## QUERY

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UniRef90 S7TTP9

1
DGLKMENTKMPVIFNHSSHSSYQCADCHHPVDG PPVPDKPLEFKGSKKTVMFPHAPHQKVECVTCHHKVQD KRPAGVLKT SFAPVSFSHGQHANIPCVTCHHMWDG A $\square$ K EP $\square \mathrm{E}$ I EAS GSKRMNVMFPHTAHKGISCFHCHHEEGS PLTLRY LIKKEVVFPHAAHGALECRTCHHPWDG $\square K K P \square E L N G G T S K R M F V T F N H S S H K G V K C R T C H H E G L P$ $P$ VPADGLVLQGPKKPVTFNHSTHKTVECVICHHPVDG K VRFAQVKFSHFDHQDVNCVKCHHTWDG DKPDEFKGSQKTVMFPHADHAKVECVTCHHQVDG A Q IE KPMTMNNTGNEKKQVIFTHSAHAKTDCAFCHHKAVE S A MEAPGDPEGVKARFSPVEFSHDAHSALDCESCHHMLAE AAPSAFASDLPAEARRPAVPFSHDSHHDRHDCMTCHHMWDG DAPSCPVKMEATKKPVVFNHASHADSKCDACHHPDDG APD D DLVINF LIKPDVVFNHSSHENYECDDCHHNIKK APPPSMQATKAPVDFSHDRHAADDCATCHHTWDG PKQFEGYTPKKGDVTFNHASHMD\ACQQCHHTVPD AVDAPGDPAGAKMTKSPVDFSHKGHAAVDCTKCHHNWDG
 AANEAPADIEPFKMRKAPVQFPHAAHEKLECTTCHHEWDG ADGAK DDF $\mathrm{D}^{2}$ GGEKNDTVVFNHSTHKDVKCDDCHHDPGD APPEEGDARRAAVVFPHGAHFDYSCQLCHHDWYG APPEEVDARRAAVEFPHGAHFDYSCQLCHHDWYG PEGVEATRGSVGFPHGLHFQFACKSCHHDWDG PEGVEATRGSVGFPHGLHFQFACKSCHHDWDG $\square K R P A K N K P V A S W V G P V K F P H G F H A D H N C K A C H H E E S D$
 TKPAVLFPHDTHGSMDCGTCHHKGYQ PEDVEAKFADVRFPHADHFSYSCMECHHMWDG $\square K R P E K$ NQPMAAWVKPVTFPHGKHAVLNCDSCHHEESD PQGAKLAKPRVPFPHEAHTAVDCGTCHHKGYV CGSCHHNSPA CGSCHHNSPA E A P D D G $\square E \square$ N H $\square \mathrm{E}$ G S $\square \mathrm{MF}$ N H S S HESYD C A

APDKPDELKSSEHKKMWVKFDHATHESVECDVCHHRSPL VDAPADPAGAKVTKAPVAFSHKGHASMDCKTCHHKWDG PAGAKATKTPVVFPHKKHGG■D | D |
| :--- |
| C |
| C |
| L |
| S | CSACHHNSPL

A VDAPADPEGMPATQSPVKFSHKGHAK DDCKACHHK CAVCHHNSPL CAVCHHNSPL C A
PSGARQTMPPARFSHQAHTP $\square$ NCT $\square$ CHHTGYV KAPADPAGATLTQGQVMFSHKTHASYDCKTCHHTWDG

PVPFSHTVHGDYDCGECHHPDDE
VTFNHSSHEGYSCSSCHHNSPA ADPVVGADGKPVKYGMPPVPFDHKYHESKNCVSCHH

NCGACHHVYDK
NCGACHHVYDK $\begin{array}{lllllllll}C & A & C & H & H & N & \text { P } \\ \text { C A A C }\end{array}$
K A VYRMSPVVFDHKQHEAASCRVCHHESLE APEEYGEMREAPSP FSHTVHGDFDCGECHHPWDG
 SAVPFNHKLHEEKNCSVCHHNASS RSAVFFPHGLHMDLECLDCHHVYEG
PAGAKMTKAPVDFSHKGHAALDCTKCHHKWDG
$\square E \square K H G T S K R M H V M F N H T T H K D U A C E Q C H H D P S P$ CQGCHHNSPA
K PATPYAGMQPVPFDHKTHEAKNCVSCHH
CAGCHHNSPA CAGCHHNSPA IK R PKNLEAQSSWVGSVQFPHGLHAVMNCRACHHMETD APKDPVEMKYKYAPIMFMHDKHLGVACATCHHKWDG ERPKGNEPLATWVKEVKFAHGFHARVPCQKCHHKESD PKKA $\square E \square K H G T S K R M H V K F N H T S H K D V A C E Q C H H T P D P$

PEGSPMTKPAVVFPHKKHADLDCLVCHHKSES C SGCHHNAKP
CSGCHHNAKP
SGM
PNGQKMSKTPVEFPHKMHVD $\square$ DCLVCHHKATS TYERKYEPVRFMHAKHAAADCASCHHRPED $\square E R$ P AN Y E K L N S WVTK V NFTHGAHARVSCNQCHHKESD

PATMERTMPRVVFSHDQHREMECSLCHH A G
PQVPADVVIDHLSNPKVKFSHKAHASLACQKCHHKWDG CQGCHHNSPV

## QUERY

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UniRef90-G1UX3 UniRef90-G1UX31 UniRef90-LORDM8 UniRef90_UPI0003B6545D UniRef90_E5Y4X2 UniRef90_C8X5K5 UniRef90_C8X085 UniRef90_UPIO0035C6857 UniRef90_UPI00035FDC4F UniRef90_C6BUH4 UniRef90-E6VZT9 UniRef90-F0JDK5 UniRef90-P38554 UniRef90-LOR8D6 UniRef90-S7USW4 UniRef90-UPIO0036DD270 UniRef90_P00133 UniRef90-E5Y2S7 UniRef90-S0G6U2 UniRef90-S0G6U2 UniRef90-15B3B3 UniRef90-15B3B3
UniRef90 C6BRKO UniRef90_Q728S8 UniRef90_UPIO003B627A8 UniRef90_UPIO003B627A8 UniRef90-A1VBG7 UniRef90-B8FKY6 UniRef90-B8DM32 UniRef90_LORBZ6 UniRef90_LORBZ6 UniRef90-COQHA6 UniRef90_Q9RN68 UniRef90_Q9RN68 UniRef90-B6WSI9
UniRef90 P00134 UniRef90-F0JIY2 UniRef90-G1UVJ3 UniRef90-C4XGG7 UniRef90_UPI00039ECBD1 UniRef90_UPI00039ECBD1 UniRef90-B6WTN7 UniRef90-B6WTN7 UniRef90-E5YBQ5 UniRef90-S7U5Y8 UniRef90-D6SNJ7 UniRef90_C8X171
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UniRef90_Q30UU9 UniRef90-M5PSE9 UniRef90-M5PSE9 UniRef90-M5PSE9 UniRef90-C8X3T7 UniRef90-P94691 UniRef90_S7TTP9





 463133B



50
60
70
80
KEDLQKCATAGCHDNMDKKDKSAKGYYHAMHDKGTKFKS C K E N L A K C A T A G C H DV F D K K D K S V
 DGRYVACT ECHATP GARERDPMSMFMAFHSKNS D R S C ENPMPKCSEA GCHDVV DAKDKSEKSFYK VVHGPGAAAPSC K D RYASCTNEECHSIKGARERDTMSSLMMYHSPDTDDRSC KESYAKCATAGCHDNL KDKKGTNSLLYYVMHAKEKKHQSC K T P V VK S C A A A G C H D E L $\quad$ VKK K G Q P Q S Y F K AF H T L Q S D KESFAKCGSS GCHDDLLAGKQGEKS LYYYVVHTKKE LTKHTN C

 SGQPASCSSSGCHDQQG RRREARSFYRAFHDR SDQRSC K ENYKKCATA GCHSAAKADKKKA GSYYK IIHNKK S G LIS T C



 GMSFTS YKCASCHSTAKK DK S DDSS YFKA DHGRKA LP T R C
 D AKYVSCASEECHSVVPRGDDGVAPSDFKAFAAKDS DH S C

 V VEEVKSCSSSSGCCHEEFWPGAPSARPWEKSFSMTGAYHKAC V E EVKK S C S S S GCHEEFWPGAPSARPWEKSFSMTGAYHKAC
 RTLGQFLPCTQCHNKPTVTE TVSSFYMAWH NDSTHS
 SKTPPSCAS
 $\begin{array}{lllllllll}G & E & P & P & R & S & C & A & T \\ S & L & T & P & P & K & C & G & S \\ S & L & T & P & P & K & C & G & S\end{array}$

 | $L$ | $P$ | $P$ | $C$ | $G$ | $S$ |
| :--- | :--- | :--- | :--- | :--- | :--- |
| $L$ | $T$ | $P$ | $K$ | $C$ | $G$ | CHTGTDINEKGYKPYFKAMHMKKNFRPSC C H T K E D K A N P GRP

C H T K


GEATQKCSSAG
S

## 

 KTLGEFVVACRQCHSADNPKDK TGFYRAWHSEGA
## \section*{K} <br> S <br> S

## R

 PKASEGYVASERGCHTDAGKKKKDPEGYYQAFHNK QSKGEEGTCRYCHGATQVENRSPF QSKGEEGTCRYCHGATQVENRSPF QSKGEEGTCRYCHGAT QVENRSPF RTAAHGAC

SKKPPKCAT CHHGKPDERTPMRPGLMMAYH K
NSSKAS TYEIK GCMSEGCHDMAVAEERDVIRYFYKA YHDQL NPEVYS CSSK GCHDLATPERKSIAYFRNAFHA


NSKASC
$\mathrm{V} T \mathrm{C}$ GKNLQD EDELYEDNPD DRCQDCHNTDASDDTTKAFHHQC
 DK P Y A S CT DNDCHATPGPRERDTMSMFVAYHAKDTD $\begin{array}{llllll}S \\ S & P & P K C A S & C H G K P D A D R G D R P G G K A A Y H\end{array}$ SSLLTS CAAK
SKTPP K C A S
SKTPP K CA CHGKPDADRGDRPGLKAAYH

SKTPP KCAS CHQPFDAAKGDRPGLKAAYH KAEIT GCRVEGCHDTSREMKREPTSYDSAFHSKDAKM EAFASCTNNDCHTPGPRERDTMSMFDAYHAKDTD RSC
 LNPPKCAS CHSKPQERTANQPGLKGAFHNQ

K S E
A C
TNE $\square$ T GCMSEG CHD S L

PK K K S D K S FY R A F H DMK
K S C

S DTPP P K C G S CHTNNI DPKNPARPTLLKAA Y K A L | C H S A | $K$ | $D$ | $P$ | $R$ | $V$ | $P$ | $E$ | $R$ | $P$ | $T$ | $L$ | $K$ | $A$ | $A$ | $Y$ | $H$ | $L$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| $C$ | $Q$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| $C$ | $H$ | $S$ | $A$ | $K$ | $D$ | $P$ | $R$ | $V$ | $P$ | $E$ | $R$ | $P$ | $T$ | $L$ | $K$ | $A$ | $A$ |

S ATPPKCGS
$D G K S L L D T S$ $\qquad$ T Y P S C H D GLLT ATLRNPFY

FY A F HG $\qquad$

SKKPPKCAS CHDMAVAEDRDIRYFEKAYHDQ
SKKPPKCAS CHGGKPDPNTPDLRPGGLKAAY H
P E A K P K F S CVS
A S LT S CAA K CH TVA G S
Y EKGKESTCRSCHE
KEGGFVTEAAMHKVDAGQS Y E K GKE S T CR S CHE
 A DLK T P P K C A S CHGIIEADKSVNKPSLKAAY H


SBCS QCHTPAGDEK GGFVVRLE QAMHAQ
TDSASC













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QUERY
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UniRef90 E1JRN1 UniRef90-B8IY72 UniRef90-T2G7I8
UniRef90-G1UX3 UniRef90-G1UX31
UniRef90 B6WXU8 UniRef90 LoRDM8 UniRef90_UPI0003B6545D UniRef90_E5Y4X2 UniRef90_C8X5K5 UniRef90_C8X085
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UniRef90 S7USW4 UniRef90-A8FWU2 UniRef90 UPI00036DD270
UniRef90-P00133
UniRef90- $5 Y 257$
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UniRef90_Q9RN68 UniRef90_B6WSI9
UniRef90 P00134 UniRef90_FOJIY2
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UniRef90-T2G9Q2
UniRef90 T2G9Q2
UniRef90 M1WM43
UniRef90 F0.JK49
UniRef90_Q30UU9
UniRef90 M5PSE9
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UniRef90-M5PSE9
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UniRef90 ${ }^{-}$P94691
UniRef90-S7TTP9
UniRef90_S7TTPs

VGChLetagadank k Kelt GCKGSKCHS
MSCHLEAAGSDKDLKKELTGCKKSKCH LECHTKVVAEKPELKKDLTGCAKSKCH L GCHMKTN $\square A R K E K G Q K P S P C S N N G C H$ L GCHKK LAAENPGKFPQFKGCR PCH
 LSCHVKVVAEKPDLKKDLTGCAKSKCH RGCHVKMNKEGKTD1ATAAPCANNACH
$\square G C H S K V V E G K P E L K K D L T A C A K S K C H$ MGCHQKAVAENPDLKEKFKGC NPCHA LGCHMAKKKAGETTGPACTQCH
LGCHMQ $\mathrm{L} K K A G E P T G P G C T T C H E K$
VGCHSTARRAGSTDVPSCGDCH
MS CHKEAAGKDKAMKKALTGCKKSKCHS LGCHVKEFGND KEMTGCDNSACH
L GCHKDMVKQGRKAPVACKDCHAN
LGCHKTAKKDGKPAPASCKQCH
VGCHREL KRQGPSDAPLACNSCH
VGCHKALKKAKEKTGPKCGDCHPKK
MGCHRVERQAGKTNVPACTACH
LS CHVNEQKTRSKDKPSLTGCTNSSCH
I KCHRQERKG
PKGCNDCH
ISCHKDKAGDDKEDKKK LTGCKGSACH
YGCHMKKRKQYTGFQKGCLPCHEKA
LPCHRELQQER
LPCHRELQQER
V SCHRDTDKKDKADLPKAAPVNCKGCH
VSCHRDTDKKDKADLPKAAPVNCKGCH
VGCH A VKQKDPNCAGCHE
MGCHRQKRLKQGEPP $\square$ SCTGCH
RDCHREELKKDKKQAAGFTGDACDGCH

MDCHAEALREKGKAAGPV G G G
LGCHRKMREAGKMPPLSCTGCHKK
NACHARLLAAGKAAGPECRDCHKVK
MTCHDAMNLEKPAS TDCTACHAKK
MTCHDAMNLEKPAS TDCTACHAKK
$\square$ C CHTREFGDD
DAMTGCATSACH
MGCHKGMDVARP RDTDCTTCHKAA
MGCHKGMDVARP RDTDCTTCHKAA
Y GCHTAMVG
$P G C R P C H$
VGCHKSMKKG
PTKCTECH
LACHKKEKKAGKNVPVSCKECH
$\begin{array}{llll}M G C H K A M K V G R P & K N T D C T T C H K \\ M G C H K A M K V G R P ~ K N T D C T T C H K\end{array}$
MGCHKKEAKG
MGCHKAMKVGRP KNTDCTECHK
MGCHKAMKVGRP KNTDCTTCHK
MGCHDGMDVARP LDTSCASCHK
MGCHDGMDVARP LDTSCASCHK
VACHTESLAAGKPAGPDCKACHQAP
LGCHREAKKAGKAKAPSCTQCHPKK
MDCHRDMTREGEETGP VACNDCHT
MGCHTRMELEKPAA TDCASCHPKK
MGCHTRMELEKPAA TDCASCHPKK
LSCHDEEFGND K KEMTGCYESACH
AGCH-ATVQKKPECAGCH
V SCHRELAEAKKETGPTCAGCHSA
VSCHRELAEAKKETGPTCAGCHSA
MACHDRMKIEKPAATDCAGCHT
MACHDRMKIEKPAATDCAGCHT
VGCHTQE TRKTDCAGCHD
MGCHRDLARAGEESGPACPECH
NGCHKSRKKEGKSAGPACKGCHTVK VGCH A QAQAKPECAGCHT

$\begin{array}{ll}V G C H K A L K K A & \text { TGPTKCGDCHPKK } \\ \text { YGCHKKMA } & \\ \text { GQHPETGCR }\end{array}$
MGCHDRMK I EKPA NTACVDCHK
MGCHDRMKIEKPANTACVDCHK
AGCH AA $\square$ QKKPECAGCHDS
MGCHNRMKDEKPAD TACAECHK
MGCHNRMKLEKPAD TACAECHK
LGCHKEKRLKREAMPPSCTGCHK LK
VGCHTANLKDNPDFKKKCNDCHSKK
LGCHTQRRAAGGDNPVGCTACHK
YGCHKQMAKDHPEFS GCRPCHMSA
MSCHKKEKKAGKAAPVACKDCHK
1 GCHQEMQVNPKAT DCQGCHKPK
I GCHQEMQVNPKAT DCQGCHKPK
LGCHKKAKKAGKSAPVSCKDCHPKK
MNCHEKMKKG PVVCEGCHTK
LGCHTDM RSKGGKNPVGCTSACH
V S CHLQKRAAAKDTGPNCAGCHSA
V S CHLQKRAAAKDTGPNCAGCHSA
$V$ S CHLQKRAAAKDTGPNCAGCHSA
LNCHTEEKTHGHNQD VPDHCASCH
QGCHKEMKTAKKTTGPACAQCHNQK
MTCHAEMKDEKPA GCTGCHKEK
$P \stackrel{20}{R} F$
P ViR F A H K R H A A Li ${ }^{20}$
C S K CHHR PK D
HA AEFAV CASCHHNSPP
HAAEFAV CASCHHNSPP
A D P AFAPRQRPGVFFMHEKHMDAECTDCHHFYEN MKAKPKGDKPGALQKTVPFPHTKHATVECVQCHHTLED

 PPASFRHDKHKDKDCSACHKTVDG PPASFRHDKHKDKDCSACHKTVVG
KA Q GLYSP VDF DHELHLDT TA CT DCHHMYEQ

| C Q S C H H N S P |
| :--- | :--- | :--- | :--- | :--- | :--- |
| C Q S C H H N S P |

KDPKYTPGQAPMGFDHSLHYDNKCEQCHHEYNK K DPKYTPGQAPMGFDHSLHYDNKCEQCHHEYNK

T Y D K K Y GPVRFMHAKHAASDCAACHHRPAD
EGRMKGVAFKHNFHEKNNCRVCHHETDK EDVYTSKKKPSVLFDHDLHMAVDCMDCHHHYKD EGEDPHPRFGKVEMSHAKHRNVSCVSCHHMFDG

PVRFMHKKHAAVDCLLKCHHAADE
PKASKMKAVVFNHR DHQAQKCRDCHH
RMKAVTFNHKAHEPRACNACHHNSLE
A D DQTKKNYMDAV AF DHKFHEAQSCKDCHHETLK
LGGTK T R GPVAFPHELHEGFECLECHHVMEN
$\square K$ APEGYSKVV TKTEVAAFSHKGHAKDDCKMCHHK
P PVE Y E EKY GEKR LA PVVFSHDVHF DYDCEYCHHT G

A A DPGPAE $\square$ T L GE G GKKPAVFPHKKHQAFKCADCHHGMA D HKGlR CTDCHQPHSG
PEGVKPRSTWWKKVQFDHETHSGVECRTCHHMEDE
PLYEK H T K G $\ V Q F T H K Q H V E \| A C G S C H H D E T G$
P A T $\square$ E F P GGKK GKVVVFNHEKHQKFKCGECHHPYK E
DAEKPCKLVEFTHKKHYKDSCGDCHHDKE G
$P \vee E F N H Q L H V D I D C S V C H H H T T G$
 K DKK GPVAFDHVAHAGVTCNECHHTWKE


GPVVFDHQKHSRLDCATCHHESLQ
A GVTCTDCHNPHTA
$P \square S F P H A V H G Q F P C M T C H H D V N D$ PKFQAAFNSSVHGSNGCRSCHHMTG TKAPVSFSHSTHEAVECLKCHHEWRP
PGDTKY GP LMFSHAKHAAQKCESCHHKMAE DLKLE GAKFPPVKFSHAKHYKVDCKVCHHKEDP A GVGCTDCHNPHT T
K K P K P S VF P HKAHQAFECGDCHHGVDA A GVTCTDCHNPHSN A GVTCSDCHNPHTG KRGPVKFNHDAHVSLDCTACHHRYEN SPVYF P HEEHMGFE CDDCHHCY K D VIF D H A K H Y GESCVACHHEDS D FDHAAHEEYDCQACHHGND ATPPTEVE IIVEGAKMAGVKFSHEKHYKVDCKVCHHKEED Q GV S C T DCHEPH T S
TRPAAVFDHDAHNEKDCAVCHHEEDG A GVT C T D CHNPHSG
KAPDEPLVFTGAGGSVVFDHATHAALTCTACHHHPQD D P AF GE GQRPPAVF EHDAHELFDCAVCHHYYED KKGPPQR K LVVTFTHKK HYN $\quad$ HS C GECHHDET G LID HAA HLA

GAFNCVVCHHNYTE A GVACSDCHEPHGG A GVVGCTDCHNPHTA A G V T C S DCHEPHSQ GAKFPGVKFNHANHYKIDCKVCHHEADP

A LP P P P S K VVRTTKTYGTVT LD HAAHRKI SCKSCH HKK G IV QFTHQKHYKI GCGECHHGDDG M PLA FP HEP HVVSV NCDTACHHDYK D R R P A AVFDHNSHNEADCAVCHHVWEN A GVT CLDCHRPHDA K H T TGPCTSCHHANDN K P E A VNEK P V G M A P VA F D H K L H E A A S C L T C H
 A A DQGPAEVTLLESGKKPAVFPHKKHQAFKCAECHHMADG H A GVSCMDCHDPHSN

DEAVFAKHTRPFVCFVHDEHAGIDCDVCHHVYED M S K LY GP VE F D H S M H VGYS CQE CHHHTTGG | $H K G \square R C D$ | V |
| ---: | :--- |
| C | C |
| N | C |
| C | H |
| S | P |
| K | H |

## UniRef90 KONPE1 UniRef90-S7T120 UniRef90-S7T120 UniRef90-P00136

 UniRef90-UPI0003B43F13 UniRef90-Q315E2 UniRef90_Q315E2 UniRef90_Q315E2 UniRef90-Q315E2UniRef90 C8X1Y2 UniRef90-E1YDG0 UniRef90-E1YDG0 UniRef90-A8ZUW4 UniRef90-A8ZUW4 UniRef90_E6VQS7 UniRef90-B5YFP9
UniRef90-D6SSM7 UniRef90_Q9R638 UniRef90_B8FFW2
UniRef90_UPI0003B5F9D4 UniRef90_I4CEC9 UniRef90_K0NK05 UniRef90-K0NTI5
UniRef90-E1JZ85 UniRef90 C6BRK7 UniRef90_D6SST1 UniRef90-F8C3E3 UniRef90-F8C3E3 UniRef90-A8ZTK8 UniRef90_K1YNL0
UniRef90-D2QYM7 UniRef90-F3YUC0 UniRef90-KONAD9 UniRef90_UPI00046CD62A UniRef90_UPI0003B6AB80 UniRef90-15B4L2 UniRef90-UPIO003734801 UniRef90_B9M5I7 UniRef90_14C8C5 UniRef90-D8FAC3 UniRef90_A8ZS59 UniRef90-S0G649
UniRef90-B6WV78 UniRef90-X7AI62 UniRef90_A0A023D6X9 UniRef90_Q2LQK1 UniRef90-F2NGH7 UniRef90_UPIO003B685BE UniRef90-B5YKW8 UniRef90 S2KEL1 UniRef90-V4IF16 UniRef90_UPIO00255704B UniRef90_R9VG68 UniRef90-S7V061
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UniRef90-UPI0003B72AFB
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UniRef90-S6AQ13
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UniRef90_C0QC90
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UniRef90-J3BRZ4
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UniRef90-M1PTB4
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UniRefgo_H8MFZ2
UniRef90_H8MFZ2

|  | MECHKTEAKG PVDCK |
| :---: | :---: |
| $\begin{aligned} & \text { UniRef90_KONPE1 } \\ & \text { UniRef90_ST120 } \end{aligned}$ | MECHKTEAK <br> MGCHTMMGIEGKDAKPVPAATDC |
| UniRef90-S7T120 | MGCHTMMGIEGKDAKP |
| UniRef90-S7UJB9 | MGCHEAKGVASPAA |
| UniRef90-P00136 | TDCHKALKKDKKPTGPTACG KCHT |
| UniRef90-UP10003B43F13 | $\checkmark$ NCHTSLTTSAS |
| UniRef90_Q315E2 | 1 GCHTEMVDNGQKSGPECRS |
| UniRef90_O315E2 | I GCHTEMVDNGQKSGPECRSC |
| UniRef90-Q315E2 | 1 GCHTEMVDNGQKSGPECRSC |
| UniRef90-Q315E2 | 1 GCHTEMVDNGQKSGPECRSC |
| UniRef90-C8X1Y2 |  |
| UniRef90-E1YDG0 | LGCHKEM |
| UniRef90-E1YDG0 | LGCHKEMKTGGKKPAGCTDCHKK |
| UniRef90-A8ZUW4 4 | TCHQAEKTKKKNSGPDCGGCH |
| UniRet90-A8ZUW4 UniRet90 E6VQS7 | MGCHADMKKGKKS |
| UniRet90-B5YFP9 | QGCHKQM TSKKECSGC |
| UniRef90-D6SSM7 | DTCHRDMAGEDATGPR |
| UniRef90-Q9R | CHKAMKAKNEQ |
| UniRef90-B8FFW2 | MTCHOKETEEA |
| UP10003B5F9D4 |  |
| UniRef90-4CEC9 | $1{ }^{\text {a }}$ |
| UniRef90-KONK05 |  |
| UniRef90-K0NT15 | VKCHDQYS |
| UniRef90- C6BRK |  |
| UniRef90-D6 |  |
| UniRef90-F8C3E3 | LSCHTKTNKEMEVSKKEAPP鸟CGKCHT |
| UniRef90-F8C3E3 | CHIKINKEMEVSKKEAPPLECGKCHT |
| UniRef90 A8ZTK8 |  |
| UniRef90_K1YNLO | LECHKEKAGSDPALKEKCDTCHPKK |
| UniRef90_D2QYM7 |  |
| UniRet90-F3YUC0 | V SCHR |
| UniRef90-KONAD9 | RDYNKEKGDPKKEPAPVSCTKC |
| UniRee90-UPIOO046CD62A | K |
| UniRef90 15B4L2 | KGCHKKMKERNAPTKCGGCM |
| UniRer90-15B4L2 | RGCHQE GQGKDGPTGCOTCH |
| UniRei90 ${ }^{\text {B9M }}$ - ${ }^{\text {a }}$ | RDCHAPKEDSGQEMNRMCAGCH |
| UniRef90-14C8 | VNCHKQFKKDKKPTGPACAK |
| UniRet90_D8FAC3 | NCHKE EKKGPYKKCNDCHAKK |
| UniRef90-A8ZS599 |  |
| UniRe990_SOG649 | MGCHNNYSLTSEPT |
| UniRet90-B6WV78 | VTCHHYTLAAKEEFGVDCTSCH |
| UniRef90_X7Al62 | V |
| UniRee90 A0A023D6X9 | RMARN |
| UniRef90 Q21 | VDCHRDUHNQRAGREQCTACH |
| UniRef90-F2NGH7 | GHKMKKG |
| UniRet90 - ${ }^{\text {B5Y KW8 }}$ | G |
| IRet90-B5YKW8 |  |
| UniRef90-S2KEL1 | $\checkmark$ CCHMPSKAAQGIGPDACTACHK PCH |
| UniRee90-V4F16 | KGCHEQVJA |
| UniRet90-UPIOO0255704B | VNCHMPS |
| UniRef90-S7V061 | CHA |
| UniReto-SM0603 | GChag |
|  | V C CMP |
| UniRef90 UPI00036F8DE5 | MGCHMENGSGP |
| UniRef90- 66 BTM | ПDCHDEMFEAGL K |
|  |  |
| UniRef90-E1SC6H5 | CHMPATTAQTGAPDACN |
| UniRef90-A0A031M105 | CHMPERARQLGTPDACTGCH |
| UniRef90-Q30UZ3 | QSAGKGPA NCGGC |
| UniRet90-F3KYS8 | CHMDA |
| UniRef90_S7TCB0 | CHK |
| UniRet90-C8X122 | Q |
| UniRef90_SOFXO5 | KQVNAGDPKGRKGPAPTSCTQCHE |
| UniRef90-Q21J34 | VQLAKG PTDCRGC |
| UniRef90-UPIO0022605F7 | EGCHLKLKREKRASGPVRACCHV CHA |
| UniRei90- T2H0P3 | VSCHMPTKASKV |
| UniRef90- UPIOOO35DC7E3 |  |
| UniRef90 UPIO0033B72AFB | VECHKKVAEEGKQAPV |
| UniRef90- ${ }^{\text {- }}$ GGB19 |  |
| UniRef90-S6AQ13 | V |
| UniRet90_B4UBF4 | C R |
| UniRef90-C0QC90 | CAKCHPKK |
| UniRef90 UPIO0004672969 | QSCHLERLLQAFHASGPSCQACHRN |
| UniRee90-UP1000477254C |  |
| UniRef90_J3BRZ4 | CTTCHAGK |
| UniRef90 Q1DEC2 | SSCHLTKESUQRTSKAACLSCHKS |
| UniRef90 UPIO0025587DB |  |
| UniRet90-FJEL9 | AGCHAARPAKA |
| UniRef90-E1QFI8 |  |
| UniRef90-K1YPP7 | ECHKTKANEDAALKEKCETCHPKK |
| UniRef90-B5JHX4 | VNCHMPER |
| UniRef90-C0QH26 | SCHLE RKEGP |
| UniRef90 S9F | DKCHLKAEGLQR |
| UniRef90-M1PTB4 | VGCHTDTSG PTGCVDCH |
| UniRef90_Oplooos7412DB UniRet90-H8MFZ2 | VECCHMKOETTQRTSKAACTSC |

SPTTVMESVHAKFACVDCHYKGHE ${ }^{20}$ VFEHGARAGVTCLDCHRPHDA K GVKCTDCHNPHSG
FPQSAHGNNACTSCHHM TG
HE GVTCSDCHDPHTL
DPAFPKHQRPAAVFAHDAHNEMDCAVCHHVWED EGVGCMDCHHRAEV A GVTCSDCHEPHGQ A GVNCVDCHNPHTM
HTRLPAVFNHDAHNEKDCTTCHHKRDP A GVTCTDCHHSTRV A GVTCSDCHEPHGL

APVAFDHKFHEAKDCSTCHDNG
APVAFDHKFHEAKDCSTCHING HA GVTCSNCHNAHTG HAGVRCTDCHQPHTL HKGVKCIINCHNPHSG
PEAPSD HKVK ARAAAYFPHDRHMEVGCNRCHHEEDE $C Q G C H H N S P A$
$C Q G C H H N S P A$
D G L WP T P T K A A VE LT HKKHAK IA CAECHHVYKD KMHEKVTCSNCHNPHSG
A QKAPESP $\square$ RFKNTGGNVLFDHKTHYGISCSDCHHDDEE ADPAFAPAKRAAVPFNHDAHNAANCAACHHAYEN A GVTCLDCHHEATL
HSGVDCVDCHQPHSV K GVRCSTCHDPHRG E GVTCVDCHDPHTQ
VKEYKVKMGKATFPHAKHAGNSCGTCHHKVNG VKKDHAKH $\square$ NLDCAVCHHHTTG A GVTCLDCHRPHEA
RAPVSFTHTAHMDLECLPCHHSGVE
PAYAKHTKG■VQFTHKKHYKI GCGECHHDEKE
 A GVVGCTMCHOPCHT
HKKHQGVDCRECHHKNTE
HSRPAASFVHDTHNEKDCGVCHHGTTE A GVACTDCHDPHRG KGVSCTNCHNPHSA
KMHAAVRCMDCHDPHTS
GLS C S T CHDPHS G

QDPAFTKRMRPEAVFNHDEHNENDCSTCHHFYKD
SVEFNHAQH ${ }^{\text {V }}$ RDCAGCHHHTTG K GVQCTDCHNPHSG
VLFTHQTHTELDCLDCHHNDDD
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HNGVVCSNCHNPHST KGLRCSTCHNPHST

F QNKAGKVLFTHSVHSDQDCLDCHHNDED HKGlRCTDCHDPHSL
KHTKG I MMFTHKKHYKLDCGECHHDDK G
EYADHSKGLVEFTHKKHYKLKCGECHHDAEG
VK E Y QVK $\square$ GKASFPHAKHAGYSCGTCHHKVEV
FDHEVHKDYDCGDCHHGKDI QGVNCMDCHNPHTL

HNDVTCTDCHHSLKG
DPAFEHPERPAAMFEHDAHNEKDCASCHHVWED GGKKGTVTFHHDKHQK $\mathrm{V}^{\mathrm{C}} \mathrm{K}$ GECHHPYKE M F N AGVTCSNCHNPHSG KMHAGVTCLNCHHSLDL HVSIKCTNCHLDPS G
HAGVTCSDCHNPHSL
 PELAPFKYAPASFDHDPHNEKDCTACHHGSQD A GVTCLDCHHAATL
FAASPHGKNGCTSCHHM TG A GVTCADCHNPHSG
RSAVTFNHDEHADEDCTECHHSFDE
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UniRef90 Q6ALR1 UniRef90_UPI000477A247 UniRef90-A0A024HAWO UniRef90_UPI000472D2F7 UniRef90_T2KR17 UniRef90-C6COZ5 UniRef90_UPI000237A00C UniRef90_I4JLC6 UniRef90-UPIOOOOE11070 UniRef90-E1JSW6 UniRef90-B1J6W1 UniRef90-UPI000367000F UniRef90_M5H5D4 UniRef90-F9U5M6 UniRef90_UPI00036F7DF9 UniRef90-UPI00036F7DF9 UniRef90-K6ZRX9 UniRef90-W4LMJ7 UniRef90-TODHB4 UniRef90-S7TBW8 UniRef90-E6VRF1
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 GLKAEGNACTQCHTETKPERQDPSGLFHTRHQAGSTGAQC EVK $\square$ A GNGC $\square Q C H N V A G K Q A K N Y D S P E H H H H K P G T P G A Q C$ REKPAAVSCGDCHAEDARE KTKYTDNRCDQCHT

PNYNEPSHHFH
A GK T D D
R
GK VVEDESCSDCHAV KAEAGKTDDRNAYH KLRAEGNACSQCHAPAA KVKTQTNGCDQCHQPQAYE SQDTSG円PCSDCHTAAGKE PGRTPSQHDMH KREGNSLCLQCHNPKPPTEKNYDSPVHHHHTPSSAGEQC K $\square$ RATGNGCLQCHS KNKYDTEKHHFHRTSSAGSQC KTKTNGL CLQCHSASEYNKPEHHR RHKHEQSTGAQC HAEGNAV CARCHTPAR YDSPEHHHHQTGSTGAAC IGNDKDGFKPLYSDMHDAKSSASC KVK $\square P G N G C L Q C H Q A G N D K D G F K P L Y S D M H D A K S S$ ASC KLRAEGTACVQCHQSQPDAAKAYDTPDHHFHP $\square G S T G A Q C$ KLKKDGNSCLQCHDLD SYEKYDHHKKESSASC LDGGEAMRCRTCHTDA SKTPPKCAS CHAPFDPATPDRPGLKAAYHGQ SKTPPKCAS CHAPFDPATPDRPGLKAAYHGQ
 EGD TPAACSECHTPDGEEAPKRV DVLHEQC GQDSVGVPCSDCHLKAGMAAAGGQDVTAETTLMNAYHRQC KAAGNALCTQCHTETAPERGEFDTPVHTHHPVGSSGAQC KLKVEGNQCVGCHSSEQFDTPKHHDLSQATNDQQTQGTQC NLKVTGNGCLQCHNPEGKKAKDYDSPTHHHHRPGQAGAQC RTHEEGNACARCH AETEFDQPTHHFHPPGSDGAQC S LKADGNACTQCHTETKPEKQDPSGLFHTHHPVGSAGAQC SSETKIVACKGCHTLREKRQGTKDYHQDKPGLKGAYHLNC GLKAEGNACTQCHTETKPDRGRFDTPAHTHHQAGSTGAQC KKAPPDQLCAQCHNP QTFDTPKHTFHKAGSVGSAC DGKFVPAPCKDCHAAEKDGEVNDAAAYHTQ GDAVQGC ECHPKASKKKGEPKSEKMEYHAE ALHANC QVKI A GNGCDQCHNPAGKPAKNYDSPEHHHHQSGTPGAQC QVRIEGNGCLQCHNAAGLQAKDYDSPAHHHHAAGSPGAQC
$K$ TRAQGNACESCHTPAV EAEPQNCAD CHDSKKAEEGTPPNLADAVHAR
 DGSSEGTPCADCHAADNGSRGKGTPLERAYH
RLRAEGNACTACH NPAPDRGRFPGLY DAFHRGGGAGSQC KLEDAEEACSQCHTQFAPEN HTFHDKDSEASKC KNPVEGNACMRCHGSPAPVIQGDVDPDHSFHKAGTAGSRC GVRVEGNGCTQCHNPSGKKARDYEAPEHDHHAAGQPGSQC QLKMAEPACAQCHVAA KTETNGDACLQCH A A AKTYTPEK KTETNGLCLQCH EAKTFDTKDHHHHDPASEAAMC
 EKAPGNGVCLQCHNTAGKQAKNYDSPEHHHHKPGSPGAQC DETYN CSDCHEETGDDYMPSRT
GKLVEGEACADCHTL KAQGSQPGDMMAYH
KLVLPNNACSQCHSPA KFDNSEHHHHQTNSSGAQC
ALRVTGNACTQCHNPSGKRAKNYDSPEHHHHAQGQAGSHC KSAPSANDCMNCHSQ DKAESPE DQK $\square Y K A \square$
RSTSASPTCVLCH DETYN CSECHEETGDE D DPSRADAFHAQ ELKHPGETCTSCHQHAAGKYDVPS HHHHAVGSEGAKC GDDVQPC LECHTKTGMPDRDKKKAEDEYHYGADHENC
 $\begin{array}{ll}K K L P A N Q T C T Q C H T S E V F D T P D H H G & H K V D T T G S Q C \\ N G K M Q S V P C K S C H G D L S R K Q ~ K K G F H K N C\end{array}$ EPEPQKCTN CH GEKSTNNLLSMREAGHKSC K LQRAGNQCTFCHQSTPPKEKQYDAPSHHFHKMDSPGANC

 GKIVEDESCSDCHTV KPESGKTGLRNAYHKL GMKIQKC SECH NKDMANKKLNSVKKAMH K LKLPGQTCTQCHSQVDFDTPEHHG HKDESEGSNC KAEGNGLCTQCHSPAGNPDADYDSPAHHFHPEGSAGATC K LRMQGNQCLQCHT ELKAKGNACGQCHQASKYDVEAHHG $K L P D E T Q C L Q C H Q S D$ NYASTQHHKHPESSTGAQC KEETLSCV GCHEDQDDADQRDEEFERSVHATSDHPKSC $K L K F S G Q T C T Q C H N A T T Y D L P E H H Q$
$G M K V Q E L P C A D C H T V$
 KADGNA C CTQCHTETAPEKNDPSGVFHTHHPVGSSGAQC NTK DDKVNCSRCHGNETKE ANDFDFEKSVHAKSGLGAC K LRLA GSACGQCH A ANVFDAPAHHHHQKDSAGAQC TVSPSNALCLGCHTKENPAGLKGTVSEHTHHAEGSKGSQC
 KDEVKKCG ECHNKFKKDKKNKK DDMVHENAMHGNC
 KPDDLK GGCAECHTKMAKDKKD DMVLENAMHG KKEGNDLCCLTCHVPDT


| UniRef90＿Q6ALR1 | $\text { TSCHSQEAFHGQNG }{ }^{90} \text { GKACRQCHNDP }$ |
| :---: | :---: |
| UniRef90－UPI000477A247 | ANCHMPER |
| UniRef90－A0A024HAW0 | T S CHMPEHAQK $\square$ GAPDAC $\square$ G C |
| UniRef90－UPI000472D2F7 | SDCHRDLKNKKRAVIEKCTECHGND |
| UniRef90－T2KR17 | V NCHM |
| UniRef90－C6C0Z5 | SNCHVKEDKGP】 $\mathrm{S}^{\text {S CAGCH}}$ |
| UniRef90－UPI000237A00C | K NCHM |
| UniRef90＿14JLC6 | V C CHMPET |
| UniRef90－UPIO000E11070 | V S CHMPT T T S F K $\square$ P R PE LTP P A C N S CH |
| UniRef90＿E1JSW6 | MGCHM AKKKGPVTCGDCH |
| UniRef90＿B1J6W1 | V ${ }^{\text {CHMPEA }}$ |
| UniRef90＿UPI000367000F | V D CHMPAKTARLGTPNACANCHANK |
| UniRef90－M5H5D4 | V S CHMPTNR |
| UniRef90－F9U5M6 | V A CHMDERADHSLRVPRPTPNACNACH |
| UniRef90－UPI00036F7DF9 | VGCH AMRIAQNPSCEGCHT |
| UniRef90＿UP100036F7DF9 | VGCH AMRIAQNPSCEGCHT |
| UniRef90－K6ZRX9 | $V \mathrm{SCHMPEK}$ |
| UniRef90－W4LMJ7 | V NCHMPAK |
| UniRef90－T0DHB4 | V D CHMPKKA |
| UniRef90－S7TBW8 | I Q C HR A LE K E G NVS GPRTC G C |
| UniRef90－E6VRF1 | MSCHTAMKLE KPSNTGCGDAGCH |
| UniRef90＿E6VRF1 | MSCHTAMKLE K K S NTGCGDAGCH |
| UniRef90－A0LQL1 | QECHKKHKQEKPDTKATCAGCHPAP |
| UniRef90－UPI000360520F | V D CHMPSTTSFKVPRPDLT P N A CNACH |
| UniRef90－D8FCE1 | KGCHDE GGSGPVECGQC |
| UniRef90－S7U237 | QTCHEQ RGKGPVM |
| UniRef90＿UPI00046FB5B0 | ANCHMPER |
| UniRef90＿W2UDV7 | 1 D CHMDYRRDHS F R $\square$ PR P T P N A CNDCHT |
| UniRef90－L8MQB9 | 1 DCHM |
| UniRef90－K0CDT5 | V N CHMPETTYMKVDPRACNGCHEDK |
| UniRef90－W6S3A9 | A C CHM |
| UniRef90－UPI0003B47789 | LT CHKEVKKS GKKVPTKCRECHDK P |
| UniRef90－A5G7N7 | MGCHTKMGG PTGCLDC |
| UniRef90－C6B781 | ANCHMPER |
| UniRef90－UPI00037CE7B1 | VS CHMPEKTEKDDTPNACNQCHDN |
| UniRef90＿UPI0003B75C08 | 1 GCHNEAKKGP】T CGECH |
| UniRef90＿S7UXM1 | 1 D CHKEHNKKNKAAPTSCTTCH |
| UniRef90－W5IM53 | TSCHMPGKAQK】GAPDACMGCH |
| UniRef90－B0KGF0 | V ${ }^{\text {CHM}}$ |
| UniRef90＿UPI0002D4E981 | V A CHM |
| UniRef90－UPI0003B52407 | $V$ TCHEEMFAAKA KGCAQCH |
| UniRef90＿G2HB29 | MGCHKQQAKGPTA CGGCHAKA |
| UniRef90＿C4K963 | VDCHMPSRGARTGAPDACTGCH |
| UniRef90－UPI00030ACB44 | TTCHMPE |
| UniRef90－B4CWD3 | VDCHMPET |
| UniRef90－U3HH47 | V D CHM |
| UniRef90－R9PMG6 | SSCHM |
| UniRef90－UPI00036FAE80 | VNCHMPET |
| UniRef90－E1YFN0 | $\square$ GCHEQE KAGP $\square$ MCSECH |
| UniRef90－B9M5X1 | MGCHEKMGG P TGCRD CH |
| UniRef90－M4WQ73 | TSCHM |
| UniRef90－S0G5P4 | $\square$ GCH ADYGAGPVDCN |
| UniRef90－C7LX18 | KS CHDES K G GPVACGECHVKK |
| UniRef90－B8CMA0 | VNCHM |
| UniRef90－S6APU0 | V D CHM |
| UniRef90＿UPI0003764B9A |  |
| UniRef90－W5ILM5 | E S CHLKRLQ S F Q A GPSCKGCHTPP |
| UniRef90－K0NH48 | KGCHEDLGAG PV |
| UniRef90＿M2AYB1 | V N CHM |
| UniRef90＿A9A054 | QGCHETYNKEKVDARKGPAPVSCTQCH |
| UniRef90－C0QD60 | IECHKEHNDSLKDPKKGPAPASCSKC |
| UniRef90－UPI0003685F1A | V D CHM |
| UniRef90－F8ABA3 | LS CHK T $\quad$ K K E GKNVP T K C K E C H $\square$ K P |
| UniRef90－F0JJH8 | GQCHEDMFDKGLSS SCKSCHSQK |
| UniRef90－A7BWE2 | V NCH |
| UniRef90－C8PH43 | KDCHQEQAAARADTHKDCASCH |
| UniRef90－U1M916 | VDCHM |
| UniRef90－M7MLA6 | 1 NCHM |
| UniRef90－LORDE2 |  |
| UniRef90－F8A8M9 | K G CHKQVK DKHPNAPTKCKECH $\mathrm{C}^{\text {K K K }}$ |
| UniRef90－Z5XS85 | VDCHM |
| UniRef90＿B6B812 | QS CHMPER |
| UniRef90－UPI00036AB4F4 | K S CHM |
| UniRef90－N6X170 | V NCHMPEK |
| UniRef90－Q47ZT2 | V N CHMPETAQKLGTPDTCLSCHENK |
| UniRef90＿W9VF27 | HS CHDPHAFRNSRVGEECLSCHKK |
| UniRef90－M4S001 | V NCHM |
| UniRef90－UPI000366CD6F | 1 T CHEKQNKG PVQCKSCH |
| UniRef90＿U1K788 | VDCHM |
| UniRef90－UPI0003B37F4E | A C CHMPER |
| UniRef90－B3E3G0 |  |
| UniRef90＿UPI000367EF74 | A N CHMPTTTASLLATPNACNACHDGK |
| UniRef90－E8X6V4 | VA CHMPK $\mathrm{C}^{\text {E A }}$ |
| UniRef90－S7V7A1 | KGCH EDQKLGPVMCGECHQKK |
| UniRef90＿A0A024HHW4 | EGCHLKNDQAFRQTGPSCKAC |
| UniRef90＿UPI0003B779BB | VECHKEAKAGDPKGRKGPAPTSCGKCHE |
| UniRef90－D4H311 | 1 KCHKEM KVKKSCSTC |
| UniRef90－S6AF05 |  |
| UniRef90－K0N3M5 | 1 V CHKDVNKKAGDPKKGPAPASCAKCH |
| UniRef90－UPI0003821C8A | INCHMDGKLDQSAQYNTPNAC NKCH |
| UniRef90－IOK4M5 | QSCHGEEKMEVVEQRSSLTMGWCDDCH |
| UniRef90＿UPI0003B53BF2 | A T C H E D M F D K Gll K G C A P CH K M K |

UniRef90 S0FWJ2 UniRef90 Q74GH2 UniRef90-UPI000382191F UniRef90-A6ULE6 UniRef90-S5YH10 UniRef90-M1WLR7
UniRef90-B8FMZ9
UniRef90-UPI00047702A9
UniRef90 Q026X7
UniRef90_LOF8M3
UniRef90_UPI00039EEF69
UniRef90_UPI000255BBF3
10
 NK I SAAAFSASVHGKNACTSCHHM K G PAFTENKRPAAVFNHDEHEVYDCTPCHHEGVE

PMTFAHKDHGSVTCATCHHEFAK HKKGlVQFTHGKHYA $\square$ C C G G C H H H D A P A QGVKCSSCHNPHTG ALSCLTCH HEDT ${ }^{\text {D }}$ A
 KMHAGVTCSNCHHSLELL



| UniRef90_S0FWJ2 | $\text { RDCH } \quad \begin{gathered} 90 \\ \mathrm{DRSQGPVLCGEC} \end{gathered}$ |
| :---: | :---: |
| UniRef90-Q74GH2 |  |
| UniRef90-UPI000382191F | $\square \mathrm{DCHTE}$ K K VGPVACGQCHVN |
| UniRef90_A6ULE6 | A N CHMPER |
| UniRef90-S5YH10 | RSCHVEDHAAGKAAGPTCLSCH |
| UniRef90_M1WLR7 | MGCHAAMKLEKPTTCDDAQGCHKKK |
| UniRef90-B8FMZ9 | RDCHKAYNKEKGFKGKDGAPTSCTAC |
| UniRef90-UPI00047702A9 | K C CHM |
| UniRef90-Q026X7 | V A CHMPQVAARA |
| UniRef90_L0F8M3 | V D CHSNSLVAKKNDPQT C A K C H D N P |
| UniRef90_UPI00039EEF69 | V NCHMPTNRAQFNTPNACTQCHDDK |
| UniRef90_UPI000255BBF3 | $\checkmark \mathrm{NCHM}$ |

lupas_21
lupas_14
lupas_28
jnet conf sol25
sol5
sol0
jhmm

jpssm


[^0]:    

[^1]:    

