F Q A T A H L K S K VLT P E E A F DVVQ FT Q S R Q GlV V I


 A A P $\square$ QP FKAFVVDST $\quad$ Q NEKI Y F DFK I A E D DYVYK

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## $F L Q P E E A F K P S A K M D D K T L G V E \| V M G D Q \square H V Y A$ $P$ VA VA A R MVGQGDSRAI EVRMRIHPGFHVYR A S $\square L S L N E A F N V K S N S Y N N S I S G D \square E L G K D \square Y L Y S$



A A AKPR $\quad$ VDVG
VFS LS T N D NKTT VKAN ITPEGYY A Y A


 NREIVVRLKI HP GYHIY S | $Q$ | $G$ | $G$ | $E$ | $L$ | $Q$ | $E$ | $M$ | $T$ | $L$ | $D$ | $D$ | $F$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| $Q$ | $A$ | $A$ | $E$ |  |  |  |  |  |  |  |  |  |
| $Q$ | $G$ | $I$ | $T$ | $V$ | $S$ | $A$ | $G$ | $P$ | $P$ | $G$ | $H$ | $K$ | A A A GD LVDSSAKDRASAA $\square$ NGHNAVVTLHIDRGWHVYA

 F S LIK TN EN K L I V V N A D LS A S R Q G N N V R VE LA I E K P WHI N A




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UniRef90-D5BE04
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UniRef90
R6XGN1
UniRet90-UPIO0035C18DF
UniRef90_UPIOO0479C0A5
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UniRef90-D6CMA9
UniRef90-SOEZS8
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UniRef90-UPIO0039F53E4
UniRef9O-U2EHL6
UniReffo-F9Z7G99
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UniRef90-UPIO0047EA918
UniRef90 U507C7
UniRet90_UPIO00248EB23
UniRef90 R6XRQ8
UniRefofo $=8 \times 313$
UniRef90-TOY6C2
UniRef90 RGNKPO
UniRef90_UPIOOO45E7308
UniR Ref9O-U2EPO3
UniRef90-C6XY42 UniRef90-D5BE04 UniRef90-H3NVP1 1
UniRef90- $\times 0 \times \mathrm{M} 3$ UniRef90-C6G412 UniRef90_W5X9R3
UniRet90_UPIO003B339E9 UniRef90-Q02CV9 UniRef90-W6TKB5 UniRef90_UPIO0036A5E91 UniRef90-C2FRU6
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 EQLKLEGSGIS AEEEKPKSTLHHEE GDHVYLESPEF DMFEVDSNGERANNTNMSEGENENDEFLGNVEVSKNDAST


DKLKDNIHPDK SDNDIKKPKPDEDD NKVSVKIKKPQTKELVLPKP DKLQVNGTKPK円KELNLPKPVEY DEF


D HKI YNKP M K K L
DKIQUKPKKVELKELTLPKPVPY DGFi VHFDDLNL


$E$
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E

KRLKFDGLPAE QHPALPEGEAHSDEFFGD
DKLQV NITKPKLTKNUNLPKPKKH DGFDKEYGYYDENFTL


DKLKVKVSDTDTTELNLPK
HKFTVQQGTQPLA LKLPKA AAQYDENYGQSDIYGGTFDL DKLKIEDDGIDFASITMADAVKHEGES GQSQVYESSPAV
 NQLKLYLNQNDUSSEILPPSTQRLNE GLGKVYYQDLNL DKLKVESDGDDFQKITMAD QPLTVEPEGCTAGELTAPAPQQ
 DKVNLDSAQTKTGDTSKTDSVPKADPERGTVNVYYDQARL DQVKMVSSEPYLYEVFFPEGKTKDDPFLEKEVSFYEYVQL NPFSVEPDGFKVAPFKDEPGVEWHDKFSKKMATLTAHTEA NTGVVQRGGHGLDQPSYPPGQDUGLRVDGKGVYENGTRG
NKLLFFADNKETNT TYPKGEKYHED
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 DRTFIEPSGDAFGPLDVPKSVERTEPDGNTARFHEDATF
 NKIKFFADNKEI NAI YPKGEK

TKVEFRPEGVRAGGLQAPEGKVYEDE DKFLIDGAGVRAQSAEYPPGHDYTDPTTKKVRVFS EPF|PTVVNIEGPSDYKKEGELKTAPVFKKTVIFE


## KTFDFQLPEGQEGEMQLPSFTQLDDNG

DYFYПEADGПEF LPVQYPEGTEHEDYY
NY GY EADGIEFLPVQYPEGTEHEDYY DSVVYTGECVF NPDK IFANGVLLTELEYPPGDELDVGLEDPIRVYSGNVKL


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QSFSFQSSKALSGKVDAAKPDVKHDPTF DMEVYYKKSVV
HQLLVEPAGLKVVSVVFPSGTDYSPAFSPKDVVYTGEFVL HKGAVQDDPANVBTVDEPKGHDYSFAFDDKDVDVYTGDGFAL

| UniRef90 E6L 809 |  |
| :---: | :---: |
| UniRef90－FOBK12 | RVRLRLKMQGCLKDQ |
| UniRef90－M5JQJ5 | SVKAGT |
| UniRef90－D2V838 | EDPEKGEI RVT |
| UniRef90－12K6Q3 | VKEIAGDYTLELE |
| UniRef90－E4U3 | G |
| UniRef90－B6BLC4 UniRef90 EOUS |  |
| UniRef90－ToZGW6 |  |
| UniRef90－T1C3L |  |
| UniRef90－W6EIF2 | V PQSL |
| UniRef90－Q30S55 | S |
| UniRef90－Q4 | D |
| Unip | SEQGLCYEPLSATY S |
| UniRei90－UPIO0046FE61D | NVSKR EFNLFYQGCKDKVICYPVQQ |
| UniRef90－－131HX5 | NSFLPPGPYHDKLKVHYQGCSDK MCFAPKDEELTMPA |
| Ref90－B9KCE2 |  |
| UniRet90＿UPIO00472CE34 |  |
|  | LIPIUSIUGKFKLEVQYQGCATSG\｜CYQPLKSHFEFD |
| Ref90－UP100047B7 | VPFKVDAKEYE ELNFQGCSKAGLCYA |
| UniRef90 A8EWS8 |  |
| UniRef90－UP10004 | DVPYKLDSNKYEI EIKFQGCSKAGLC |
|  |  |
| 㤑 |  |
| UniRet90 UPIOOO37BBFF8 | $N$ |
| UniRet90－G2HS47 | TIPFK\DTNKYE\EDHFQGCSKAGLCYAP可SKYLLT |
|  |  |
|  |  |
| UP1 | 俍 |
|  | DLPAGSYNVQLKVHYQGCSDK |
| UniRet90－A71200 |  |
| UniRef90－W0D774 | FIPLKNNSNEFDIQLNYQGCSYSGF |
| UniRef90－X1XED5 |  |
| UniRef90－K7TO50 | GLSKSSQTLKVKLSYQGCS |
| UniRef90－K1 KQ 24 | YLVEDAKKTGNLNVKYQGCSEETMCFMPKNKSFKLGST |
| Ret | ALPNLAQNQNNQ1 KLQFQGCSEQGL |
|  | Q1 PVKNPKSPYSLTIDYQGCQPGRYCYPP |
| E4U315 | TVGwosc indy |
| UniRef90 R6VQ84 | VGWQS DD |
| UniRef90－K2BXH7 | QLHYQGCSQDGFCYPPMEK |
| UniRet90＿K1ZW34 |  |
|  | TDNVPEGPYTIRLKVNYQGCSET CFAPKTEELLLP |
| UniRef90 UPIOOO3752559 | DLPGLT ${ }^{\text {P }}$ |
| $\begin{aligned} & \text { UniRef90_O6N } \\ & \text { UniRef90_R6) } \end{aligned}$ | DQ FCLPPTTKLTEVPIVV |
| UniRef90－D6CTN2 | ， |
| V2W209 | Q1 KLRYQG |
| UniRef90－J9UV | SLNDIDDNKINFLKTVYQLCCAKDNVCLQPQSKVI |
|  |  |
| UniRei90－UPIO0035C18DF | QLRYQAC DANACMPPRKDPVAVSLDA |
| UniRet90－UPIOOO479C0A5 | CTEGGV |
|  | RYEIISSGQGEAKCTVSYQ CCDA HICFPP EKELKV |
| UniRet90－S7TGX2 | R |
| UniRef90－D6C | － |
|  | G！PVKAGAQKALIEVHYQLCTDRN－CLDPTTVSVPLSFRV |
| UniRef90－R5LJT8 | TLILDAKNTNFIKANYQLCSEDNTCLPPKSEII |
| UP1 | KD APNGVYVDVI |
| UniRef90 F9Z | EIKGDGTGVSCIMTYQ CCDNHI |
| UniRef90－R6BJB7 | RQKYTGKQGKVTVTDNYQAC NNHACLMPKSKTLEL |
| UniRef90－UPIO00374B2 | N |
| UniRef90－UPIO0047EA918 | 㑑 |
| Unif | GEVKCTVSYQVCCDNTDCFPPAEKI |
| UniRef90－R6XRQ8 | RVKLQALPATIKCNIECQ CCDANVCMPPYSKD |
| UniRetg | RLPVQSTPGPHTATLRYQAC DRAAC |
|  | TLPFSAPAGLQHLAVTYQGCHAVA CFPPHTEALSLTUGG |
| UniRef90－MPIOOO45E7308 | N A |
| UniRef90－U2E | ALLDEVADNPTVKAVVQACNDSGLCLPPSTLSTSIPSTT |
| UniRet90＿C6XY42 | KVKLTGANAIVKGTLEYMVCDDSQ |
| ef90－D5BE04 | QQV ERKTAEAKASVEFMVCNDES CLPPTIKEFELEATA |
| UniRef90 H3NVP1 | PPADAPNVEGTSLKMT |
| UniRef90－C6G412 | AVPLSQNTVALEARVTYQVCC ST |
|  | TVPVTLGPDPDKLSVHFQTC DD |
| ef90＿UP10003B339E9 | TVPLASGTHAVRVAVGYQAC DEGSCYPPSEVTGEVNLSV |
|  | K |
| UniRei90＇UPIO0036A5E91 | KLPVVAEAGTHTVTLRYQAC DHA |
| UniRet90 | KIKLTGNAPTVKGTLTYMVCNDKQ CLPPEEL |
|  | RMHVQAGEHLMQAALRYQAC DANSCYPPKKA |
|  |  |

UniRef90 E6L809
UniRef90- FOBKI2
UniRef90-M5JQJ5
UniRef90-D6V838
UniRef90 I2K6Q3
UniRef90-E4U3X0
UniRef90-B6BLC4
UniRef90_E0US13
UniRef90-TOZGW6
UniRef90-T1C3U8
UniRef90 ${ }^{-}$W6EIF2
UniRef90 Q30S55
UniRef90_Q4PKB3
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UniRef90 A7GXU7
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UniRef90- G2HS47
UniRef90 U6C3K1
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UniRef90 K1ZW34
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UniRef90_Q6MIB0
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UniRef90 UPI000374B2B2
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UniRef90 R6NKP0
UniRef90-UPIO0045E7308 S

UniRef90-U2EP03
UniRef90-C6XY42
UniRef90-D5BE04
UniRef90-H3NVP1
UniRef90-X0XMX3
UniRef90 C6G412 UniRef90 ${ }^{-}$W5X9R3 UniRef90_UPI0003B339E9
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UniRef90 WGTKB5
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UniRef90_C2FRU6
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UniRef90-E8V5G8
UniRef90_G8P253
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A

UniRef90 UPI0002D96544 UniRef90_UPI0003B692DB UniRef90-UPI00047CCF82 UniRef90 Q312P4 UniRef90-13ZMJ8 UniRef90-UPI00047615DE UniRef90-I3ZCV9 UniRef90-A9IEZ2 UniRef90_D6SLI8 UniRef90-E8WYF4 UniRef90 UPI000371B2CD UniRef90-W4LWJ8 UniRef90-F0RHK5 UniRef90-A4A2G9

1
10
10 SAAGTPDLVTASQAFSVSDNGPARRVVVIRTVVMDGFHVNS ATVTKSTATSASVALTATDQE GWHVYA GLAA A H A R L Q P G WH I Y A QA TVKL HFRVEPGFHNS



ELTA VQPGWH Y A



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lupas }2
lupas_14
lupas_28
jnet
conf
sol25
sol5
sol0
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jhmmm
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110
120 KVPVVAAA GDHTVKLRYQACDHAACYPPKSLPVEI AVTA $K V P V V$ VRAGQHTLVLRYQAC DNAACYPPRSLDVQIAVTA YVPLPDDPAVQTLSVSALMCSAANCWP
TVDLKTPTETVPVDVRYQACNDRLCLPPYTAHLT
LLPFSAMQRGAELAVSWLVCKD A I CIDP $P V H Q \square E I D A G E H L V M R V Q A C N D Q G \square C D A P A T \square S V$ PLPADIPLEGLTAELSLLLCSDRN CWPVHKEEFDLDRV. QLDKSASAHPIQQLIRYQSCNDRVCLPPHTDTVDV Y 1 PGSSGPVTLEASAYYQACSDVK CLLRPVTKKVD RLALLSGPVPESPHKLRYQDCNDVNTCFPPRTVQVP
 $K A P F V W P P K T L T L A A G K Y S G C S D A G T C$ VPQRRELQ

[^0]
sol5
solo
sol
jhmm
jpssm


[^0]:    lupas_21
    lupas_14
    lupas_28
    jnet
    conf sol25
    sol5
    sol0
    jhmm
    jpssm

