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 MQTAVFGLTNYLDFDDNTDVKD GWLKCK DSNAKDL STA AVNGGEHFLSVAPKVRGEADGHITCQLVEVVSEQE ATATINSLDVLQKVEF ESVEAEEGNTSCIVKDDC ALNFINSVEVLTGKSPKMELADE GGFLKVEKPQELAPHQ SLTGVFASEKLCNLSKQTTQPEE GLLSLEMPTGLDEQQ SLGGAVNA DYSLLNVELDVVEMKED GGYLFCSVPAGLLADT A QTALGALEEVCDLKNKGYVKD
A TINTINSI DKFTEEKTSLVSDEESGGFIDYKIIDGRP


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## S <br> \section*{A}

## A Y T A A GA L G DL I G $\quad \mathrm{K}$

 SDAAANGGEHFLSRAPQVR
AQTALLGGLGQYLHRDMDYRVKSG


T D S A V L G L E R H L G RK LT L Q A G S G G L





 I K S Y GLE N GYMDCSDP MINTVNA E Q Y T GDKF SLETEESSGSMECM VQMA A N GLT ECA G L P A DL S VHE G D $\square$ S L R C
TQTAVLGL DRHLRARMHLDIA S GN D F
$\begin{array}{llllll}\mathrm{T} & \mathrm{Q} T \mathrm{~A} & \mathrm{~V} & \mathrm{~L} & \mathrm{G} \\ \mathrm{S} & \mathrm{L} & \mathrm{L} \\ \mathrm{V} & \mathrm{V} & \mathrm{N} & \mathrm{L}\end{array}$

 $V Y S T E N A L D K L T N A R Y S Y G V D D$










 VINAI NSMEKLAEEDMEI KA EEAAG円IRCRIKRTL

 VINTLNAI EQFANEKTDVVAKD
 AI STVNSLDALAGVIPNDETNDDEGGYLHVNLDDSMTQEQ



 VI NTVNSSLETFTDDQFEV A YL V A N A I T D V V Q i K A D L S V S E

GDMSLRIP EKAVF



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UniRef9__R5ST7W

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 SSDGKV $\mathrm{V}^{\mathrm{G}} \mathrm{S}$ RF E MI CVGKNACOT MICVILGKNKAGQFTSFSVNGHA MVCVVFLTPPE
$M T K V L V R K T S T$
$M T R E F F$ M门TVT THRS ML K N Y Y MUH DMKRDKHKHFCGFTVSGHAGYNEAGNDV MTNTTFYПKSDKSIIGFKSKGHAGFGVRGTDVVCASVSAL MDKVTVYKKANSDTMGFYVSGHAGYADSGSDIVCSAVSAL MTTITITKSKNGEYKEVICKGHAGYADAGNDI MПKAHMTYGTGNRDTGFELTGHAGYGDEGEDI MI NAT YVNSRHAYMGFDLEGHAGYSEAGTDI


 MI K KYHYTDEFRRKGFTIKGHADFAP MIKIKYHYTDDFRLKGFTLKGHAEYAEYGYD MTRCEF D MTKVRIGKK HNGSYCSV
 MIEATVKKKNENIVSFKIEGHAGFDEYNKD
 VCAAVSVL
 $\checkmark C A G T S V L$
C SAA V SAL VCAAVSSA MTNIIFYR KSGDIIGFOSKGHAGHGMRGTDVV M NVSLFEDSNKNVVGFEVSGHAEYADPGMD MIRVTVEKSNDTYVSFRSKGHAGYAPAGED
$M V E V R G R K N E R L V S Y S T D G H A E Y S E P G S D$
 MI HVRIQQNQHEDIVAFVMSGHAEAGPHGQD MVKVKFS TQDGKFVGVECSGHCEYAEEGADI MIKVKYYQNDDFKLKGFCLKGHADYAELGSDIVCAAVSS MVSVTVFVDSEQHYTGTQMLGHAGLKDEGQELVCAAVSAL MTTVTII QSKDGEYKGFVCAGHAGYANSGSDI
 MI KVKIKKR MISVKI QGKNNIATEIIVSGHAGFADHGED MVTVKVYRKEEGKITAEIEIIGGHGYAPSGQDLVCAGVSVL MVNVVFKKENNDVVFVKMSGHAGSTDQGYDMVCSAMSAV
 MI QITYKKPDNQYKGFQVIGHADSVEEGAD
 MTRIEVKKQNSDIIGFEVKGHAGSDEYGKD M R V ${ }^{\top} V T R E A D G K V S G Y Q V E G H S G T A A H G E D$ MIKVKIGRDTHGMVQEDTITGHAGQGPYGHD
$M T K V E F F T Q D D R I T G F C S G H S G Y G E A G T D$ MTVVKIFRKNGNITGYRANGHTSYSDYGTD MDEVRSVDKKG PLSVQARGHSGYAPRGKDV MDTVTFFKSAD YFKGGKUSGGSADLADA


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 THAINSIDERTDSRYGLEENAEQG CLLTANT TEVHLPAAAEMKD GGFQQRLSGDTAP


ADNFYNSVETFTDDGF AYNGINALDELVGIKN YGIESGYKKTLP PLSKER

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 | $V$ | $N$ | $L$ | $T$ | $E$ | $C$ | $T$ | $V$ | $N$ | $D$ | $V$ | $C$ | $G$ | $A$ | $R$ | $A$ | $K$ | $V$ | $R$ | $V$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| $L$ | $Q$ | $E$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| $M$ | $L$ | $P$ | $L$ | $L$ | $S$ | 1 | $Q$ | $E$ | $L$ | $L$ | $K$ | $A$ | $E$ | $V$ | $K$ | $F$ | $N$ | $I$ | $N$ |

$M H A L L E G L R D V L G D Q D V Y E I D E E E P$




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 VISNTVNCLEDLLHEDIEVVDYDAVNGGDLICNFRSD IV Q M T A N A L T E V V L K L K A D V S VVG E N V I F T A DN S L E S V A K Y E S E L N I K M A E N G Y M S F S L L I N D N D P R VIN T I N A L E E L A G E T F K T TV S N E K D G F I R C D F E T P L





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MTAVNVRKG
MITVTVSKNEDGTYNRICMHGHAGYAEYGQDi MVTVTVNTVQD EMAAVETSGHADYAEHGQDL MI RVDFFTAKDGGLNGFQLKGHSGSAASGEDIVCAGVSSI MIKVSTHQNKDGKINGFRCKGHAGYAESGQDIVCSAVSSA MI SVQLARAKDGSLASCRALGHARFAEKGNDIVCSAVTVI MIRAEVRRSGTGEIHSLRVRGHADAGPYGHDLVCAAVSAV
 MIKVHLIK KDDQYISLEVVGHANYAEH
 MIRAEFY
MTHEIY EKQGLLTG
SHNGEVG RATFYF TAN


MIVKVEMVK NNNDESVSTNGHANSSE

$\qquad$ FKKK

MI KIRVWRRANGGLSGFSLSGHAGWGESGSDVVCASVSSA MILVDVTFNADGTPTRLLMTGHANRGEYGSDIVCAAASAL MIKVQIKRNNKYITQUKIKGHAQFGEYGKDLVCAGVSAV MTTVSFL VEQSRIVGFDVQGHSGWGDEGADVVCAAVTSA MVKVVFYKSGNN YYGFREYGHAGFDEAGKDIVCAALSAM ML TVYYDEKGKDMSLQASGHAGYAPKGQDVVCAAVSAML

 MI KAAFA MI R R V $\mathrm{H} V \mathrm{~L}$

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\section*{A <br> | S |
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## $\stackrel{M}{N}$

 MII QVV VI KEKNQQI DDIEV ${ }^{20}$ GHANSAEHGKDLLVCAGIVIST MI KIVNFIKTENGKIKKVEITGHADQAESGKDLVCAAVTAMI MI DTRYTQHDNHIISGEMSGHAGFAESGKDLVCAAVSGII MI NINFTKKKSSEVVSRVNGHAEYYDRQEDVVCGCVSNLI M| SVLLERGTRNEFFSVTASGHAETAPRGFDIVCAAVSVL



 MTHVKFILEEDSPVGYVAEGHSGAGTRGNDVVCAAVSTL MIKVILTYHED QVSSI QITGHAQSGPYGQDIVCAGVSVA MTKVVISR RNGLIYALSVEGHSGYAPSGEDVVCAAVTTML M N I T TKDDDNGKHDS:KGHAQSAKYGNDLVCCAVSVL
 MTKVTFYQNSENQCVGFTSEDHAGYAQEGEDVVCAGISAL MIKITI YRNADKQYVRLHCDGHAGFARAGEDIVCAGVSAL MINVTIKQGK KQYTIASGHAGYAPKGYDIVCAAFSVL MTVVYLSRSRNGQFRSCRAEGHSGYAPSGSDIVCAAVTVL MII IAEFLRSG KMWTGFRIKGHAGFAESGHDIVCASVSSA MLTVTFHLAAESQLNGFSLVGHAGYAKPGQDVVCAALTAGL MIKVVFYKS NGNYYGFRESGHSGYDEAGKDVVCAAVSAM M品VTVKY KANQAGYRVGHSGTAARGQDIVCAGVSAL
 MITCKFFK ENGFFEKFLITGHSMYAQKGKDIVCAAVSAL $M V T V K V V C A K D$
$M$ MVNVEVKRQSD FLQEVLI HNHAEYGEHGQDLVCAGVSSI
 MINIKIFK RNANYIGFESFGHAEYSD GDDIVCAGVSTL MI N: NLLINKKDEDVGFEDEGHANYDEYGKDLVCSAVTIL

 MIEYKL IKKQKGRLGLMVRGHANFAGPGKDIVCASVSSA MITAKFKKKNNQIYWYQVTGHAGFANDGNDIVCAGVSVL MVTVDGFRKDGKIVKVKSTGHADHGSYGNDVVCSAVSVY MIKASFKKDYDG FYAYEVSGHANYAEKGKDIVAAVSM MI DVTLKITKKKRLGGFAIKGHANFAEKGKDVVCAAVSSL M SYFFQKNANNHIIDAEVSGHALYAPKGFD




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 $G \vee R$ $\square^{20}$

20
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MTTVR』

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M H ITTKYDEWFDIGQDEGHAGYAEKGKDIVCAAVSTL
MITA FKKENNQYVSYSI SGHANFAPKGQDIICAGVSAL MVKAKIVQ KDNKINSFVINGHANSGDYGHDLVCAGI TAl MTKYEFKKVK TVVELTVSGHANYKKHGEDIVCASVSTA MUE YFFQKEKNIIQQIEIKGHALYAVKNKDIVCASVSTA

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$M \Pi K \nabla$ NKNGNV $\square K \square T G H S G Y D N A G$

 MTTATIWK NGFVNRICTKGHSGYDENGRDLICAAI S MLS DQI WWSKEKFVELNVVGHAGYAEHGKDVVCASAVS VVYNSDGVIIKVTI EGHAGDKKSGEYVCDALSAL

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