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60
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 Q A P E S L L K K K H Y A DLLK D R P F F P G L V S Y M A S G P V V A M V W W G QASEDLLLKEHYADLSSSKPFFAGLVEYMKS GPVVVMVWEGD
 Q $\square P R E V A E K H Y$ GEHA GKPFFQGLVDFDTS GPVVAMVDEGK

 QPSEKLLSEHYADLSKKPFFPGLVKFMATSPVVCMMVWEGQ T P GKEHLEQHYSDLSDKPFFKGLVEYMNSGPICAMVWEGR
 QASEDLLKKEHY DDLKDRPFYAGLVKYMHSGPVVAMVWEGL TPTENLLKKQHYAEHVDDKPFFPKMLSYMMS GPVLATVWE GK
 $T P T K E F A M L H Y$ D D L K E R P F F N G L C D F L S S GPVVVAMVVWEGQ

 DP TK K TVVNVHYA E HTEKPFFHKLVSF LT S GPVVAMVVEGE
 NVERSFAEKHYADLSSKPFFQGLVDYDISGPVVAMVWEGK

 A P D Q GLLT Q HYA DLK ERPFFPS L L KYVVSSGPV AAMVWE GK
 H A S E QL LQ THY E A L K S L P F F T NLV V Y M S S GPVVVPMVV E GR QP GK AK F E Q H Y S DL S T K P F F G GLVVNYA A S G P V V A MV WE G N




## L

LLP
L P S S EQA K QHY GDLAS KPFFK DLVVYFSSGPVVGMVWEGL


 HATRPHLLEVHY QE LKKDKPFFKTVI DYMS S GP VVVAMVWE GL QDD R E L A E K H Y A E H $\square$ G K S F F E D L I GF DT S G P V FA M V L E G D VPPSRALVVE E H Y A E H K G K P F Y E K L V V G F M A S G P V C A M I W W G K

 SVSQSLAE QHYAEHKERPFFQELVSFITSSPVFAMVLEGE
 VPPMELLEQHYAEHKGKPFYPS LTGFMSSSAVVAMVWEGN QPSEDLLKKHYA DLASRPFFPGLVKYMSSGPVMPMVWEGL
 KPTEDLLAKHHYDDLKDKPFYAGLCKKTVSAGPFVAMCWE GQ
 Q $\square$ S D S L A K T H Y A E H D GK P F F P DL I H F IT S GPVVAMVLE G D QATKELLDTHYGDLVDKPFFPALRDYMMSGPVVCMVWEGK TASEELLKKQHYADLAGKGFFNGLDAYMASGPVVAMVWEGK QVSRQLAEKHYE EHRDKPFFGELVSFDTSGPVVAMVWEGQQ A P GK E HLLEKHYE DLK DKKFFP GLI A YMTS GP V CA MVVWE GK Q MTK GQAE THY A E HK DKPFF GE LV GF DT S GPV L A MV WQ G D
 QPSREHMEKHYA DLLSSKGFFKGMM DYMMTAP VVGCMVWAGE WLDRDLAERHYEMHK GKPFFDDDI A F DT S GPVVVAMVWEGE Q A T T E HLLELHY A D L K GK P F F P G L I K Y M A S G P VV V A MV WE G L

 QASDELLRREHYAELRERPFFGRLVDYMGSGPVVAMVWQGL
 HS T K E H L L K K H Y G D L K D M P F F P G L I K Y MA S G P V V A IV V I E G L A MPEAKAKEHYAE HVGKK F Y Q D L V D F I T T S G P S V S LVVVEGR
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## QUERY

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$N$ VVKKTGRVM| ${ }^{90}$ GETNPADSK ${ }^{100}$
110
NVM

 DAVSTVREMMGATNPLKAAPGTIRATYGMDIGRNVVHGSD EVVKMGRMMLGATNPLQSNPGTIRGDFCDDVGRNI CHGSD
 DVVKTGRAMLGATNPDA $N V V K T G R Q M L G A T N P A D S L P G T I$ EAVATAREMMGATNPDKA DAVKTGRFILGATDPKNS GVVAAARKLIGATNPLTAEPGTI
 GAVKTGRVMLGETNPADSKPGTI
 DVVKQGRTILGATNPLDSSPGTI

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 DVVKTGRVMVGETSPADSKPGTV DVVKQGRALLGAANPLASAPGTI
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$N V I$ \& $A T A R Q M M G K T N P A D A D P G T$ <br>
$N A$ \& $A D G R K M M G A T N P K D S E P G T$

 EVI SAARQMMGKTNPQEADPGTI 

$G V I$ \& $E V R K L M G K T N P L E A E P G T$ <br>
$N V V$ \& $R G G R V L L G A T N P A D S Q P G T$
\end{tabular} DVVAQGRTVLGATNPSNADPGT

$E A V A V V R N L L G P T D G A K A A P G T$ DVVKQARKMLGETRPLDSQPGT GVVDTARNMMGKTRPHEAAPGTI $N V V A L A R E L M G A T N P A N A N P G S$

 QAVKTGRLMLGASNPLDSAPGTI NVVE ETRQLI GKTNPKDALDPGI | $D A V$ | $R G R V L L G A T N P L D S A P G T$ |
| :--- | :--- | :--- | :--- | :--- | :--- |
| $G V V A S A R K M U G A T N P L D S E P G T$ |  | NAI SVARTMMGKTNPAESAPGTI $N V V K T S R A M V G E T N P L D A R P G T I$

 DVVKTGRVMLGETNPKDSAPGT: DVVKQGRAMLGQTNPLTSAPGTI $N V V K M G D M M G E T N P A D S K P G T I$
$K A V V V R N I I G A T D P A E A A P G T V$


## GVVKTGRKMLGATNPLDSEPGT

GAI AVVRQMMGKTNPQEAAPGT
GAVKEGRKMLGATMPSESAMGTI
 NAVAVVRQTMGATRPYEAAPGTI GAVVTGRKMLGATMPSQSEPGTI
 GAVATGRKMLGATNPGASAPGT,
 DATRQVRRMVGATDPGEAAPGTI DVRASRALGATDPADAAPGT DVVKMSRKMLGGETNPADSDPGT, DAVKTGRAMLGAATKPLEASAPGSI GAI AAMRKMNGATNPVEAVPGTI
$\stackrel{\mathrm{D}}{\mathrm{D}} \mathrm{N}$




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 HI SRQQAEVHYGEHKEKPFYQELVDFI TS S GPVFAFVLQGE NVEKS LAEKHYEDLSARPFFPALVDYICSGPVVAMVWEGK HLSREQAEGFYAVHKERPFFKDLVDFMDSGPVIVQALEGE MVTKEHAEKHYADLSSKPFFA GLVTYMCSGPVVCMVWEGK RPSVEHLEAHYSDLSGRPFFPALIKYMSSGPVTCMVWEGT SLPEERVVEQYREHLEKPFFPGLKAYDMSGPCFDMVWEGK
 QATEDFIKKHYEDLSDKPFYGKLCTFASSGPVFAMVWEGL HPTKELLEEHYCDLKGKGFFPSLVEYMDSGPVVGMVWTGK QDS
$V$ S K K QCTKEVAEEHYKDLSSKPFFPALVEYILSGPVVCMVWEGE R LAEATVNEHYREHDERSFYPSLKQYI MSGPCFLMVWEGK QV TREQAQRHYAEHEGKVFYDNLVRFIVTSSPVFAMVWEGD RI SQELAERHYAEHREKPFFPGLVRFITSGPVVAMVLEGP $Q L S K N D A A K F Y E V H K E R P F F N D L V D D Y M T S G P V V V M V L E G G E$
$M I T$ KTPKELAEEHYGELKEKPFYGKLVAYIDSGPVVCIALEGP KVDREVAERHYAEHRGKPFYEDLLAFIVTSGVVVGVVEGP KLPEQRVTEHYKEHDSKPFFPSMKEYIMSGPVFFLMVWEGR QTPREVAEEHYKDLSAKPFYKDLVNYILSGPVVCMVWEGK H LTRDQAEGFYAEHQGKPFFDGLVSFMTSGPVVVQVLEGE RVTRPRLPFPRCSRCFPSSFRCREIPLPSAPAHGGVWEGLD RLTPEIAGKFYEVHKARPFYNDLCSYMSSGPVVVQVLEGD
 QASEGILGEHYHDLRRKPFYPALLSRYMAAGPVVAMVWEGR HLSEEKAKGFYAEHEGKPFYNDLVEFMTSGPVMVQTLEGD TATEARASDHYADLAGRPFFGGLVKYFASGPIVCMVWEGT HLSKEQAEGFYAEHSERPFFGALVEFMTSGPIVVSVLEGD QASRSLAESHYAVHKGKPFYESLI QFLTSGPTVAMVWEGR Q 1 SRETAEDHYAEHHGKPFYERLI TFITSGPVVAMVWQGD WI SKE LA EKKHYAEHREKPFFKSLVVEYYVTRTPVVVMVVEGR

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 TVSEDQAKRHYAEHAGKDFFSELVSFDTSGPVFAMVWEGD QASEDLAKQHY DDLKDQPFYAGLVKYTSSGPLLAMVWEGD
 QLSRADAEAFYAVHAERPFFGELVEFMDSGPVVVSVLEGV QVTEADARDHYGEHEGKPFFQGLVEYITSAPVVVAVVVGK


 WASEEHLKQLYPDLKDCPFLPGLVKYVNSWPVFVQWEGL
 TAPKELLEMHYHELRERPFFPKLMAYMGSGPVVAMGMD

 HLSQEQAAGFYAEHKGKPFYDNLKKFMTSGPVVVQVLEGQ
 QLT DE L LNQHYA HLSAKPFFQRVKDSMMVTPVI VCCYEGV IPSSRPFAQKHYADLKDKPFYVGLVAYWSSGPVVAMVWEGE
 QLDRRQAEGFYYVHRSRPFFHSLI EYMTSGPVVLMVLKGP QLTPAQAEGFYAEHKGRSFYDELVAFMTSGPVVVQVLYGE QVTEE DAA QHYEEHDNKPFYPNLVKY TTSGPIVAMVLQGY VADRETAEKHYEEHKDKPFFKDLIEFITSGPLVAGVVEGP HLDENMAGGFYAVHKDRPFFKDLIAFMTSGPVMVQVLEGK QLSQAEAEGFYAVHKERPFFADLVKFMDSGPVVIQVLEGE KLSKEQAGQFYAVHKERPFYTDLVEFMSSGPVVILVLKGE QLTREQAEGFYAEHKERPFFPSLVEFMMSGPVCVQVLQGE LMSKATAEAFYAVHRERPFFQELTEFMSSGPVVVQVLEGE HLS RE QAERFYAI HKERRFFNDLVSFMDSGPVMVQVLEGK


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E VVV|KASRAMVVGTNPAEARPGT|L|RGDFSVH|V|SRN|VV|HASD D ATRQVRRMMGATDPAEAAPGTIRGDEGLDLGRNVVHGSD $K A V E V T R N T V G A T N P V D A A P G S I P R G T Y G L T I G Q N L I M A S D$



 GAAAVVRTMIGATNPAEAAPGSIRGDFGLD\|GRNVIMASD





 NAI A GWRSMMGATNPANAAPGTI R $\left.\begin{array}{lllllllll}N & A \\ N & A & K & N & R & E & L & M & G A\end{array}\right)$


 $N A V Q R H R D D M G A T N P A N A L A G T L L$


 | $N$ | $A$ | $S$ | $R$ | $R$ | $E$ |
| :--- | :--- | :--- | :--- | :--- | :--- |
| $D V$ | $L$ | $M$ | $G K$ | $T$ | $N$ | D AV KRHREIMMGATNPTEALAGT$L$



 | $G V I$ | $S$ | $S$ | $A$ | $R$ | $K$ | $M$ | $G$ | $G$ | $T$ | $K$ | $P$ | $L$ | $E$ | $A$ | $E$ | $P$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| $G$ | $T$ | $I$ | $R$ |  |  |  |  |  |  |  |  |  |  |  |  |  |
| $D$ | $M$ | $V$ | $K$ | $Q$ | $G$ | $R$ | $K$ | $L$ | $G$ | $E$ | $T$ | $N$ | $P$ | $L$ | $T$ | $S$ |

 | $N A$ | $L$ | $K$ | $R$ | $E$ | $M G A$ | $N$ | $P$ | $A$ | $E$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | A E A G T



 | $N$ | $A$ | $Q$ | $R$ | $H$ | $R$ | $D$ | $M$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| $N$ | $G A$ | $T$ | $N$ | $P$ | $D$ | $N$ | $A$ |
| $N$ | $L$ | $A$ | $G$ | $T$ | $R$ |  |  |
| L | $R$ | $N$ | $R$ | $E$ | $L$ | $M$ | $G A$ |




 E A V D AI AKHRDIMGATNPKNAAPGTI

 NAVVLAHREELLGATNPMKEAAAPGTI IVA Q GRAMLGAT DPLKK SAPGT


 GVI KY GRKLIGAT DP QKSEPGTI
 NAI DRWRTLMGATDPRKADPGT NAI A L NRELMGAT DPTKAAPGTI K A V Q G V R H L L G V T D P C E A DVVG S I HA A I E A WR QL A G GT D P K K A T P G S il N AVV K H R E I M G A T N P K E A A S G T V NAVLAHRDLMGATDPKKAAPGTI D AV AKNRALMGATDPSKAEKNTL $\begin{array}{llllllllllll}A & A & S & R & N & R & E & L & M G A & T & N & P \\ N & E & A & A & P & G & T \\ N & A & L & K & N & R & E & M & M & A & T & D \\ P & S & K & A & D & P & G & T\end{array}$




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49

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## 

## T



 $Q L T L A G A Q Q E F Y$ G V H S E R P F F G E L C E F M I S E P I V V Q V L E G E Q M T D E M IDD N WY A H H K E K S F F K D L K T F M S WT P I V A M V WE G L R M D R A K A E GF Y A V H K E R P F F G E L V D Y M I S G P V V V S C L G $\quad$ D HASE E LLLQD HYF E HK EK H F F K D L L E Y M Q S D L V V A MVV E G D




 RMSEA QAKTFYEVHAERPFYGELVSFMTSGPVVVQVLEGE

 YP S E E L VM E H Y K E H E GK D F Y E R L I N F LT S G MVI A I V L E G K


 QLS R D E A E R H Y S V H K G K P F F A D L V N Y NA T R E M LAA A HY A E H E GK P F Y E P L V E F MM S G P V V A A I F E G E

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 H M S K A T A Q G F Y A V H K A R P F F D S L C T F M S S S G P C V V L V L Q G D

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90
N A V|ARHRE|LLGATNPKDAAPGT\||RADFAES\|IEANAAHGS D GAI AKNREVMGATDPEKAEEGTIVKDFAVSMQQNSVHGSD




 D GI A AVVRRLVGPTDTRESLPGTI RGDYAYRTRLNIVHASD G A I A K N R E L M G A T N P K D A A P G T I R A D F A E S DAI AKNREIMGPTDPENAPAGSI

 RAI A A WRQLA GGTDPDKAAPGSI
 NAV EKNRELMMGATDPKVAKPGTI RADFADNDDANVVVGGS NAI KAVRGQMGATNPLDSVPGSI R NAVARNREDMGATNPQEAAPGTI
 DIVKHGRAMLFGDDDLNPMPGTI
 NAV QRNRD M GATNPANAEPRGTI
 GAIINAYRKLMGATDYKQAAKGTI RHDYAADVGENAVGS D NAVLKNRELMGATNPQEAAPGTIRADFAASIIDANAVHGSD NII I L A RVMMGATKPEEAQPGTIVRDYAVTISENVI HGSD
 $N V I D D G V R K M A G K T N P D L E S A D L G T I$
 NAI SVVRKMIGKTNPAEAEPGTI

 NAVEKMRKAMGATDPAKSEKGTV | $D$ | $A$ | $I$ | $K$ | $K$ | $Y$ | $R$ | $D$ | $V$ | $L$ | $G A$ | $T$ | $D$ | $P$ | $A$ | $K$ | $A$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| $N$ | $P$ | $G$ | $T$ | 1 |  |  |  |  |  |  |  |  |  |  |  |  |
| $N$ | $A$ | $A$ | $E$ | $W$ | $R$ | $E$ | $L$ | $M$ | $G$ | $A$ | $T$ | $N$ | $P$ | $E$ | $D$ | $A$ |
| $E$ | $E$ | $N$ | $T$ |  |  |  |  |  |  |  |  |  |  |  |  |  |

 \begin{tabular}{l|lllllllllllllll}
$N$ \& $A$ \& $A$ \& $K$ \& $Y$ \& $R$ \& $E$ \& $V$ \& $M$ \& $G$ \& $A$ \& $T$ \& $D$ \& $P$ \& $K$ \& $E$

$A$ 

$R V V$ \& $D$ \& $S R T M M G V T N P K D A G P G T$ <br>
$N$ \& $A$ \& $A$ <br>
\hline
\end{tabular}




 NAI VKNRGIMGATNPDDAQEGTI NA A QA YR DLMGATNPANAAEGTI DAVSTYRKIMGATNPADAEEGTI $\begin{array}{ll}D A T R Q V R G M V G E T D P A E S P A G T \\ N A & L \\ N & R E M G A T N P S E A D A G T\end{array}$ | $E$ | $P$ | $A$ | $V$ | $R$ | $K$ | $V$ | $G$ | $S$ | $T$ | $K$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| $G$ | $R$ | $E$ | $A$ | $E$ | $A$ | $G$ | $S$ |  |  |  |
| $N$ | $A$ | $E$ | $R$ | $Y$ | $R$ | $A$ | $L$ | $M$ | $G$ | $A$ |








 NAI S RNRQIMGATDPSKAEAGTI



 $R$ VI E GFRS L A G T T DP T T A AP G T I RVI E GFRALAGATDPTAALPGTI | $N$ | $A$ | $K$ | $K$ | $N$ | $R$ | $E$ | $L$ | $M$ | $G$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |$A$ D AV SVVRA LVGKTNGREAAA GT I

 NAIA DYRNLMGATDPAKAQPGTIRALVYGKNIIEENAVVHGSD $N V V K T G R Q M L G E T N P A D S K P G T I R G D F C D Q V G R N V M H G S D$ RVI E GFRSLAGATDPTTAAPGTL RGDLGRDWGERVVHGSD
 NAI SRYREI MGATNPANAAEGTIIRKKYAVSLDEANSVVGGS

| UniRef90 B2I7V6 | $\text { S VIENAA } \square \mathrm{E}\|\mathrm{~V}\| \mathrm{A} \text { Y F F A A T E } \\| \square^{140} \quad 150$ |
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| UniRef90_V9HLH5 |  |
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M

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## \section*{} <br> 

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    \(T L_{L}^{L}\)
    
MEM
MS
SK
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$M 1 K P D A V A D G N S G A D L K M D E E A G F R I V A L K L T$
$E K T \| V I V K P D G V Q R S L V G E V V K R L E M K G L K V V G L K M M$
CTCCII K P H A I S E GLL GKI L MA I R DA GFEI SAMQMF

K M


STL
S G


            E R T \(\quad\) L \begin{tabular}{llllllllllll}
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T $\square \mathrm{E}$
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 K S I A S WRELDGPTNPFKAHPGCLIRDTYGKDQTRNAVHGSD
 D A I C EWK RLL GPANSVVARPGSI RADF GAD GI RNAA H GP D EPPKVFRDFCGPSDPE\|ARPGTMRAVFGKNKIQNAVHCTD

 NAVQAWRDLIGPTDVEMAFPRSLRALYGTSKTYNACHGSS D SVV P F R QLSGPMDPELARPDSLRARFGLNTDKNGVHCS D H A V D H WR N MLGGPA S LRRQA A P D S L R A Q F A N D S D I N QL H G S S | $N$ | $A$ | $V$ | $E$ | $E$ | $W$ | $R$ | $Q$ | $L$ | $M$ | $G$ | $P$ | $D$ | $D$ | $E$ | $E$ | $A$ | $C$ | $P$ | $E$ | $S$ | 1 | $R$ | $A$ | $Q$ | $F$ | $A$ | $H$ | $D$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| $H$ | $L$ | $S$ | $N$ | $A$ | $V$ | $H$ | $G$ | $S$ | $S$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| $H$ | $S$ | $K$ | $\square$ | $C$ | $R$ | $K$ | $V$ | $T$ | $G$ | $G$ | $T$ | $E$ | $P$ | $L$ | $T$ | $S$ | $D$ | $V$ | $G$ | $S$ | $I$ | $R$ | $G$ | $D$ | $Y$ | $Q$ | $M$ | $S$ |
| $D$ | $V$ | $R$ | $N$ | $L$ | $I$ | $H$ | $A$ | $S$ | $G$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |

 RAI A TWK GLLMGPSDSTQAYPECLRARFGTS DLRRNAVHGSE NAI E E WRNTIGPVVDPDVARPNSLRAQFARSVLENALHGSS $N A I S D L R E L I G P Q N A F K A K P E S S L R A D Y G T D D Q Q N G I H A S D$
 KAI E H WRKII GPTNSAKARPTSIIRA LYGTDNYKNAVMGSD

 GAVKTGRVMLGETNPADSKPGTIRGDFClDVSK
DAVATVRKIVGSTYPNEAAAGTVRGDFSHYSRANLVHASG
 N MV A TSRAMVGMTDSTQAPAGTDRGDMGTSSMKNDCHASD

 NVVK GRVMLGETNPADSKPGTIRRDFCIQVGRNIIHGSD
DVVVT GR RI G A T R P WE A A P G T I R G D H A V E V G R NVI H G S D

| UniRef90_K0YXQ3 | S P T S A E R E I K I |
| :--- | :--- | :--- |
| Un W |  |

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UniRef90 J9HLG9 UniRef90-R1EUW1 UniRef90-W7XGD1 UniRef90_UPI0003C28E4F UniRef90 UPI00042B9894 UniRef90-V4AV25 UniRef90-A0A024UW35 UniRef90-W4Z1L5 UniRef90-K2E3D9 UniRef90-K7IMG6 UniRef90_UPI00032B1153 UniRef90_UPI0003C16995 UniRef90-H2SKK6 UniRef90-НЗВ3С0 UniRef90_UPIO00393E3EE UniRef90_UPI0003292262 UniRef90-H2ZDH8 UniRef90-K2E4B7 UniRef90-W4H978 UniRef90_U3IGC3 UniRef90-W4XUI8 UniRef90-UPIO00265460C UniRef90-K2A692 UniRef90-T2M335 UniRef90-H2RMP2
UniRef90- X1MED7 UniRef90-H2TP58 UniRef90_S4R4Z0
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VVVGPD GLS G K VVD DV L K R I K D E G F V V TV A I L K P D A V A H G KV N E I MM K I Q N A G F QI LA QE ER



> QQTLAVIKPDAME EHREA ILEEDRGRGFSVTRLKET

$\qquad$
E K

NLEK TLAL $\operatorname{L}$ R P SLLLKERK NS
TKKEKTLGLI KPDAI NKMGSILKRIIKDDGFTIAMMQRE I


E K T L GL I K P D A I S K
$\square \mathrm{Q}$


S K K ERTLALIKPDAVTK

K I Q E A GF Q


GEVLEKI YASNLIVTHKAKMT Q VE Q T L AV I K P D A MDEK | $L$ | $G$ | $A$ | $F$ | $Q$ | $M$ | $V$ | $Y$ | $D$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| $A$ | $G$ | $L$ | $T$ | $T$ | $K$ | $A$ | $K$ | $M$ |





## E K

$\qquad$

 ERTLALI R PDVARERNAEEILSQIGKAGFTVALQREV


 Q Q T LAMV I K P D A L L R R E H K E T M L | $F$ | $S$ | $V$ | $V$ | $Q$ | $S$ | $K$ | $E$ | $M$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| $M$ | $R$ | $V$ | $V$ | $A$ | $A$ | $K$ | $M$ |  |
| $V$ |  |  |  |  |  |  |  |  |

CTLATIK P DAVMHGNSEEVVMKDQETGFDILTNEER L Q L T L A L I K P D A V A H P L I L L E V H Q Q: L N N K F L I V R M R E L V E R T L A A L L V E N T M A L I 1 E R T V A V L

 A EK K
MK ML
$\square G \square I K P D D V S E G R V E E \prod K S Q \square T N A G F S \prod E A E E E R$ MI K P D G V Q R G L V G E D DK R FE Q K G Y K L DA M K L T

 TVAI I K P DVVAHGKADEI DMKI QD GFVI LA HEER IE R T L S I I K P D A V A K N V I G Q I Y A R F E G A G L K V V A A K M I
 ERTYMVKPDGVQRGLVSEVIRRFEQRGYKLVALKMK

 ERTLALLKPDCVASGAAGEVEALIAEHGFEVVRRRRW $K$ K E K I L V L L L

 C T C C I I K P H A I N E GLLT GKK I K T $\square$ F D A GF E I SA LAQLF

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UniRef90-|1GHX0
UniRef90-J9HLG9
UniRef90-R1EUW1 UniRef90-W7XGD1 UniRef90_UPI0003C28E4F UniRef90-UPI00042B9894 UniRef90-V4AV25 UniRef90-A0A024UW35 UniRef90-W4Z1L5 UniRef90-K2E3D9 UniRef90-K7IMG6 UniRef90_UPI00032B1153 UniRef90_UPI0003C16995 UniRef90-H2SKK6 UniRef90-H3B3C0 UniRef90_UPIO00393E3EE UniRef90-UPI0003292262 UniRef90-H2ZDH8 UniRef90-K2E4B7 UniRef90-W4H978 UniRef90_U3IGC3 UniRef90-W4XUI8 UniRef90_UPIO00265460C UniRef90-K2A692 UniRef90-T2M335 UniRef90-H2RMP2 UniRef90-X1MED7 UniRef90-H2TP58 UniRef90_S4R4Z0
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UniRef90_R7T472
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UniRef90_UPIOO03332292 UniRef90_Q1JTK8 UniRef90-UPIO00258EA51 UniRef90-T2M930 UniRef90-R5N865 UniRef90-R1FSL1 UniRef90_UPIO0035A23BB UniRef90_UPI000443981C UniRef90_UPI000298AD4F UniRef90_A2DJE8 UniRef90_UPIO00274580A UniRef90_U6I176 UniRef90_K7F2E6 UniRef90-T1AWC0

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 QAI S Y WKELLGPSNS LRAHPESLRALYGTD DLRNALHGSK N AV S Y WK ELL G P S N S IVKA H P H S L R A I Y G T D E L R N A L H G S L

 D A V E G WR Q A I G P T D P E K A K P E S L R A $\square Y G$ G K DV L H N AVVH G S S D A I LKVVREITCGKTNPE EADSVVSIMRAYGRVTETENVMHASA R A V GA F R Q I A G G T D P D K A A P G S I R G D F A L EVQ E NIVVH G S D GAVHHWKNI LGPADVNKARPECLRAQFAVENEPNQLHGSE NAI SHWKELIGGANSLLKAQPDSSLRADYGTD DLRNALHGS Y
 QSI KA WRGLLMGPTNSNRAHPWTLRARFGDDGTRNATHGSD GAI AVIRRKTMGATDPAKAKAGTIRGDFGVDVEHNSVHGSD
 $G V V T T G R K I G A T N P A E S A P G T I R G M S S T E A T R S R A Q G$ GAVEHWRNLLGPKDPKQAQPESLIRAQFGESECVNQLHGSQ
 DAI ATARRMMGKTNPDEADPGTIVADYADHTNRNVIMGSD NAVEEWRAMMGPTDPNKARPESLRARFAADILHNSVHGSS

 SAI GAWRSLIGPTSTETARPRSLIRAMYGSDNTRNAAHGSD NAIK R WRELLGBTNTVVARPNSIMRGLFGTD GTRNACHGSD Q A A D D WK E L S E S D E NE S Q K L S R L G E V L E N E N M L NVVC DV Q D





 E AVVSLWRKVLGBTDDSSAARPESDIRAKF GTDGTKNAAAGGSD
 N A V D K L
 NAVQEWR QLMGPTDPEVAKPESIVALFAQD\|LSNAVHGSS $G \boxtimes V S D \square R N L L G P K D \square E V A K P E S L R A Q F G T D K K M N A M H G A D$ D AV D GWRT L M G P T D P D Y A R P E S L R A L L G K DVVMQNAVVGGS S HSVDMVRKIVGNTLPSKADMGTVRGDFSVDSIRNVVHASG NTPTKFREFVGPSDPEIARPHTLRAQFGVDKVKNAVHCTD DAVA HWNHLI GPANSVVIAHPDSLRAKYGTSELQNALHGSE NATQK $\square R D \| A G N T D P Q K A E Q K T L R R L Y A Q$
NAVE E WRAMMGPADPE QACPNSMRARFASN LIHNSLHGAS DAVR GWRELMMGTKVFRAVPESDRACYGLT DTRNTMTHGS D NAI SHWRMLVGGPTSVTEAHPDSLRALYGTDQQRNGLHGS D DAVDGWRSLI GPTDPEKAKPESLRAQFGKDTLANAVHGSS NAIA EWRELIGPTDPSKAHPQTLRAKWGDDTQRNCVHGSD NAVEEWRAMMGPADPEQACPNSMRARFASNVLHNSLHGAS D A I A K N R E L M G A T D P K K A D A G T l



 D AVAKWREMLGPASVNEAKPESLGRAQFTMTSGKNLLHGSA G A E G V I E D L $N \vee \vee Q R$
 DAI PNWKELDEMSKSVDEAEKPDRGEETVETGANVCDMED NVVLE GRKMLGATMPAE SACGTDRGDGCIEVSK
 D A V S A WR DL I GP T D SVVKARPK S I R A L

 SAVAKNRDLMGAN
N A I E E WK R 1 IGPT TVTEAFP DS $\square R$ A K Y GRD D L K N A $\square H G S Y$ Q
DVVK Q GRRMLGETRPLESNP GKHPSRRLLH N AI T R W $\square E V I$ GP E D S E E A R P S S L R A C Y GK D K $\square H N A \square H G S E$
 GAI GAWRELLGPGDPAAARPRSVRARWGRDKQANAAHGSD CVAEAFREACGPADPRHLRPRTVRALFGMNKVQNGVHCTD


 Q AVK



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UniRef90 НЗА143

## 由のにのに

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NSEDAEREI K LLWFKPDE I DE
SAESAAREI ELWFP
SHKEAKREI DFFFPKQ
CFTSAEREI RFMFPE
S IE EASRQLAFFFPDF




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CTLC\IKPDADKAGLSAK I Q D DLSEGLE ITGLGLY LE ETVALI KPHAYQD LREE RFKS LVSQCLTCKE C LQLTLAALKKPHVVK SPFVQK RRDLIDDNNFKVVRSRRT
 KLEKILALL RPDLFQ EKKENVLGII HDEGFKI LMQRPI
 SHKEKTLGLI KPDVISK QLGDIVNSLRDHNDQLSKAKMD


## EDTFALIKPH VTHEQRGETLKLIKDEGFELTQLKEL

 $K$ VEKILALLRPALFHERKEAVLNUIEDEGFKILMQRQI

T
NIERSLVLIKPDAMQRGLAAGIITHLEKQGLKMVALKML
T
1 E
D L

$\qquad$ T

$\begin{array}{llllllllll}R & L & A & A & I & K & P & D & L & L \\ R & K & D \\ R & L & V & L & I & K & P & D & G & I \\ E & R\end{array}$


C T C C V i

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 VEKT LAAI K P DVVDKE EEDQD IILRSGFTIVQRRKL
 E C S VI L CK P D A V E R GLLVA E V L D R V R A A G F T V S D P V D V QTTLALIRPDLCQANK EDVLNVIQEGGFKIMMQQQI
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 TCVVVVK P H A I D D G Q A GY I I E CI Q NLGYR I S A Y L M C

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 $C T L C D V K P H A I L E G G L G A A L E A D D K G G F N V T A L Y M L$ R TVAA I I K N H A L L N H R F

VE K T LAALIK P DVVD K E E E I Q D II LG S GF TII Q R R K L

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 QLSRAEATDFHADHLSKPYYNELLEF DTSGPVVVAHVVWEGE Q L S R A E A T D F H A D H L S K P Y Y N E L L E F DT S GPDVVA L E L L G D AMREAEMXLFYQHRDGEEAFKKLVHHMCSGPSHLLNLTRE
 TLLTE E EVVK I I Y K DQQFRRDFFPD LV QHLLTS GP LMALCCL E RE $N A S K D L L N E H Y Q D L A D K P F F G G L V A Y M S S G P V I A M V W E G L$
 TLLTHELAD KI YP K ITTGKDFYNDVLEMLSEGPSMVMILTKW QAPES LLA EHYHDLQRKPFYPALI SYMSSGPVVAMVWEGP V L S K E E A Q T L C K E Y E N E D Y F E NVVI E N M T S G P SL A L V L V R E
 MS R A E A E G F Y A AVMHS E R P F F K K A L V V
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 VLSKEEA QIVGKEYENEHYFENLI ENMTSGPSLALVLLGD
 VLT E E QVR DF Y S R $\square$ A D QP DF E DFVVSFMTSSS L HI LVVVSQG $M K A A P V V A M C V E G C$

 LLDDP E E A D K DY S K DK GK D F Y K DV L E V L S E GP S V VM I L T K W



 QLTTVA DA E DF LDEVV N G VVP E Y K K L V E Q M S S G P C WA V E V C A E










 EL T A E EAE EFY QD HVK QNYYEKGVRWLSSSKLCALVLEKE
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 HLLTP E QV A E DY E Q H Y G CP S F P N M V V S M S L G P CL V L S L A G M
 RFSENDVVAFYE E H DGK AFFPNL A A MVMA GPACVI ELIGP



 TMTEAEMRLFY QHRA GEDAFEKLVVYMCS GP SHLLLLL TR T

 | $M$ | $V$ | $E$ | $G R$ |
| :--- | :--- | :--- | :--- |
| $M$ | $V$ | $F$ | G E |



 LVK VMTCSPVVAMVWEGN HLSPEHCSNFYVE QY GKMFFPNLTAYMSSGPLVAMMLARH

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$N V S R S L A E P G A L R A V D A P V P C A A S G D$
$N G L Q H W K E L I G P S T V E E A H P N S L C V Q F A M S D P D N Q L Y G S D$ KATDTDPLWKELHEISESMPGEAEAEETDSMSKNICDVPD EATARWRSLMGPTKVYRARPESTRGTLGLTDTRNTVHGSD NAVAKHRELMGATNPKEAAPGTIRADFADSDDANAVHGSD NAQAQFRQLCGPPDPEMARPHTLRAKYGKNKDENGVHCTD
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EAVQVVRT门TGATNGRKAAPGTTRGDYSMSGQENIVHASD NAIADWRRLMGPVDPEEAKPDSIRARFGINILKNGVHGAT NGLRHWRQLDGPSTVKEAEEYFPQSLCVQFAMEGLLHGSD DAISRWRELMGPTKTFRAQFTHTRGQFGLTDTRNCTHGTD DA: PEWRRLMGPTDPEEARPDSIRAQYGVSMLLRNAVHGSS NGLKHWKELLGPKSVEEAREYAPESLCAQFARENLLYGSD
 EAVKVVRKAMGATDP
N SVEYWKQLMGPKTVEEAYPESLCAQFATNFPTNQFYGSS HSVERWKELIGPKTVEEAHPDSLCVRFASNFPVNQFYGSS DAI RQWRELMGPTKVFRARPDSIRAQFGLTDTRNTAHG NAVPADRAVCGPQDPEVCHPHTLRSMYGVDRDRNAVHCTD MRYSVTAICLAP PIVRTRWRVR
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| UniRef90_F7FVB7 | S P E A E K E $~$ R S F F P L E H TVA $\square$ K P H A Y Q D L R E |
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| UniRef90-UPI000333B02E | SLEVVAKEI Q F F FPPQNTVDYEKKD |
| UniRef90-UPI00046B39E2 | SVFEAAENINALIFAES |
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Y D R F E R A G X R DA A X R ML
 $T V L L L K P R D W S S A L L G K D L G K V Y Q Q K F I L V G M K P V$ TLLA I I K P E ATMAQLA A T L E R E F GI S $R \square Q L T L A D I K P D \square S \square Y P F R L R V R H L L L K N D F L I V R S R K M$





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UniRef90_M7CIP2 UniRef90-DONBP2 UniRef90 UPI000477D76F UniRef90-UPI0003743303 UniRef90-UPIO0035F3E7B ero
niRef90 UPI0003837D5A UniRef90_UPI00045DCAF8 UniRefgo J91AQ
UniRef90-I3IHN0
UniRef90-FORZF8
UniRef9- X8Z9Q
UniRef90_UPI00034F9815

- 90 LIJC47

V
UniRef90-T1JFH3
UniRef90 ${ }^{-}$X1G0L6
UniRef90_K2BZH8
UniRef90-GOROJO UniRef90_UPI0003EE005A

100
110
120
NVHSEFRMFCGPSDSD\|ARPATLLRARFGCDKYKNAVVHCTD DADGVMKRLVGRTDPQAAGKGTIVRGDFSSDSVMNDIHASD TSPPIDKKLKGHYRPEYAKDGTI RSRFGDNRIFNLFHSSD RSVEVWNALMGNRDPEVARPNSLRALYGVSLKQNALMGSP NAVKKLMDMLGPEDPKVAQPCLWRAQYGUDAVQNGFYGSE NAVKKLMDLLGPEDPQLAQPCLWRAQYGTSTVQNGFYGSR TAVQHFRESCGPWDVNFARPQTIRAKFGDDGVKNAVHCTD

 NAVKKLVDLLGPEDPTLAQPCFWRAQYGASSVQNGFYGSR $D V V A Q F R A S A G P W D V E M A K P Q T V R A K F G H D R V R N A V H C T D$ YEGSDDRTLAGKTNPQDAGADTI RGKYSDDSYENVI HASD GTPEHVRGLLGHYDPARAAPDSIIRGHFGADSIAQAIHTSD DTPERVRALLGHYDPSRAAPDTIRGHYGTDSMANLIHTSD
A ASVVSGRLKGDFRPSRAQPGTMRADLGANPDVNLVHSACD
 NAVMRLWELMGPEDPDEARTVDWRAQFGSDRLHNGIYGSR NAVAEWRRLMGPTDPEEAKPDSMRARFGMSKLKNVVHGAS NAI GYWRELCGPTNSMKAHPDSURGMYGIDDQQN $N V V Q G L K A V C G C Y D P$
DAI RKI RNI LGPTDSKKGEPGKVRRDYGEDDMKNAAHASD DAVQKIRTILGATDPNKATAGTIRREFGSNIRVNAAHASD DAVQKNRDLMGATDPARAC
RSASALSALAGPDDPRRSDPECLRADYGANRWENGVWTTK GAVLKLLDLLGPEDPKAQDQF LWRAQYGTSAVQNGMYGST NAVAVLLKLCGDEDPSLAREFALRAMFGLSATANGLHASR DAVVKIRDVLGPTDPTKAPGGTVRREFGSNVMVNTAHASD HAI AYWKELIGPSNS $\square$ RAHPDSLRA $\square Y G T D D L R N A V H G S \square$ DAI GAWRRLMGP
NGLQHWKELIGPSTVGEAKQYFP NAITKDRKVLGETNPKEAAPGTVRKDFGLDIIKNGAHASD NAQGIMQTVKGNDDPDGNPVDAIQSGGRLYVFKNGIHVSD NPDEKWKRI DGKSDPE IAN MNCLRADYGKNI DKNELHGSD R L MLVWEKI

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130
LP E D T R L E L E Y F F
AANAENEIKL
CTKAAKREASLLFTAEET
DPEGARREYGTWYGPGQUNDFDRC

SYAMAVRDVKLFFPE
LPDDGESEVRYFF EILN
PVTGAQRESRLWFQSNELVDWAAH DHP AANKK $\square$ NLYFTKKELVS
CSEDGMLESYFFF
SYHAAVRDMK LFFPE
LAEDGVLECQYFF
SHESAQRELDNW
DAEGSEREFNVWFGP
DAPGTAREFGVWFGPDN
DPASVRRESRVLFDDGPVFD
SAASATLE I AVVFGADEV
SYRNAl EDMKLLFPS
NAYEAKEVIN RLLFNPEE

A VE NAERE DG L W E NKEPYE LKK E DE DY L SAENAQREMGVL KAEENFCYSLISE

DEASAERVVSEDFSGSP\ME
SYQAA DRDI QTFFPEGLLFE
SFLSAQEEURSIFPE
SPENAKREMKUL K I E ENNCGRVLKDY


DGMTSERERD ${ }_{\text {F }}^{\text {F }}$



[^0]:    

[^1]:    lupas_21
    lupas_14
    lupas_28
    jnet
    conf
    sol25
    sol5
    sol0
    jhmm
    jpssm

[^2]:    lupas_21
    lupas_14
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    jhmm
    jpssm

