

## QUERY

## UniRef90_G5H5K9

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UniRef90_A5M8L7 UniRef90_UP10002FAEA5F UniRef90_G1WHZ1 UniRef90_R5GXLO
UniRef90- XOTL67 UniRef90-I6RYL4 UniRef90-G7RZW4 UniRef90-U2TCE4 UniRef90_A9HPK3
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UniRef90_SOFMN6

A A DDGTGEQ ${ }^{50} K D G K P \nabla V K P V^{60}$
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A A S L D GS GE Q T K D G K P A M K P F WE I T E A E A E K C L K A T T WY P A

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 S AL DF C GEK K EN GNGlVVKPFWEMT ATLDGT GQEMLD GKPAMKPYWE IEDKD DKACTDATTWNAA TTLD GS GQS DAEGKPVVK QAWEMTEK DVVEACCLIKATTWWYPA ATLDGTGKQKYEGEPVMKPHWEISDEEVEACLAATTWHPA
 A ALD GTGHQQ $\square$ D GKPA $\square$ K P F WEV SA L D A T GEMYKD GIP A MKP F WE I
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 A T L D A T G D Q $\square R D$ G K R V M K P Y WE V T T T L D A T A QQQDEEGNPTMKPFWD A A L D GA GREVN E GVDPVIKP FWE I AT LD GCCQKDEQGHPTMKPHWEI
 A ALDGTGRQSKDGKPAMKPFWDIT T TMDGCGQSDAE GQPVMKPFWEMT
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 SALDAS GRQTRNGQPAMKPYWEDTPYEAQACLDATRWCPA
 A $\quad$ L D A T A Q Q T A A G R P DMK P F WE I T E G E A Q K C L D A T L WP Q A A CLL D A C G E C D EN G N A VVMK K WW EVV T D E D
 ATLDGTGCQRVDGKPAMKPFWEITE E EVVEKCLEATRWCPA A ALDGTGACDENGTAVMKPWWEVSTDDIQAMLAATDWAPA A T L D GT GE QK LDDGQPAMKPFWDITEE EAAA CLLAATSWRPA A A L DAT GASDENGNHVMKKWYDMTDAD DDGCLKATDWCHA A ALD GT GEQE $\square D G K P A M K P W W Q V T P E E A Q R C L S A T K W R A A$ CTLLDATGEQTKYGKPVMKPFWEIWEEEVNAMLKATKWGPS A A L D G C M A C T K D G Q P V M K K H W DVVT P D DAAK TMM M A A V WC P A CTI DACGEKDKNGNAVMKPFWEMTEADI DACLKKATEWCTS

 A A L D GTGAATRD GKPVI KPFWEMTEQD DES CLLESVT WCPA A T L D F A GVVQKK D GEAVVKPFWEDT DE ECQQEME KT S WYP A A ALD GTGQHDET GNHLIVKVWEITEEDGKRCLAHTRWCAA T TMDAT GQKDGEGNSVMKHFWEITEKDVKAVLDNTTESVA T A LDDCT GAKDENGNGTQKEWWNMTDE DI QNCIDDAT D WCRA A ALD GT GEQDPD GNPTMKPYWE IS DQEVSACNQATTWHPS SAI DYTAQVDKDGNRC $\quad$ KT T Y EMTQEDADNCLKATTWNPA VA L D S C GE Q Q $\square D G Q P A M K P H W D \square S Q D E V D R C L Q A T T W H A A$

 T T L D G T F A A V V E G Q H V I K P WW E V T E E D Q K A M L E G T T F H P A
 A A L D WT GEQTVDDGK A I K P WWEVVEEEAKRCLEATRWCPA S A L D F S G V C T A D G K P E I K K W W W E I TME E D Q K K I L E A T T WN P A QTLDATGEEKVDGNPAIKPFWQVTEDEAKRSLEKTRWDPA SCLD A CGEKNDKGEACMKPWYDVTDEDVDKMLKKATTWNAA T T L D A T GMKDA DGNPCTKKPFWEVTEE DQKAVVEATTAFAA TALDGSGAKNAKGENCMKPFWQMTEA DVKACLSATAWCAA CTLDGTGQATRDGKPVMKPFWEM TCLDATGAKDEDGNSIMMHFWDLK T ALLDAS GARDERGESIMKPFWEMMD DEDI KAMLLDNTD WCQA A A D A GAR D A A A A D WCRA A A L D G T G A K D E K GE SVVMK P WWDMT DK DI D A C L K A T T WP C A
 S ALD GAA ARDEA GQPTMKEFWNMTE EDIKACLEATDWCRA S CLD GAAAKNEAGEGCMKEWWKLDAEDMHALLEKTDDCRA A A L D GSCARDA QGAPTVKPHWQVDED DVSALLAAATD WCP G A A L D G T G E Q QRD G QPVL K P H WE IV D S EVVR R CL D A V S WR P T A A L D G S M A S DVVDGHPGI K P A WEMTETMD S R C L D A TAR WGA G


## QUERY

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UniRef90 U2UMD9 UniRef90-R6K2W4 UniRef90-T2KQP3 UniRef90-G2Z8W1 UniRef90-B6VMR0 UniRef90_UPI000475DA1A UniRef90_-16YW57 UniRef90-D3EM02 UniRef90-R5B6X2 UniRef90-S6EZD1 UniRef90-D5EY63 UniRef90-F8F285 UniRef90_GOGEM3 UniRef90-E9S7C9 UniRef90_UPI00046F3C1E UniRef90_UPI000479853C UniRef90_R5C412 UniRef90_UPI000479F4BC UniRef90_B5JDH6 UniRef90-R5ICG7 UniRef90-C0EDQ6 UniRef90-F5SB52 UniRef90_U2KD09
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UniRet90_E2ZFZO
UniRef90 SOFMN6




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$K ~ N$ $\begin{array}{llll}A & F & K & D \\ A & H & A & S\end{array}$ D E V R D I I E G R T D P L

 \begin{tabular}{lllllll}
$T$ \& $T$ \& $W$ \& $F$ \& $V$ \& $P$ \& $R$ <br>
$T$ \& $T$ \& $W$ \& $F$ \& $A$ \& $P$ \& $T$ <br>
\hline \& $T$ \& $T$ \& $W$ \& $F$ \& $V$ \& $P$ <br>
\hline

 $\begin{array}{lllll}C & T & W & F & A \\ P & R & V & D \\ T & T & F & F & A \\ P & R & L\end{array}$ 

$P$ \& $T$ \& $T$ \& $F$ \& $F$ \& $A$ \& $P$ \& $R$ <br>
$P$ \& $S$ \& $T$ \& $F$ \& $F$ \& $A$ \& $P$ \& $L$ <br>
\hline
\end{tabular}

GKGA
GPFK
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$\begin{array}{ll}T & S \\ K & N \\ T & S \\ R & S\end{array}$EGHTVTLLDETHRTLDERTDRTWPTTWC G

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\begin{array}{lllll}
G & S A & F & C & T \\
K & G & F & T & S \\
G & G & A & F & K \\
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\end{array}
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P C T WF T P I C

## QUERY UniRef90_G5H5K9 UniRef90 W6TRI6 UniRef90 B5YC67

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UniRef90- XOTL67 UniRef90-I6RYL4 UniRef90_G7RZW4 UniRef90_U2TCE4
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DE

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 AQLFSDLRTFWSTDSIKRVADYDLQAKKADGLI LLLNSGS AQUFCDVRTYWSPDAIKKATQKDVSSLIPTGAIHLLNSGS QFSSDVRTYWSPEAVRRVTGHELTGRAAGGFVDFRNSGA







A VMFHDVR WGAEY
$\square$ FADVNHL
SVMFHDVR WGAE
FSDपR YWDP
$\square$ FADVNHL
SVMFHDVR WGAE
FSDपR YWDP
$\square$ FADVNHL
SVMFHDVR WGAE
FSDपR YWDP


DDDVVITANCGA DMEQGVLRLVNCGT DDEILLI ANCGA $\begin{array}{llll}D D E L V L I A N C G A \\ G R W L W V & A N S G S\end{array}$ HE GRWLLWVLLNSGS DDDLVLIVANGA

Y WF VNS GA

A G S

[^2]

 QT DDATGAEKDDGQPAMKPFWDLTDEEVQANDDQTDWVPA TTLDATGQQKEDGKSTMKPFWEITEEEVKRCLENTKWSPA

| H A |
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| H A |
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| A AFDWAAEPGAS |
| H T |
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ASFNWA
QS
MCTYYA GRC
QS
SS L

CGGYAMSGDP

| S S L |  |
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| A AT | T L A R D K |
| S S L |  |
| S S L |  |
| C G | A F NQDV |
| S S L |  |

LA P Y F A Y GRHDS LEGA

T WY A A R S D D P A V N L A R VHLLY P TYFAGGSFDYRDNLPKVVFYPE T WY A A R S D D P A E N L R R V $V$ H L $L$ Y P
 TWFAARSDDPAENLARVVTHP T WY A A A S D D F K E N L L A K V T T L Y P P A TWFAERSDDPAENLLRRVHLYPE TWYAARSDDPAENMRHVVFLP

 TYFAGKSNKPE ENLLKNVII HPE

P WF A T R T DDVES NLDDKI SLDKQ
TWFAERSADAAENMAKVVQL LP G
SWY A S Q S S DP S ENFKK V T L Y P A MATFYAARSCNLEENLKRVSLYP Q
 TYYATGTDDHRENLGRTTLYPA T WF A ERSEN HVDNLLKKV T L $L$ YYARMSDDPEENLLKATTI Q GQ
 A T L K GVVHSYRQ Y Y A A L S D D P E E NLLDR R V $K D V D W G L Q$ Y Y A G L S D D P Y E NLLKRTM IQ GQ Y Y A KLLS EN P E ENL K ETET LQ G Q L N GVH SYR Q

EGTSSQRQQ TYMY $\square$ A GATGYRQ

EKSSLLMSA

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\|LE Y F LGDGLS SHYRTPGEDPMTAYRYNVI MPFTMTRVNIIKGNMDCSVV NVEYFRGGGFSSHYLTKGGMPVTMTRINLVKGVGPVLQLA



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VY SVMANWGANHGVTVYGHVGADLDTLASMDRDPVTLHNV TFLSRFGANHIHAVPGDRRAELRTVCELMGISLD
IE
RFI EKFPCNHI HAVYGDYVNELDMVCEILGIPWEI
EFLSRFGANHIHAVPGDYVAELKQVCRLLDVDFE
HSVFLDRFDANHCHAVYGDHVDDLKMICSMLNVEVEM
DFLSRFGANHTHAVPGDHRATVRAACDLLGVTVD
TFLARFGANHIHAVPGDRRPELREVCELLGITLD
H
IFLDRFGANHCHAVYGDHVDELKMICKMLDV DVVMYN
EVLSRYGSNHIHAVPGDRVQELRAVCELLDID
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PANHFHMTWNLPVARL
HPCNHDQVAEDCVAGLVEDCEKLGDKVQDKNL ADELLARYGSNHI HALPGDHVDALRETSRLLDVEFD FNEFI SVFPCNHILGVAGDYLDSLIVGCEISGI PV Q T LI E T

PANHFHMTWDLEPARLMDFCNVLSV
HQ $\quad$ F LDKFDANHCHAVYGDKTEELKMICEMLDIDVEI
$\square$ KKFVDRFNSNH $\square$ S GVA GHYKKELLDLCSLLDI E KLI DVWPANHFHAVS GNY RQDFINVMGANHLSLVPGDYTQELKFVAKVWGI KVDLNDL Y EKFYEKINSNHLHMVYGKYAKVLKMFCEIVGIE RETLMAHYLSNHDAVAYGDILEEMAADSQTLGFRV YP PANHFHMTWNDPVARLQHWMDLTGT AQDFDQVMGANHDSLVPGDFTEEI KFVAKI WGIPDDLTDP PKEFLQNIRSNHLHMAYGNFKSELLDFCEILGI K

 KEKFLNAMGANHLSAVSGDFRSEIEFSARLWGIR QMMGRHKANHIHVVYAENEAEANKLCRILGQQV
 QMMARHKANHIQVVYANHKACRIKAAALLGGIQVHFGNV VDDWSACGPTHHCAMGRGYHAQAFHKLGQLLKI PV RDQMMARHKANH $\square$ QVAYARSAAD $\square$ AA LIARELGLEVAL




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[^1]:[^2]:    lupas_21
    lupas_14
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[^3]:    lupas_21
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[^4]:    lupas_21
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