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GWLHKRGE
K V I K Q GY LEKE GKVFRTWRRRFFVLMDDNLGYYD KA GYLWKK GERRKTWKKRWFVLRP AHLAYYK DNAVI KA GYCVKQGAVMKNWKRRYFQLDEN TI GYFK V L K C G WL K K Q R S I M K N WQ QR WF V L P G D Q



 $N \square P V V M R G W L H K Q S S G M R D W K R R W F V L T D D C L Y Y Y Y$ K A GY L WK K GERRKTWKRRWFVVLRP A H L A Y Y K VSK Q GWLHKQSS GVKQWNKRWFVLT DRCLEYYK NTPVTKA GWLFK QS S GVVKQWNKRWFVL VDRCLEYYK HE GYLYKRGALL K GWKQRWFVLDS RRYFI LDQANSLLWYM NAPVIKNGWLHKQSTGMKMWKKRWFVL SDMCLFYYR
 EP Q Q I REGYLVGK GTLLNSWKAVWVVLSDDG LEFYK QVTLLRGWLYREGGAIKQWKRRWFVL GDYCLFYYK $P \vee L K A G W L R K Q R G D M K N W Q Q R W F V L C G D Q \quad L \quad F Y Y K$ NS PVL DR GWLYK QS S GLLK M WKRRWFVL

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 IVVK Q GY L T K K GAMRRNWTKRWFVL K Q G Y L L F Y F K
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$S G T V V K$
S G A

$K A G Y L S K L G G R V R A W K K R W F V L Q Q D A B V Y Y K$ S M D S S Q SVVH S GWL R K S P P EKK A WR R R WFV L R S S Q EL E Y Y K

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A N A PVA LA G WL Y K Q G S D G K V WR R R WF V L S S Y V L Y Y Y K LDERQVLWS GFLTKRGRFRKNWKRRYFELYDNHELAYFK

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| UniRef90－UPI0000E477B5 | DKPKGKSGDSDALNNFTVAQCQILKTEKPAPNTFKVRCLQ |
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| UniRef90－A0A024MRX9 | AKPDPDQPLPDPLNDFTVKNVQLMKVERPKPNTFLVRGLQ |
| UniRef90＿T1JPP3 | QKPELGKEAGEPLNRFTVRGCQVFKLEKSKPYTFLLKGLH |
| UniRef90－ROKV46 | ERPETSDHSLPPLNN |
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| UniRef90－UPI0003C28DDD | DEEETKPQGFIPLQGNQVSELAPNLDEPGKPGGSGDREKM |
| UniRef90－H2T5A7 | S DLEEREALKVVPLREDHKVQECKQSEDMMRD |
| UniRef90－L8－ | NTKIK T P L GFIPVDRCSVELLNTDDDFKKRKQGFVFCSAF |
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|  | DSAEYKLHR LLDLTEVHSCTPVELKKHDNT F |
| UniRef90－UPI000440CF34 | D DKEEVVLGSLPLLSFKlGAVQSSDNISRK FAFKVE |
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| UniRef90－E0V997 | SHCKGYIDLGEVVSVISATPGPGTPKKADEKAFFDLL |
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| UniRef90－UPI000383612E | DEKEEGIL GSILLPSFHISMLSVDDHIT RKYAFKATHPN |
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| UniRef90－UPI00033188D8 | CEQDREPLRTILLKDVLKTHECLVKSGDMRDN |
| UniRef90－T1HH24 | DAMEDSHCKGFIDLAEVVSVSQVAPQGAPKKTDDKSFFDL |
| UniRef90－G7DYE7 | NAKEY QLLRDVPVADISTCAEVQVKKHDFV FGI |
| UniRef90 | EKRGDKE PLGMI ILEGC |
| UniRef90－UPI00038BC8D5 | DS GEDTS CGHIDLAEVEMVIPA GPSMGAPKYTSDKAFFDL |
| UniRef90－F0Z675 | S PKDTTPAGVIPVNEIDISLKCEDS Q Q E G |
| UniRef90－V8NVK7 | HKKDEGKCR PLEVTKVEGAEIGMMDNSLGKPFVFKCVP |
| UniRef90－UPI0004572F7A | DEKEEAVLAS $\mathrm{S}^{\text {PLL }} \mathrm{F}$ SFRISSVQPTDN |
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| UniRef90－UPI0001CBBDFS |  |
| UniRef90－F0ZDA6 | TSKDKKPKGII QL N NVSVTRSYYKPN CMADK SNS |
| UniRef90－H2V2Y7 | DEKEEG\LGSILLPSFHVSMLSVDDHVNRKPLQNNPKQAA |
| UniRef90－F2U6N0 | NKAARGVI DL R T ClS lie EaHHEPEKSR YNFRI |
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| UniRef90－M4C409 | GNSDATPLGVVSLRRVTVQTCAPHEVNARNKCHFFK\｜VPP |
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| UniRef90－UPI000328AB8D | LKGGRKE TPPRGQ LLLDGCTIT CPCLEYENRPLIIKLK |
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| UniRef90－B0ES05 |  |
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| UniRef90－W5KRU7 | DRQDKECKGVIELAEVESVIPGTPTMGAPKNDEEKAFFDL |
| UniRef90－UPI0001DEA422 | VSEDLKDKKGDILLDENCCVESLPDKD GKKCLFLIMKCFD |
| UniRef90－UPI0003EAI | DSREEAVLGS $\triangle$ PLPSYVVSPVEPEDHDS RKYAFKATHAE |
| UniRef90－R9XJ80 | DQDDTANACRGSLNMSTCYLHL DSSEKLKFE\｜IGGG |
| UniRef90－H3C194 | HRRDEGKQR PLLAVRLEGAEVRADASLGKP FVFRCCP |
| UniRef90－V9FAK6 | A A MVA H S TIQ T F HVS GVS NWDGHATLMHYDHGFLLQ |
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| UniRef90－K2H3Z9 | GKNDTEAKGWIDLTPGTEVKDLTQPGQK K K F F S I NSRG |
| UniRef90－B4KJY4 | V SKSAFGMACLHGYKVASMSA NASGKKNS |
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| UniRef90＿UPI00021A440D | D DAPQGS D DKEVHQ A $\square$ A GAAKKG F F GFDLT |
| UniRef90－UPI000153900A |  |
|  |  |



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$\begin{array}{ll}V V E G H \\ L & \text { K Q G Y M K K T S K V }\end{array}$ REGF L WKRGRDNA QF LLKRRFVLDAREGLLLKYY
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N A P QVL
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K I K S F Y F A A D NIE DMNRWLSLMMNAAAAHSEQKR K VK S F Y F A S DCLLD EMNHWMNRLGGAADGYSPEEKD

LIDS S S RHVLLCAE T DEERDDSWVE EVLVRYVT GTYSEEP $V$ SP T K S CMMQA DS EKGLR QA WI TA V QN S I A T Y T K S K A Y ILAA D N Q $\square$ TMES WV K A L S R A
 K T Y D R LIF F LVAS S A V S M R $\square$ WMDVIVVTATDEHSR SSSHVKHVLCAEDDKERDDWVDALLQWID ETPDRTYLLSARQSDDDRSQWINVI QKVIDK D NKTRNIFVYHENGKE V VDWFNAI RSV KTYDRLYFMVAPTPEAMRIWMDVIVTGAEGYTQ K S FEDSS A S DKKKK QEWDQAI QTCLLQHLS TSSHERHVLCA ES DKERRDLWWVEALLLQWVVDYRDPE DLLE K T H D R LYYMVAPSPEAMRIWMDVIV T GAE GY T Q DAKPFVLTMLATTEAERTSWDEALRKV LS P SK S H M L Q A D S E EMLN S WI A A L Q N GI R S A IQQ Q Q Q GKNGLRWHLKKANHPDETNRWVWTMQNA DNDKKRN TTHERRVY LYYA DTGDSAHEWVKQLQQKVI FRSHNDGD VT T N R T W V L Q A E S E A E Y M K WVV NAALS H TVVS
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KVLDV SVLEPNPTEKGALVLVDVTQNFAUANGLLPYCFQIVERFALH DPLDAFAKGELFLGSSQYEVKPVVVPSTHGNCTWNYGISI

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F C

## VMM

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$\mathrm{D} \square \mathrm{V}$
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VK G GF LF K R GK H I K K WE K Y Y AV L T R D H DY FF E GVLSSKYTNLLQQ GWQNRYFVLLDPELSQLQYF
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K E GFL F A T SKP SK T WHKY WCVVL S G GQLL HEYS

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

L K E G Y $\begin{aligned} & \text { K } \\ & \text { I K K K }\end{aligned}$
$S \| N L S E C Q M T G F L Y K M G Q L N K E W K K R F F F I Q Q G T N G Y Y A$ Q GFL L L R G S GRK L WKR F F CF L L R R S GL Y Y S S K


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K Q Q D K D P L Y il I P I K N $\quad$ L





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QTPERTF LLKADSQELMLYWLQQLQ
KTSARVYWLVCQK LDEKQKWVQVIKSQD

SDT TFKFAAPSSEDLQLWVDNLE
NSYGDVYLFQASNQTDLENWVTAA HSASAS LKRQGKE
$\square P G E Q H F Y V R A D N A A E R Q R W L V A L$
H GVYYRSESEYTFERWMEVI RSAARGLNNKES
SDENKTWHFEANNAEERDSWITAIQQQLSSLQNSD
IHTSKSFTVFADNDKEKATWLMALDTQIKFLSQNE
TNQKTDTLQADSVPSRDEWVKAI RKVS

GRQLVLSAVTAGIRTNWMSAGRKAA
LREGGRLHLCAETKDDADAWKTALLEA
IPNEQYFYLRTENAAERQKWLVALGSAKACL
YTRAESFAVAAENEADQDEWYQAMVELQCRSKKH
VT TSRTWE VAE EKEYHK WLKEFCQAVK
KK S QY DK YLCCDDVRTDHQWINGDRDAK

LLMAKRTYYI S C ESEADKLSWMEVI GAAIES

KTVVV FYFAAENTQEMEVWMNLGNAALIQAARSS
SNTKTFKFAALTAQDLQEWVD

YS SNNSYFFQTNSLK DLQ DWTNS
GKKVHD $\ K F Q A P T L E E K E S W D K A L N E G \square N R G K N K$ HKKTLFYSFKAEDUSTAQRWMEAMEDAS ETPDRQFLDMCENERDQREW
FTQPRTHYFAVDNKEEMRGWMAAL
GDEDFYQAAGKQDRDGWAHAVVAVIRSLS GSKQD
EKTLY IQQANNCVEANEWREAJLCRVTRCNQRR

$\square$ D NGGTQTFHDASEVERQSWVTALEDAKAK
HPAYPVLLLQAENEEDRTKWINNIKE
DTPCRKLILCADNRKEMEDWI AALKTVQNR
EGQLHTMYI QCKNVNELNQWL SAU RKAS
TQDFCEYLLQAEDREDMLDWIKVI RENSKTGSEE
YKAGGTRYLYADSEQSRTCWMTA RLLKRSVSE

GDMLKK LRATDARARQEWVDGLRAIAE
FRTVNEYLLQSDNEAVSQDWYQAIKGVIHRLDQENP
$F T$ T Y YQAFTEDDMRSWIAA I NNALQS
GRSYVLTAPSDSEFKKWVNGL
TTARTWVI A AESDQDYQRWLKVLCGVVK

| $\square Q K K S Q Y Y L C C D D V R T$ |
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 MQDKST FLLLAADSEAEMEEWDGTLXNGUL
RVLERDYFLIGENSSEMDSWFNAIFAALKT
STPQKKYFLCAETPGAARAWVTTLHATQLVLKAHKE
VSNDCVWYLRASNPEEKQQWI DVLES
KY ILTTKY ILSAHDEPDRQSWI DAI RCAS
YEKSKSFYFAAENASDMSKWINHLIDSADNKYK


K $\square K S F F F A T D C D E E M N R$
$\square$ QKESQYY LCCEDPWLLHQWVT G $\square R \square A K$
GQGNQSLTVAANNEEEKERWI EDLLNMAI
M
YK RVYQATSEEDMNNWI LAA NNA ES
ISKKDKTYFLSAKSSEEKLSWMKLLNEVIDKKVGQNPE
QKTGKNCLISCENDEEFQNWFALLPQIQQUVLKREQ
NSFGDVYLFQATSQTDLENWDTAG HSACASL
TAKGKNYYILDAETEQDLVQWLEVLYQAK
DKKVI LSADTSGDRTNWVSAIRRSAGLTEEN
SAFRRNDLLATADKPEDKEKWVKALISUASGMQVVQKEE
$\square T P E R K F I F T C P S E K E Q R E W L E S L R D V L S$
RERTYHLQADSETADAKRWMLAALKREI T R VKKAK
GNKVSDDKLQAPS GE E K E S WDK A L N N E G I S R GKN K


## UniRef90 J9EHA6 UniRef90_K1P942 UniRef90_K0RVS8 UniRef90-X6NFW9 UniRef90_X1WF60 UniRef90_I3JGZ6 UniRef90 E9CGF3 UniRef90-W5N9R5 UniRef90_UPI00046B77BA UniRef90_A0A024JLC8 UniRef90_W4FLE6 <br> D















D QE D LK A E GVLLHLPA F QV
ESKEHGTPA KLMLDDVSSAMVE IVDDDDDTLAKLCFPPVKGG
DERDPDERAVINLANAHVEYSEDQEQMVRMPNTFSV
K SPECQQEHSCIPVGHDRAVERVDEGTFPQPHVMQVVAQDG
D S RE GRS VEAAP KHELI I V D G A I M Q P DP E H P K R D Y I F C L S
 RTPQDVSPLGTTEGRTY L Q CKDQHTMM


D P K S Q T P S S WK P GNS R P ES S V V D L R H W A D D L




Y EK DK Q QRGAFY SNAY L MQNA







 D E

Q
T


GSAAAFGAGDSPAGRLP LAGVASVA
QNDLECRQYFTTRNPDLVVKDALQLR
D R


D Q GP F G T N N E T Q L D M K I DI S Q S L A H P V H D H M K R D H L F C L S



K QE E R HFR GVI T L E ECN I E E A A D E E EPPAAK SF KL T S K V P Y





EYK P E K HL S E E DL K N A I S I H H A L A I K A T D Y EK R P NVL KL K








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[^1]UniRef90_UPI00038C3D9B UniRef90- UPI000440E6B1
UniRef90_G5B0D8 UniRef90_U3K8B2 UniRef90- G7YNIS UniRef90_G7YNI9
UniRef90_Q6C9X UniRef90 UPI000387B71A
UniRef90 H2T1Y0
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UniRef90-M5XTN7 UniRef90 C0PG99 UniRef90-UPI00034F8006
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UniRef90-D8QSV0
UniRef90-FOWSD2
UniRef90 X6NSH5
UniRef90-B2VZ94
UniRef90-T0RZP3
UniRef90_LOPB30
$H$
lupas 21
lupas_14
lupas_28
jnet
conf
sol25
sol5
sol0
jhmm
ssm



[^0]:    lupas_21
    lupas_14
    lupas 28
    jnet
    conf
    sol25
    sol5
    ol0
    jhmm
    jpssm
    

[^1]:    lupas_21
    lupas_14
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    sol0
    jhmm
    jpssm

