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$D V S W L V F Y T G V R S S E Y A E R K H$
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 EQFNDLDETVAKRARHVITENDRTLQATKALSENNIRRLMG A QF NQLDPVVARRVRHVITENVRTLEAAEALARGDLKVDG
 GKVSELDEVVFKRARHVITENNRTLEATDALDAGBMKRMG AELTALD GEAAKRARHVVTENQRVLDAVDAL NQGDDSRGG E QFNELDEVVAKRARHVI SENARTLEAADALRENDMPRLAA
 EQFHELDPLLRKRARHVITENYRTLDAAQALSQHDMQRMA A QFA QLTPLVAKRAKHVISENDRTLAAAKALTENNUPLLS EQFEDLDPVVRRRAKHAVYENARTI RAAEALKRNDLQLGG

 D T F N A Q DS E V L L R R R V R H V I T E N N Q R V L E E F E L H H D P V R Q R R A K H A V Y E N Q R T I R A V E A L R E N D V E L



 DEFEQHDKVLEKRAKHAVCENQRTLKAFKALQEGRLEDFG EEFEANKEVRQRRAKHAVYENQRTLKAVDALKAGNLLEEGG K HKD QLAAVNVYKRAH HVVSENERVL ES QKALLQAD DMS K FG A E F HELLDPLVAKRVRHVLTENARTLEAADVLARGDLARLA ETEEQYDPAL $\square K R A R H A V T E N E R T L L A K Q A L T E G D L E E F G$


 AARSRLGEEVYYRRARHVI GE DARTVQAAQALQDRDYRMF G E QFPELDKTVAKRARHVI TENYRTLEAAQALNRNDMKRMG

 E T F E K Y D P I K E Q R A K H A V A E N Q R T I D A V A A L K E D N I E L F G
 KHKDEMSDVVYRRARHIITENARTINAAKALTNNDIVTMS E Q F H Q LD P L V A K R A R HVI S E NARTMLAAADALAA GDL QRMG



 STFDEYDEARLKRARHAVLENQRTLKAEKALEAGDLERFG
 E QF E I S S D E TN NQK R A K H A V Y E N A R T I D A L L K K L EVKS RLD DVIFRRARHVI EETERTLQAAEALRRGAYKEFG



 EKFEEVKPICRKRAKHAVYENQRTIQAVKALKENDLETEG V $K$ AQFDNFKDVLRRRARHAVTENERTLKAVKELKHGNLEAFG $N L F F E L D P V V A K R A R H V I T E N E R T L T A A D A L A A G D L L G M G$



 A MKESLTKTCYKRARHVI SE DARTTDAADALQRGDYKAFG
 SEFEQNDEVLEKRAKHAVYENQRDLRAFQALEHD GLASS G NKGLIKNEVNRKRAKHAVYENQRTI KAQKELSAGNLEEFG A QK DKLP EVVYRRAHHVVSDSERTLQAADALRTGDAKTGLS EQ L E A S E E KVYYE R A L H V V E E N N R V L H A V S A L Q E R D Y V R F G L
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## QUERY

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| $Q$ | $V$ | $R$ | $K$ | $A$ | $V$ | $E$ | $N$ | $Q$ | $Y$ | $H$ | $T$ | $Q$ | $T$ | $G$ | $L$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| $T$ | $K$ | $E$ |  |  |  |  |  |  |  |  |  |  |  |  |  |
| $T$ | $R$ | $A$ | $A$ | $V$ | $A$ | $H$ | $E$ | $Y$ | $R$ | $Q$ | $T$ | $G$ | $G$ | $L$ | $Q$ |

[^0]| QUEF | E P S D GVGT |
| :---: | :---: |
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| UniRef90-E6WC55 | KASEGA |
| UniRef90-UPI0003A993D8 | QAKDGASL |
| UniRef90-128B2\% | TPSEGA |
| UniRef90-х2H | KPSK |
| UniRef90-UPI00031A2003 | QAKQGAGL |
| UniRef90-Q65UV5 | TASQGVHL |
| UniRef90_E8MD08 | QAKAGAGV |
| UniRef90-UPIO00319D245 | RATDGAGV |
| UniRef90-R8ARW9 | QAMDGA |
| UniRef90-10DQF8 | QPSDGAGL |
| UniRef90-UPIO0035E8CCE | T $\square$ GDGA |
| UniRef90-UPI0003B77550 | QAKQGA |
| UniRet90_A8AJ37 | KPSQGA |
| UniRef90-F3BDN7 | D 1 GDGA |
| UniRef90-UPIO00255D674 | DVGDGA |
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| UniRef90-G2SN74 | E |
| UniRet90_W3Js 7 |  |
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| UniRef90-H2MWA6 | GARV |
| UniRef90-UPIOOO2F9B0A5 | K\GDGA |
| UniRet90-L1H15 | HASAGAH■ |
| UniRef90-UP10003B650 | RPSQ |
| UniRet90-EONH81 | K V ${ }^{\text {d }}$ |
| UniRef90-M1 N4N1 | N I SGG |
| UniRef90-W0RRO3 | S G |
| UniRef90-R6NM47 | a |
| UniRe990-H1PV63 | K GDG |
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| UniRet90-A1JR | QASQGAGL |
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| UniRef90 A3QFA5 | QS |
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| UniRef90_C3FB63 |  |
| V6Y |  |
| H3CAN5 |  |
| UniRef90_R5YLP9 | S $\square$ GD GP S |

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 ETYEKINDETLLKRARHAVSENQRTIKAVDALNKNKDLEKGG EDFDANKHVIRKRARHVVYENARTLDAVQSLQEHDLEKQFG QSFDQYDANRIKRARHAVLENGRTMQAYEALQAENVPEFG
 HQDLIEDEVKRRRARHAVSENERTKKAHRALAENNFNQFG NKHLISNPDNKKRAKHAVYENQRTLKAVEALKNDD HTEG AALARDPERMKRVRHVVTENDRTRAALELVKGNLAGDA HQDKDTDEVVRRARHVVEENQRVLDSVEVLKNNDLKQFG KESKFDDKVVYRRAYHAVSENERTKRASEVLKSGDINAFG VKHFUKDEEQLKRARHAVRENERTKEAVSVLKAGN|EKFG AHKNEDTKEMQKRVKHVIYENDRVKESEKALKSGDVKRFG QTFDEYDENRLKRARHAVWENQRTLQARQALEAGKLEEFG EENNFSDQTVFNRARHVIGEDKRTTDAATAGQKGDYDLFG DDLIEVDETLFRRARHAI TENERTLKAKSALETNDLDTFG ENRAQLDDLAYRRARHVLTENVRTLSCAEALKAGDMPRVY DMHYITNDTNRRRARHAVTENQRTKDAYQALQQDDFVTFG EQFASLDETVRRRAKHVVEENQRVLDSVKSLRAGDLEAFG EALDRLPDDLVPLVRHVVTENQRVLDTVELLRAGRLADDG AHKNNLNETTFKRARHVITEDQRTESAAQALKKGDYNTFG NDDEVVLPEEMDKRTRHVVTEDKRTIDAVEALEKGDLKKFG AEFDENDEVERRRARHAVYENRRTLEAVEALEQNDVKRFG EEFEKLNDTVYRRARHAVTENERVNEADEVLQKGNLINRGG DMLNAIKDEDYMRAEYVIGEKQRVLDVCDALLKGDYETVG AGLSELDGVLLRRARHVTTENDRVLAVAELLRAGQVREVG EEFENNDDIRRKRAKHAVYENVRTLKAKEALENNDLKLEG QEKTGLDDLAARRARHVITENARTLSAATALTAGDMPSLS ESKNKMTDEVFRRAKHVITENQRTIDAAEALANKNYTLLN GALGRLSGDERRRTQHVVTENHRVEALI GLLRAGAVRE G HKDADKDDDRRDKAKHAVYENQRTIDAVAALKAGDLTKFG ARWGEVEPVLERRVTHCVNENQRVLDAVDALRAGDLARLG TEFGQYDPIVYRRAKHAVHEHARVKQALDALKAGKLTTFG ADKASLDAEVYRRARYVITEDKRTTEAAEALKTGNYGQFG ELEAIVEPVAKRRAMHIVTENRRVLDTVFALEKEDLDAFG DDREVLDDETFRRVRHVVENQRVLDTVRALREEGPRADG EQFEQHKHLLQKRAKHAVYENERTLEALKALKAGDRTEFG DDFLGLDEVVRKRARHVISEDKRTRDAVDALKAGKLEEFG DQASEISDIEFKRARHVITENQRTLDTAQALVDGDLKQVG GNLSDLKEVLVKRARHVITEDKRTQDAAEALKNHDFAKMG RDLSGLDPVDVRRARHVLTENGRVLDFVAALRDSDFSAAG KYEAELAPKVYQRARHVISENQRVVASKNYLAANEMDKFG


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A GF G GCA|V|AlLV|APGQAEAFVQG|V|HAKYTQATGNTPQCYVIC A GFGGCAIA AVHQTQDESFKENLTESYTKEFGFPPSFYEA
 GGFGGCVVCFVESDKAEKVMKALEKEYKERTGDECSAFVVT


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A GF GGGCGIIALVVKESEWA DFEASVVKAA Y LAAEDGYATDDYAA

GGFGGCVVALVDHALTDAVVEVVEKQYAEKTGLEA
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 EYRLVIANSNVSGSKYNERRAECEAALAVVKVQALAEI EP E LRVVVI DTRAKHSGYGDRRRGTERAAGLLGGNSLREI ET ELAAVVVLDTGLLDGSPHERRAQCQNAARTLGVPALRDATE DYHLIIMNTNKKDSKFNERKGECDAALAEL QKHSLGTRED YKIVI LNTNLKDSKYNERRAECDEGLSTLKINNLGELLQE LVILITNSNVRHEYPDRRRQCETAASLLGKDSLRDATE NYTLVIMSTNKKHSAYNERVKQSQEALHLLGLTSLSDLDA VAVLVVDTKVAHSGYAARRASCERGAAALGAVSLRDVPD DYQVLLLNTNVSHSAYNTRMAECASVVEATRKTSLREMDA LAALVIDSNAPHGEYAARRNSCEKAAALLDVPALRDVDD VALLVVDTRVKHGEYGKRRAGCEKGAALLGVDALRDVPH DYELLLVNSNVKHSAYNDRRVVCEQVSAQLNVPFLRDVTK LVVLITNSNVKHSEYPVRRAQCEEAARMGGSKLRDNE RVVLGNTKKKESYNE YKLVLVDSCVKHSAYNKRRQSCENVCAAIRKNHVEF L VSVLVLDSAVKHSAYGQRRKQCEEASSMMGVPSLREATE NYVLAI INTNLADSKYNERVAECQTAGKLLNEEYLCELDA DYSLLLFDSNVKHSAYNERREQCEEGIAIIKITSFRDCTE
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 DVGVLVVTVNS LALLVVDSMVKSEYNERRAEVEEGSRALRAANLRDARE DYRVI SSEVKRSAYNERRAQCEEGVALFKVRALRDVTL LELVVIDTGTRHSEYADRRQQCQTAASLLEVHSLGAVID NSEVVMDNKGGYGERRASCEKGAQLMGVPMLRDLSD

 LRLLVVDTQVKHGEYGKRRAGCEEGAALLGVTALRDVPY DIVILNTNVRHSEYNDRHAECLRGAEAFGLASLRDLSE DTEELVIDSGESHSGYNKRRAECEESARILGVKALRDIDP DVRVVI ANSMVKHGEYGNRRDEVEAGQEVLGRELLRDATE DR V N
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 पAD DGDEEEVVRRTVRHVVTTE

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80 EEFEKDSPVNQRRARHAVAENERTVLAKKALSSGDVMARGG EEFEANDETLIKKRARHAVYENNRTKVAQKAFVAGNGTKFG QYQQVLS SVQTYQRCLYYLQENERVLOAAASAGLNKNDLREFG EMKEGMSETVYKRCSFIVVKENDRVLAMMVDAVLKKNNLLDEVGGG A WPYVTDTTLRRRARHVVLENQRAKRAVEVLRVGDTARFG TNQARLDKTTYQRARHVI SENARVLAVKQALLNHDTARFG ERLRQLETVFDNRVKHVIEENARVLASVEALKANDLHRFG A QEK E ADLPRLSEPLGRRVRHVVTENARVLATVEALRKGDLPGLG EMLDQHEPDIYRRCAYVVQENARLLAGVVEALEANDIAAFG E QFEKHDKVNCKRARHAVTENQRTLKALDELHAGNLAEFG
 GQWSEAEALLRRRARHVVMESHRAREAARLLADGNLEAFG【DLFPLGDLYYRRALHVVSENMRVLEAMKMMSNSNFENLG A A A D RLPADLVVRVRHVVTENRRVLDMVELLRANRLAETG RRGGMLDDTTLRRARHVITENARTKEAAEALRSGAYERLG
 KDFYLDNRLDRKRVQHVVEENKRVIKAQQALSSNDLKTGG EARGRMDDVTYRRARHVI EEDQRTVQAAKSLERGEYKEFG A LARVEDPGLRPLVRHVVTEDRRVERVVELLETGDARAGG YDYLINDAVLDRRVRHVVTENERTREATKAMQVNDMERLGG
 EVTSEITPDAYKRARFVLEENERVNKAVEALKSKDLNTFG ALARLPDDETRRRVRHVVTENQRVYDTVDLLRARQUQQDG
 TARKDLSTEDYDKALYYIVVEENERVLAFSKAMVDEDDLGTAGG

 AKRVWLPEEDF LDRAEYVI GEVQRVLDVADALERGDYETVG SFREQLGDVRYRRARHVIGENARVNAFANAGSRGDWDEAG EKHVITTDETILNRATHVVKENDRVHLAAKALNSGNLEEFG
 AQQDQFTTVVFKRVHFVVKEINRVIKACEALDKGNIELGG
 TDLPRLPEPLNRRARHVVTENQRVLEGVAALEQGDMQRFG
 A GLPRIDPVDARRARHVLTENQRVLDCVAALASEDFGSVG EARPDI SEEDYMRAEYVIEEDQRVLDVCEALERGDYETVG ATLPKLPEEDRPLLRHVVTENDRVLTTVDELRAGRYAEIGG
 DHLKALEEVLVRRARHVITEI QRTTEAANALKANDFVKMG A HLIPEADETIRRYVRHVVTDNARVERTDALLDAGDVRAVG EYKDEMDDVVYRRCSFVINENQRLDDGCAALDKGDYVTFG DEQDRFEEVIYRRARHVVSENERVHKAAAAFRAGDFGYVG
 LKY KI KDNRVLNRATHVVYENERVKRAYYLLSKRNLKEFG KAFE!LPSPLKERVAHVYNENKRVLNFVNDLKNNKWENDP DESRMDDEVQRRRVRHVVTENARVLEVVRRLRDGDLDGIG
 EVRSEVSKEDALRAHYVI GEKERVMEVSAALEKEDYEEVG EVTEKLPEPLRRRARHVVTERNRVL EEVLQGVSSERFG EATEDLPEPLRRRARHVVTENNRVLEVLQILETRHLQRFG EVRSEVSEEDALDRAKYVIGEKYRVLEVCVALEEGDYEEVG
 EARGSMSENAFDRCRHII TENVRVEEAATALAAGDMKRMG PEALLADDEVVARRARHVLTEIERVTSCVTALSSGDVEAFG


 AGDTVADPMLRARARHVVTENARVREVVADLQSRDPRSTG RAAGILDDVTFRRVRHIVTENQRVLDTVRTLREQGPRAGG SQLDVLPPLLRRRARHVVSENARVQEALGA DAARFG AVKDKVSAEDYQRAHFVLGEKERVLAVCDALVAGDYETVG AEQDWLD VVSRRVRHVITEDSRTTTAIEKLQAGDFEAFG AE LDDSADEEI RRLVKHVVTEDERVEQVVALMSGADARETGG AEAPALDPVVRRRVRHVLSENLRTLEAAKAMAANQPERLG QAVEVLPEPLRKRARHVVTENNRVLEAVKSLSAMAKRLG ACRQKMSPESFARCKHII TENARVLIAAREALLKADMKRFG AHREELTTTQYARARHVISEDGRTAAAASALLDGDDRRFG EQLPDADQTLRRRVRHVVTEUARTRQCCDALEHGDFQQVG YMLDVLKEVLWKRAKFVVEENKRVLTAAKEAAMKGGDKETIGS EDNRLDVNRRRARHAVTENQRTLDAARALSLADLRTFG
 DDLSELTDEQRRRVRHVIAENARVRSSVELLEDNRIRALGG DQHNVLPAVLYRRARHVI SETTRTVNAHTALVNKDMKTLS $\begin{array}{llll}D D L D E D D D I & R A R A R H V I S E N D R V L A F A A A L A S D D R Q T A G \\ E A E H D P D G V L R R R V Q H V V T E I A R T A E F A E L V R A G K V D E V G\end{array}$

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 RILTAGHASLRDDFRESCPELDAVDTALSAGALGARMTG TLLYQSHEGMRDKFKISCKELDFLVDITVSNHILGARMMG QLMLQSHSSLRDDYAVSCPELDALVNLAMNMDVYGSRMTG ALMGESHRSLRDSYEVSCEELDFVVETAESVDALGSRMTG RLMVEANDSI RDLYEVTGRELDIMTAEADKVDVIGARMTG EMMYQTHFGLSHLYEVSCPELDFLNKLARKCDVTGSRVMG VAMRGSHASLRDDYEVSCAEVDTLVSLCDRDPIYGARMTG RLMYASHQSLRNDYEVSGKELDAVVDFCTDEHVIGARMTG KLMFETHEGLSTDYEVSCAELDMVNTLKSEDVVGSRLMG QLLFETHYGLSQEYEVSCAELDMLVDTAKDEALIGARVMG QRMFETHEGLEKQYEVSCKELDLLVQLVKNQYVLGARMMG ELMVASHRSLRDDYEVSTPELDRLVEAELRHGAVGARLTG QLLYQTHEGLRHKYEVSCPELDFLVDFAKKTDVLGARMMG ELWTASHASMRDDFEVTPPHLDLIADTAVRAGALGARMTG QKMYETHEGMSKLYEVSCEELDFLNECARQHGVTGSRVMG PLLINASHASLRDDYRVSSPELDLAVDSALAAGALGARMIGG ALLYQSHNGLSTQFKVSCKELDFLVDKTKNNNTLGARMMG RLMYLSHESLSDDYEVSCAELDLMVEDARSPGVYGARMMG QLMTESHRSLSEDFDVSCHEVDLLVEATLGARVLGTRMTG PVLTEGHASLRDDLRVSCPELDLVVEAADAAGALGARMTG QKVFGSHEGLSKWYEVSCKELDFLVDDARKNPVLGARMMG ELMNESHWSLRDDYEVSGPELDQLTELLRDMPVWGARLTG ELLKASHASLRYDYEVTGLELDTLADAANEPCCLGARMTG VLMNASHDSLRNDYETTGLELDTLHEEANKAGCLGSRVTGG KLLAESHFSLRDLYEVTGKELDALVGEALNVSCIGARMIG PLLTASHVSLRDDYEVTVGELDTAVDTALATGALGARMTG ALFTASHASMRDDFEITTEHIDLIADSAVAAGALGARMTG ALLYQAHEGMRTEYEISCKELDFLVEFTKKDFVVGSRLMG KKMYETHDGLSREYEVSCPELDFLVDVARDCGVSGARVMG ELMNASHASLRDDYEVSVPALDTLVELLQKPGVFGARLTG GLMNASHASLRDDYEVSVPALDTLVEMLQKPGVFGARLTG KKMFETHYGLSKEYEVSCPEIDFLNDVAKSCGVTGSRIMG DAMLAAHDSLRDDYEVSCPELDAAVELAAEPGVYGARMTG KLMIESHCSLRDDYEVSCSELDELVKLAMEVDVFGSRMTG ELMAEAHASYRDDFAASCPEVDLLVELAAAEPGLIARLTG ARLTESHASLREDFEVSCAELDVAVETCLAQGALGARMTG ESMQASHASLRDDYEVTVPQLDVAVDVAMAQGALGARMTG KLLNDSHQSLSKDYEVSCAELDFLAEKAQCDFVLGSRMMG ALMNASHQSLRDDFEVSCDELDCLVDDAQRTDVLGSRMTG KLMYGSHDSLKNLYEVTGBELDTIVDYCKDSNVIGARMTG PILTRGHGSLRDDFEVSTPQLDAAVEAACDSGAHGARMTG SLLDASHVSMRDDFElSVPELDLAVETARGAGADGARMTG QLMNASHASLRDDYEVSTPEVDRLVELLQAHPVYGARLTG QKMYETHHGLSKEYEVSCEELDFLNDVAKENGVTGSRIMG TLMDASHLSLLRDDYESCPELDLAVDKSLEQGAIGARMTG PVLVAGHASLRDDFRISCPELDLVVDSSLAAGALGSRMTG LLMNESHESLRVDFEVSCAELDMSEEARQLGCYGARMTG ELMNASHASQRDDYEVSVPAVDTLAADLQSPGVFGARLTG ELLMAEAHASFRDDFAASCEEVDQLVALARQPGCIGARITG ELFNQSHASLRDDYEVSSPELDLAVAAARAGALGAMT ELFNQSHASLRDDYEVSSPELDLAVAAARGAGALGARMTG LLMRASHLSLRDDYEVTCPELDALAELAWQPGVIGARMTG KLMYASHEGLSKKYGVSCQELDFLVDQSKSLFVIGSRMMG DLLVASHESLAHDYEVSCAELDVAVAAAMGAGALGARMDG RLMQESHHSLRDDFEVSTPELDFLVEUINKVNKGGVRMTG PLMLASHTSLRVYEVSARELDLVVEAATSAGALGARMTG

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GGFGGSAlA 130 EASAlVDT $\square^{140}$ KAlIKEAFAHAGYRAPRTF 160 AGFGGSAI ALVAKSEADNFAAAVGKVYDEK円GYQASFFVA AGFGGCAVALVAHDKVADFEAKVGAEYEKVVGYPASFYVA


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A GF GGCVVALVRREALAAFIDATCSAYLAETGRRPHAYPC

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L VML $\operatorname{V}$ A N S M V $K$ K H S A Y R D R R E E V
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A Da OTDDEO
KRTRHVVSEIARTHAFVELLEGKRLLDVAA AVLPQLPEGLRPLVRHVVTENDRVRQTVALLESGDVAGLG SELDAVDEVVYRRARHVITENERTVAAVEHURARQYAEAG ACRSLPAESYLRCKHVITENARVREAAEAMKRGDATALGG SHADALEPVHYRRAKHVVTENERVNVARDALSAGDFERVG EHRDEMGDVVYRRCLYVVQENFRVEAACQHLKDGDLKAFG DVLAALPEESRPLVRHVVTENARVDSVVRLLRQGRYADGG DHLDDVDESVVRYVRHVVSDNHRVERVIELLDAGDLRAAG AVKSEVSEEDYTRAHFVLGEKDRVLAVCEALEKGDYETVG ENKEKLDAEVFRRADHVINETEYTLEFVEFLKEKNYVEAG ALGR DDETVRRRARHVVTEDRRVEEFDRLLAREHLDELGG EVKEEVSQVVYTRCKYVIEENERLLKGCELLEKGDDKGFG QYVLPVNELVYRRSRFVVDEMERLNDACKYLEQNDIASMG DDRGRMSDASYRRCRHIVTENARVLEARSALFGGDMRKLG EASGLLDEVTFRRVRHVVTENDRVLRTVELLESEGPGA1 G Ahtoagrpral g DAASRLDPVALRRARFVIAENARVDAVAEALTRGDAEAAG DAIFRLDPTLLKRTRHVITETTRVRGAVKALRSGSWAQLG SCKPQLSKTQYNRVLYVLQENARVDEAADFMKSGNLQAFG DTLPDVGESQRRRVRHVVTENERVSRVVARMRAGELEAAFG TDLSALNEEEQKRARHVITETARTLDFVELLEDGALTDDGG ACRDLMSSASFKRCRHIITENARVVKARAALDQGDLERFG EAVGGLPEPLRRRARHVVSENNRVL EAVRGVPAERFG DEVRNLPDELQPLARHVVSEEARVDDVVELCDAGDFEK円G EVKA IVGEEDYNRAHFVLGEKDRVLAVCDALEKGDYETVG KELAKLKDEDSAADYYVI QENHRVKLFAKAIRNNELKKLG KARPQMRDNVYRRCRHIITENSRTVAAAAALDAGNLKEMG AGLDAADDETRRRVRHVVTENQRVLDTVALLRAGRVRDTG A WKDEVDEKVFKRARHVVSEDRRTVCCAMALRRKDYG\|VG ESAAEMSKTIYNRCLIFIISENERVLATAKALQENNDELVG EDLTLDPDVRQRAKHVITENARVLAFQAALSARDPATAG RGREGADDVLRRRAWHVISEDARVRLAVAAAEFGDFAEFG ITRDCGLNSEELKRTLHVVTERERTKLVIQGLETNQFAQVGG EACEAVSLTSGRRLHHVLTENQRVLDAVDALKTADCMNLG LLCMVELPPPLDRRARHIVGENLRVLRGVEALRRGDARAFG AVREQVSKEDYSRAHFVLGEKERVLKVCDSLQKGDYETVG DTFNKNKPIVQNRAKHVVEENDRVIDAAKALAAHKLDEFG E LTEELKEDMYKRCLYLVKEDKRVEEAVKALSENNPEYGG QISTQUSPTVYNRCKYVIEENARVLKASALLEKNDGKGFG KCKSKMDQETYNRALHGVEEDKRTLEAAEVLDKGDFNRFG DDVKKMNK USYHRALHVVNENVRTKECKVALKMGLWDHVG A ELPELGETFARRLRHVVTEDARVEQFVEILNQEPLPRLG RVNAADDPVERRRVRHVITEDQRVGQFVEALKACGDIIKRAG PHERLDDPVELKRVRHVVTEIERVDAFVALLRAGDYDRTG RADEILDDVTFRRVRHVVTENQRVLDTVRILSEKGPNAIG E D
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G G F G G C T I N L
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GGF GGAA
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$E L V L V D S G V A E S G Y N E R R A Q V E E A S R L L G V K E L R D V N V$ EYCWVLOSKVKHSEYNERRKEVESVRNKFRSVSLKNVYN DSGVLVIDTRVRHSAYGDRRRGCEEAARLLGLPSLRE KE KMLLARSGLRHAGYNRRVTECKEAAQLGLEASGDEVYE NFEFYLIDSKKKHSGYNERRTEVESATEKI KRLHDPNLYE EFYLIDSKTKHSAYNERRKEVESAFQKIKVKTLYQVEL YK LLAFSGLKNSGYNTRVSECREAAETLGNGSLEDSFE NYKFLVAFSGLQHSGYNTRVFECKEAARALLCASGCEIYE YKVAl FFSGMKSKYNDRQDECKAAAYALENTRLRDVPE GALLVVDTREVDGQYGRRRASCEAAASALGVSTLREAVE YKULVADSGLKDAGYNSRVAECQEAARFLGKASGNDAYQ

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E
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 K I L L A F S GLK KA GYN SRVSECREAARGLLKASEPEAYE $N Y N F L V I N S N L I E S K Y N E R V Y E T T K A L E Q Q N K K N L C S D P D$ LLALLVIDDRAKHSQYGARRASCERSARVLLGVDLLADVDPD
 $A K M L L A L S G L K Q A G Y N R R V A E C K E A A Q I L L E A S G D E V Y E$ LDNLV DDSRLVDGQYADRRAACDTAAGYLGVQSLRELTR


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 E E I CK L L GU T TE E LL S A EEVCRCLGISRTRLLSP $\square A L L V D D T R L V D G Q Y A A R R D T C E A A A A V L G V E T D R E V T D$ PYSTEEICTCLGI SLEELRT AHFVVAHSGVAAAKYNRRVEEVRRAAELMGAKTVGDVAM YEDADFFSGVAGSKFNMRVDECRSAAYALKETNDRDVPV $\square S L L V I D T C L A D G Q Y G A R R A A C E A A A Q V L G V K E L G E L T D$ EEDCNCLEI SLLEELSK

E



## LVVTEKY LPDDGYTR

 EQ QAEMLGV SVDELKQ
DYQ VICNSGLIDSEYNKRREQCKKATEFFKIQALRDVTK


## LVDAHLKKGGYN HDKCK

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 ADLPQLDGQTYRRVRHI TENQRVLDTVTALRTSGAATIG ADAAAWQEVVGKRVRHIVSEDARTATALEQLQRGDLAGFG AVAHELDDMVTKRVRHVLSENARTLAAADALAADDLRMG A AEEKLPVELRPLLRHVVTEDQRVLDADALLDAGEPRALG DEWSNYDPVIRSRARHAISENERTQQAKMALTQRDLATFG AGLDAASDEVRRRVRHVVTEDERVTLAVAALESDDLEDG SAKADLDEVLYRRAKHVVTENQRTQSAARALEQNNIQKFS DQFETYDATNQKRAKHAVYENARTDEAGKKLQKADLEGQG EACGRLDDLALRARHVCGDNRRTREAARALASGDLVAMG KSCDRLDPLLKKRARHVISENDRVQKAADALEAGKADEGG NPADGLNEAETKRARHVTSERFRTERMLHNLKNGNAKEAG EQVDAMEDPLRRRARHAATEAKRSLDAIAALRAGDRSRFG ADVETLPSPLRERARHVVSENQRVQQAVQP GIGGAAEFG ADPNLLEKVERKRADHLLGEKARVQNVIRCLAQKDADGLG IDHISAVKNVTQKRIKHYVLENKRVQDAI AALEQQDLTHFG DQGDSLPAPLRPLVRHVLTENARVLATVSALRAGG D DG AHKCKLKPDLAKRADHYFSENMRVVKGI EAWEMGN DEDGG AEEKWLETVEKKRADHVIGEKKRTNDVIRNLVSGNIPK円G EDSFGLSEIEKKRAKHVIGERLRTSKAI ENLKNGNVQELG AHKSLLK PLAKRALHYFSENKRVKEGLKAWASGNLDDFG DAERLADPVLRRRARHVITENARVLQAEQLRGADPRADG AQKCILEENLARRAEHYFSEMKRVVKGRDAWARGDLREFG DQDNLIKDLIRKRAKHDISENDRVIEATSCDEENDAKKFG AYKSRLPELWKRRAEHYYSEFYRAEKGAELWRKGDLDGYG GELGQLGDLLARRGRHVVTEMERVRETLRLLESGNVRDMG MYKCKLDAALARLTADVTTE AGGQLTSAVLRRRVRHVVTEDERVRGSVAALEAGELARLG ADYTYLDKTLMRRVKHCITENARTSKAVMALEKGDMDLLG THKHILEPNLAKRAEHYFSENMRVSKGFKAWASGRFEDFG ADQERMDDVTFRRVRHDVTENRVLDTVRLLDERGVRADG NTRKFTDFKLRSRARHVFSEAHRVELFEKACEEKNGQEMG GCREQMTSEQYRRARHVISEDQRTAQAAKLLRKGDFRGLG QILSPNTQDVRQRAKHVYSEAARVLQFKKI CEDNMVQLL G RHKNKLQPNIAKRAEHYFSENDRVNKGLEFWASGNLEDLG ESMGDLPEELRRRSMHVVQERRVYDAEDALKHQDFPLLS AENNMDPVVVAARVRHVVSEDDRTAEATRQVSAGDMEAFG TETNNVDETLRKRAKYAIEEQERVKQFINNLDSNNFQECG AHLTRLSDELVKRTRHVVTEDERTRRLVALLRDDRLEEVG ANREHLDPPLDRRARHVTTE:RRVKEAAEALAADDVERVG HCADADDPALRRVLRHVLTE ERRVAAAATALRSGELAAVG AHKCKLEPDLAKRAEHYFSENMRVMKGVEAWETGSDEDFG LTEVDVEDSLRRRVRHVVTETARTAELAELVREAAAARAG LYSFPLPAEVKSRLDDVIEETERTLQAAEALRRGAYKEFG AEAGENEGVVKRRVAHVVEETARTVEAASDLEAGNLDRFR AHKGKLEATLARRAEHYFSENMRVIKGLEAWASGNLEDFG DRQGALEEVLRKRVRHVDTEDGRVPRFI EAFGRGDVDEAA ETSDYDPETVHRRVRHVATETARTIAAVDALKAGDLEQFG ELGRIEDPVVRRRARHVVTENARVLAVAETLRSGDPRELGG EMADTLPEAVRARARHIVSENRVLNAVADLRRGDVVFG NTQHVTHFRLYQRARHVYSEADRVLQFKAVCDSSPVLQLG $N T Q D V S A F K L Y Q R A K H V Y S E A A R V L Q F K K I C E D G P V A L L$
$D$
$D A S G Q D P E V T A R R V K V V S E D A R V G E V V A L D S G R T S E V G$ QILSQNTQDVYQRSKHVYSEAARVLEFQKICSEAPDQLLG EAAKKLPEVLRDRAVHVVGEAHRVLKMDGA SSLQQG EFGDRLTEKLYKRAKHVYSEAARVLQFKK CEEDNTVQLLG LEFADYPPEOVSRBHV NTQDVLLFKLYQRAKHVFSEAARVLQFKKICEENSIOLLG LHQRAVHVYEEANRVWQFKDVCEGSHLSKLG RFMSRFPVRLRQRADHVFTEALRVVQFLALLETQFNQELG RYMSRFPVRLQQRARHVFGEARRVLNFMELLSTEYNEKLG QYMSKFP DRLRQRALHVLEEAGRVLKFRDTLTTSKLLYLG RFTSRFPVRMASCARHVFSEADRVLEFMSLGDGQAREVG RYMARFPVRLRQRATHVFSEALRVLEFMELLESGSTDGGG LYQRAKHVYAEAARVLEFKDI CVRAPISLLG RFESKFAVRAERRAMHVES KYQRQLPEVIARRARHVISENYRVQAGIQALRTGNYSTFG RFTSRFPVRLRQRADHVYSEALRVLKFMSLLRQQPDRQLG SFPDRADKELYKRSKHVFTEALRVLQFRSLCSPDVYRQLGG RFTQNFSFHLHERAAHVFSEASRVYEFKNVCEDESLQKLG DEFYSGKDLKQKRAMHFFEENKRVEDAVKALEKADEAAFL QFLSMFPVLLRQRADHCFKEARRVLDFKSCLTRHNVKYGG

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A GWG C CTVSMVPTERMDSFLRTVKEQYYRAALVQQSLFVT A GWGGCTVSLVPVDALPGFLARVHEAYYQGSGGQHSLFAT GGFGGSAI ALVVRVDAEQVAGEI A A AFESEGDTAPGFLVA
 A GWGGCTVSIVPADKDSSFLANVMEAYYQRSPEKQSLFAT GGFGGCVIALVRSDQLEATAQSVLEAFVKAGFNQPRFLCA A GWGGC A V S L L V P T D K
A


 | A $G W$ | $G G G$ | $C$ | $S$ | $V$ | $H$ | $L$ | $V$ | $P$ | $A$ | $D$ | $K$ | $V$ | $D$ | $A$ | $V$ | $K$ | $E$ | $A$ | $W$ | $E$ | $K$ | $E$ | $Y$ | $Y$ | $S$ | $K$ |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| $A$ | $M$ | $D$ | $L$ | $T$ | $P$ | $E$ | $V$ | $V$ | $V$ | $S$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| $A$ | $G$ | $G$ | $C$ | $S$ | $V$ | $H$ | $L$ | $V$ | $P$ | $V$ | $D$ | $R$ | $V$ | $E$ | $A$ | $I$ | $R$ | $A$ | $A$ | $W$ | $D$ | $E$ | $H$ | $Y$ | $Y$ | $S$ |



A GWGGCSVHLVPADKVVAVRQAWEEKYYSK LKLEAAVVVS A GWGGCTVVHVPKKKV EVVFDSAMRTQYYAKRFLNAACFDT A G W G G C T V H M L P Q S K V F A V T K A L K E E Y Y L K H F L A E A MVI S A GWG GC S V H L V P A D KVV A V RE A WD R D Y Y S K K ELLEAAVVVS GGFAGTII CFVEKSGSERFDEADK PVFGE
A GWG G C TVV H M V P Q D KVVA DVVTEA L K K E Y Y K K F P L E V A MVIS
 A G W G G C T $\operatorname{C}$ V S L V

| UniRef90_UP10003F57D66 |  |
| :---: | :---: |
| UniRef90 UPI00047E9E18 UniRef90-UPI000370F4C7 | TASEGA |
| iRef9 |  |
| UniRef90-E4N | V |
| UniRet90-K11 | GD |
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| UniRef90-A3D8Y9 | SASAG |
| UniRet90-UPIOOO40C4C94 | SMGEG |
| UniRef90-M6GSH8 | R1SD |
| UniRef90-02 | V |
| , |  |
| UniRef90-D5U | KPSNGTRV |
| UniRef90-V6H1 | $\mathrm{R} \square \mathrm{S}$ D G VG |
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| UniRet90 D DLI | KPGFGSR |
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| UniRef90-S3UC |  |
| UniRet90-Q1 |  |
| UniRef90-UP10003F8: | TP |
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| UniRet90_W8AT36 |  |
| UniRef90_Q0JAB2 | D GA |
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| nireego Rikul |  |
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| UniRet90_W9GQB0 | RAGGP |
| UniRef90-M2C8 | KSAT |
| UniRef90-UP1000 | TP |
| UniRef90 V4UJV3 |  |
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| UniRef90-L7MEJ8 | EPSEG |
| niRef90-S9WHR6 | , |
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| UniR |  |
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| UniRe90-G7LJJO |  |
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| UniRef90_B8LS15 | ETGDKAC\\| |
| UniR |  |
| UniRef90-C2CQK | G D |
| UniRef90-F61603 | EAGHSARV |
| UniReef90-UP10003B74904 | VA |
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| Unip |  |
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| UniRef90-D1BL | VAGG |
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| UniRef90 W4UFU0 |  |
| UniRef90-G4MLRO | In |
| UniRet90_S3CR47 | RPGS |
| UniRet90_V8NVL3 | $N P G A G$ |
| UniRet90-K2RCE8 | KPGSGSSIV |
|  |  |
| UniRef90-GOS | V |
| UniRet90_R9NVZ2 | KPAGGACV |
| UniRef90-G3Y5Z8 | K |
| UniRet90_J9HRR9 | QPGKGACV |
| UniRef90-W9CKR7 | GGGSA |
| niRe |  |
| Ref90 |  |
| UniRet90_UP100035977F6 |  |
| Ref90_UP10004621637 | KPSSGA |
| UniRef90-T5AN90 | R P A G G S AL |



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DYRLLVVNTGGSHADYAS $\square$ PHEMRQVASA $\square$ GQKVCREVDE YA LCOVDTHGNHSDYAAVRSEMEDVAKAFGKKVLREVED EELATVLGDTVDELNA

EE $\square$ AAVLGD STDELHA LLAVLVTNSNVRHSEYPVRRRQCEEAARALGKESLRDLSE
 KS GEEAAR\RVLEEKVEALGREKVEELTG


$$
\mathrm{KD} \square \mathrm{AE} \square \mathrm{~L} \mathrm{G} \square \mathrm{~S} \square \mathrm{AE} \mathrm{~V} \mathrm{EK}
$$


$E \square M \square V F G G L A G S A Y N T R V D E C K A A A Y A D Q A Y A G M D V Y \square$ K E FRE HS I LTD S RVAREELTARGDECVEAYELVKKKNMMH LTE

LS A
 NYR V


CEE $\square \mathrm{NC} \square \mathrm{GG} \square \mathrm{DND} \quad \mathrm{S} V$

VALVDKHLASEQGYTR

E $\square \mathrm{AELLQ} \quad \square T \mathrm{~V} P$ QVEE
$A E \square E D A D R K \square G E G S \backslash R A D A V$
E EMAQLLGMTVAELEA YE TNLVDSNVKHEEYNRRRQECEEVAQKTLGKKSFREVTK VALLLIDDSARHSEYAARRASCDRAAADLRVSSGREVDD
$K \square H D G Y Y S \quad R G E L E E S L D S D F A G S P A S L L V$ NETGGYLEEIAAMVVGC TVWDLKK
 ETSEI AK QLGGSVEE K K K

TTEEAATD $\square G T$ T EQFTQ
 LTLCLTDTGGSHAGYARIPADMRYDASFFGKELLGEVDP DNVVVIMNTNKRDSKYNERRAECEKAVEELQVATGGELDE NAKIVIACSNKKDSKYNERRSECETALAELQUKTLGDLDE

TEELAKELG $\square \mathrm{L}$ VEEFTK HS CDVDTKGSHADYAADPYEMKKVANYFNQEAALREVDK DHSLCIVNTGKGSHAAYGA DPTEMKDVAHYFGKKFLREVDE YEFAVI FSGLDANAFNSRVDEAKAASFALRRDVPKEVFD EYALMIVNTGSGHAAYSADPAEMQSVARALGKEHLRDTSF


 Y SLLCITTDTKGNHSEYAAVTKEMKAVAEEFRKAVLVNVPE

LSVLVADTRVAHGGYGSRRGLVDDVARRLAAEGLGAVRD
UniRef90_GOQP35 UniRef90-M3AWG4 UniRef90-19EP62 UniRef90_U2Q1P5 UniRef90 H9GMT6 6 UniRef90_S8FWH7
UniRet90
R5Y252 UniRef90-D8QZX0 UniRef90_UPI0001FFF144 UniRef90-U1GCA9 UniRef90-V5EYN8 UniRef90-M5XSH7 UniRef90-O6ZP37
UniRef90 H1XWZ5 UniRef90-Q9SEE5 UniRef90-R7BHA7 UniRef90-G206M6 UniRef90-UP10003B3C31D UniRef90-R8BEJ7 UniRef90-F7AXEO UniRef90-H3D853 UniRef90-Q2NBZ9 UniRef90-Q4PD51 UniRef90-W1PWH7 UniRef90_UPI00035F38E1 UniRef90-J4VVA1 UniRef90-W9YE84 UniRef90-s7RIQ6 UniRef90-F4S410 UniRef90-W1TM28 UniRef90-T1F3P2
UniRef90 N1PJY2 UniRef90-N1PJY2
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UniRef90_R5YK9 UniRef90-S3L3M7 UniRef90-10YR09 UniRef90-D5U600
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YERALluly 60
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80 KYMSKFPDRDRQRAFHVFSEASRVDAFMELLNSPPLKSLG KTRD IMPARFSRRSEHFYSEYRRVRQGVTAWETGNLQLFD AAEARLDDVLFRRLRHVVTENARAVRMADDDEAGDWRAVA NTQDVTTFKLYQRAKHVYSEAARVLEFKKTCMEVPVQLLG YKRAKHVFEEALRVIEFRAALPESVLHELG DEFMDNTSLVRERAMHFYGENRRAEAEVKALEDGDFEKFK EEYTRLKAKLARRAEHFFSERARVEASVKAWVAGDLAEFG DAVERLDDRVRRRARHVVTENDRVRAVVELLSGADPREMGG SFPDRADAELYKRSKHVFTEADRVLQFQALCKQHQYSQLG

LYQRASHVYSEAKRVHAFKEAVSSNKLKHLLG FRRARHVVGEDRTAQAAAALRRGDYRAGG DTFFRADPQLRKRAFHFIRENQRVLEQVKALHDKDFDLFD LIORAAHVYSEARRVHGFKDTVNSNKLLKKLG RFTSRFPVRLDKRAFHFFGENDRVDKAVTSLENNDFESFK RFTSRFPVRLRQRAQHVYSEADRVDQFLDLLESSYNARLG QFTSRVPVRAERRAYHVFAEALRVLQFMDLLEDPYNKKGG AGKGLLSEVEFRRARHVVGEDGRARAATALGAGDYATFG NTQHVTHFKLHQRAKHVYGEAARVRQFQSVCEGDSDQTDG EAATGDDETIRRRARHCATEHRRVIAAREADDRGDMAVFG YTGEEFDKELYKRSKHVFTEALRVLQFQALCKQNPYKQLG AYKGELEANLARRAEHYFSENNRVLEGLKAWGSGNLEDFG
EEDEQLDD RRRAYYS
QY $\square S K F P \nabla R$ RQRALHVLE KD TTPSDNMVFNRMRVLTEAKRVEEFKDLITEHLLEKLG RSRHVKDNLTDSRVRHVLTEUARCQRLCELLEAGDLDQWR RSRHVKTFKLRQRAEHVYSESLRVLNFKDVCNKFTLKRLG QYKDRLPENWRKRAEHFYGERQRVKDGDKAWKKGDIEEFG VYLSWI DVELYKRAKHVLTEALRVLEFRDTCLRDILREDG KEEDDIPESVRRRVTYLIRESLSVDEAVDALKRKDNIMLS NTQDVQDFNVFKRADHVYLEALRVVQFENICSDNTLEKLG RSQMVDLFNLRQRSDHVFSEADRVLKFKDMCSKSPLLQLG EEFKYLPEPIVRRARHVISENYRVQTGVQALREENFSAFG
KHEATMPARFANRARHFYTECERVNEGVNAWREGNLTKFG EELEAGRELVSKRARHVVGEDGRAGAWRGDLRT EELEAGRELVSKRARHVVGEIDRTAQAAAALISRGDYRTFG LHQRATHVYSEAKRVYAFRDAGEEEALKRLG DSNLSVIGKLDRRLRYVDTEKMRCSEDPDIQQGNYVEFG LYKRIKHVTEEALRVLEFRDSCLSPPDTTLG DYMSTFPVRLRQRALHVFSEALRVLDFMEVLEAARAATLG QYFPVAGQRAAHVYSEAKRVYAFKDVVSSNTLKKGG QF LSSFPVQLRQRADHCFKEARRVLDFKSCLAQHNLEYLG AEADKYGFPDRKRAVHVFSEAQRVLDFRTVCEDERLTRLG TFLSAFPVQLRQRALHCFKEARRVLDFKACLSKKRLHYLG EDLERLSGTLRKRAQHVLEENERVLKSVQALKEGDFETLG A LTSVTEPVDARRARHVLTENQRVLDFVAALNGSDFGEAG LFQRATHVFSEAGRVLSFKRICDEADPQKGG DEHNKTGFKLRDRSMHVYSEAARVHQFSNECKEDPLKALG KFMSVFPVMLRQRALHVFSEASRVLEFDNLLNNQTNADLG ADVMKLGDRAVLRALHFFGDNERVVKQAAALRSHDFKKFL
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 EFKGK DPPAWRRRAEHFYGEFSRVQRGAKAWENGDDTEFG EEVYENLCRMRKRAFHFFEENERVDKEVRALKTVDFTSFG KELIDNASKLRERAYHFIEENERVINODNALKNDDUKTYI IVDDWATATAHRRIRHVVEEDHRTADVTVTLTDETAGLLGG ALGRKLGDRAVLRALHFYEENKRVIKQVDALKNNDFATFL
EQPTPDDEKTR DR पRHSMTEMVRSVQDADEDHSDSPDWTG

 a GPELGEELVEARVRHVVEETARAARATQLAAAGRAELG LYKRAKHTFEESLRVARFAELCANAGAEELG NNYLTDFPVLYQRARHVYTEALRVLKSDRLLTDDFFTSFG

Unipgoposs

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## A GFKGAC $\square$ A LVDPAMENE $\square R Q F$ VT TEY $\square K R Y P E Y K D T F E C$

A GWGGCVVA LVHEASVPAFDLS LKEGFYQKGDVKPEVFAS A GWGGCSV HLVVPADKAEAVKKA WMEKYYGKMELSEEVVVS GGGGTLTLLHRNAVEKY $\quad$ KKKVVEKYGPVF A GWGGCTVSLVKAGEVESFMKKVVEGYADDEKDKGAMFAT A GWGGCSV HEVVANKVDAVRKAWDDEYYSKRPRAAAVVAS A GWGGCAVALVKESVVPQFILALKKHYYKKNDLGGYVFAS
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A GWGGCAVSMVPTKEVMNFLVKVKKNFYEKEPSRAALFAT A GWGGCAVHLVKEEALEEFMKALREEYYVKHGFDEELFAS


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A GWGGC VVT LVKKDEVSNFMKRワRSSYYHDRGRAATMFES A GWGGSAVHLTTHDKLAK LVEALTEQYYKKQFPKAAVVVS GGFAGTVQAFVPNDIVSMYKKNMEHVFGE GACHVLVV
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A GFN GSSVVALVVDPAKKE QDEA F L KK E EY LLKVYP YK DDFAVV GGFA GVI QAFI PRSLCGAYTSYMEKAY DAGTGQKSPVFAM A G WG G C TVVS LVP E G KVES $\operatorname{F} \square A E V K Q K Y Y$
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NAVFVIDAHSQAYHADYNLRVAECRFAAQDIDAKKRDIDDQE $H$ SLLCIVDTKGSHSEYAAI RTEME SVAEFFGKKVLVREVSE DHCFLI IDDSHVRHDLFNAVRS DQENAQKKDEVGTLTVLDR

E पS LCIVDTKGSHAEYAA $\square$ PNEMKKVARFFEKEVLRDVDV
पR LVVVDTGARRDLRAARSAELERAQRTDGDRAVQDLSG HAGCIIDSGADHAEYAADPRELKEDCNFFGKDVLREVRE $\square S F L V I D T N D D G G F A A R R A A S E A C D A D G L R F D R D A D P$ YS C TDTRGSHADYAADPAEMKOAAECLGREVLGQTTE LILCLDSGADHAEYAA】PAECRAVAAVCGGEVLRDVPF
HS LCTTDTKGSHADYALTPQEMKQVADFFGKGF\REVKA YTLCITDTKGSHAEYAAIIPAEMKAVAAYFGKEVLSQVSE LDIFSNLHDGHYTK DDEVKLLDVSPTESLPP NYELLI PTGANHAEYSSDPTEMKAVAGKLGGEVLRDLS DHSLCIIDTKGSHAEYAA乌PEEMKKVAAYFGKEVLHQUDK

HS LClVDTKGSHAEYAA $\square$ PSEMKKVANYFGKEF $\square R E$ VDE
$Y K L C I V D T G G N H A D Y A S A A V E M C S V A R A L G K E V L R D$
Y
YDLIIVNTGKGHAEYSSVP DEMKKAAQVMGKEVLADVDE
NHSLIIVNTGKGHADYSLVP LEMKKVAEFFGKEVCAD：TE
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YD VIVNTGKGHAEYSSVPGEMREVAKE पGVKNDCESSV
HSLClVDVHASHADYASTPYEMKKVAQYFGKEVLRDVEE YALVITDTGGSHAEYASLPDEMKSVASQLGSNFLREVSL YALCIVDSGADHAQYAA』PRDLKAVCRLFDAEVLRQVDE
YQLCIVDTGGNHGDYAAAPAEMKSVAAQFGKQVLREVSP DYTLCCVNSNVCRSDYDADPKEMDKVANFFKKDVLRDVSY HA LC $\|$ DAKGDHAEYAA $\square$ PREMKAVAGYFTCDTLRQLSK
$V T D C V T D T Q A S H S E F A A D R R E M E A V A A C D G Q N V L G R V P E$ DYSLCIVDTKGDHDDYAMIPSEMKKVANFLGEDYLRDCDS


RLVIVNTGKGHAEYSS $\square$ PMEMREAAKAMGVELLCESSM
NTSLCTTDVHASHADYADTPYELKEVDQFFGQEFLRDVNE ALCLVDTRQDHAEYGLMPAEMRQVAAYFGAEVLRQVDP

YS $\square Y \square V N T G G S H A D Y A S V P A E M K A V A K F F G K E V L C G \square T E$ YS LCTTDTKGSHADYAAVPAEMCAVAEMLGKPVLNELSE LDFIVVDSGVRHSTMNVRQREDREALSMLKESIPRSLDE LLLCIVDTKGSHAEYAAVPYEMKQVANLLGRDVLRHVTG


RSGDQADT
DLVVLICNSDVRHSEYPTRRKQCSEADK LMGLKSYRHATE YSMCIVDTKASHADYAS $\square$ PEEMKKVAKFFGKTY LREVEE
YALCLVNVGSDHDEYAAVPTEMQEVARAFGKTRLCEVEP DEAQLLN』SRNEIAS
 YRLCIVHTRSDHAAYASIPREMRQI A GWFGAEVLRGVDP NAYLIVADTGDKG DGREAVKAVACLF DSKKATM DAFLLVADTGIKGQTRAAVRSVAE LDATQQ

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UniRef90
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50
EDTRRKLDC $\mathrm{D}^{60}$
$\stackrel{60}{8}_{8}^{\circ}$
70
80
 QCKFF LTESEF LTRSLYYQVEEQRRTIEFRKVVLADHLKADG

 Q Q F Y D NLP Q L Y QRA K H V Y S E S L R V V L K AL K M M T E D F FT D F G $L R Q R A L H V F Q E A G R V L A F R R A M E H E K L Q Q L G$
 K E F M E A L P NL R L R A I H F Y Q E N K R V M E Q K K A L L E N D F D G F L EDLSSCAGMDARA WVRHAFHDMTVVGRAADELGGAAFEAAG RFTSRFPVVRAERRAQHVFGEA LVRVVQF LAALLESGTTNQLG EVLA HLTEIRRQKRALHYVCENERVQQETEALKQDRFAEF L


E E FYK N T P EEFIGKN AN RKRA LIGFGEE K TRA HYFEENKRVDKEAKALE ENNF DAF L NTRHLTKY L L H PRVMHVVSEA QRV NTWVK LTAASVSPA E L G
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 A L N RMPDA GMARRVVYAVTEVHRLNAAVGLLDADRF GEVGG K E F M E N L N K V Y R K R A L L H V F F A E A N N R V V L T T A K K I C C S S E A S E L L L





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 STQHMQQFKLRQRALHVIQES GRVIKFRQDCEQQDIEQLG AEFYAK $\square$ GELRKRAAHFFAD DARVPQEVAALEAGDFDHFR


T A F WQA LPR L R T RA $\quad$ H Y Y QENARALA QKNA LQRGDFAEF L NEFYDRLDEDRERADHFFNE EHNVKNAVSSLENGKFDEFL A RVVPLDEEARHDCEYVLMESHLANDAASLLTTKDAS LYG ENLLAHNDR $R A V L R A M H F Y E E$
$E V G R R M S H V F S E$ DEFYARLVELRNRAMHVFEENVRVEDFVRVSMDHNLQQFG AEFYRRLREVRSRADHFFEG DARVGKAVAALNNNDLTGGE
 $N D \square Y Q N \square S A L R Q R A D H F F A D S H K A V Q E A E A L K N N N F E T F K$
 E E F Y K H I D E V E K R A T H Y F E E NK R V A R A Y E A LTV N D F K T F F
 E K FMERVA DL R L R A Y H F F K E N K R V E K L V E A L K K K D F D L F K
 A T L D S D D V TM S A H A LITTSENF EMLY GRLAA ELRR ERVRHFLD D E R E D L G D R A L L R A L L H F V RENNERVVK Q I S S Y I A S G D I K G F L

 DD WD WL S DVHRRRLEFT ILMNESTKKAIVELRKSKPDRGG



AA $\square$ G S MPEEVRRGCKHLRDDEARDAEAVQALEQADPDRFG GNLSDLKEVLVKRARHVITEDKRTQDAAEALKNHDFAKV D DF Y K L G D R P L R A L H F F E E ERRRVE GEVVKALGRGD FK E F L Y N Y $\square$ G
SAFNKLGDRCLLDRADHYWYENGLVDSRWQALQAND DTAFL E Y LT I YP VR LYQRAKHVYSEALRRVLRSVELLSNHEMNRFG TKKLICHI
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GGFAGT TETFVENSFVDEYKTRMDRVFGK
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| UniRef90-W4WQE3 | SPNQGA I |
| UniRef90-R5N3R4 | R PVGG T E I |
| UniRef90-S9VKP1 | EPSNGAKL |
| UniRef90-E1FUE5 |  |
| UniRef90-L1K342 | LP S D G A |
| UniRef90-R9LPI2 | RPEGGTLV |
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| UniRef90-UPI0003B3226E | EPADGASL |
| UniRef90_D2NQC8 | V A G |
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| UniRef90-R5JZE4 |  |
| UniRef90-D4JZ30 | R P E G G A |
| UniRef90_S4RK13 | QP VMGAA L |
| UniRef90-S0J1X9 | R P F G G $\square \mathrm{K} \mathrm{V}$ |
| UniRef90_UP10003B53E1B |  |
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| UniRef90-R6RGQ5 |  |
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| UniRef90-C1E0H6 | R P G R G A |
| UniRef90-R711R7 |  |
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| UniRef90-H6CF85 |  |
| UniRef90-C2NSD4 | H ${ }^{\text {G G S G T }}$ |
| UniRef90_10X9N2 | N! |
| UniRef90-F4LLG6 | K P A R GVHL |
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| UniRef90-R7QUC6 | T P N S G G C |
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| UniRef90-B9YC59 | R P E G G VRL |
| UniRef90_Q7QL29 | SPQRGAEI |
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| UniRef90-R5JJR8 | R P Y G G $\square \mathrm{K}$ V |
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| UniRef90 C6D3E1 |  |
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| UniRef90-A0A024FJZ9 | K E S E G H $\square$ |
| UniRef90-R5K3F2 |  |
| UniRef90-A8MCB4 | S 1 D Q G V |
| UniRef90-16XP09 | RKY G Glir ${ }^{\text {K }}$ |
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| UniRef90-A0A011QV56 |  |
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## YALCLVDVGCDHDEYAAVPIEMQKVAAAFGKTRLSEVPV

YA पMLIDDGCDHSDYAAVPDEMQKVAANFGKKR■CEVDP A LCVVDTHGNHADYAA $\square$ PGEMKAVAAHFGKAVLREVDE YA LCLVDVGSDHDEYAAVPVEMQVAAEFGKDRपCEVSP

DAY LVVGDTGKTGHT $\square$
EAVQDAVV
NHSLCCVDTKDVPAEYRADIEEMRSVAKYFGKEF RERKE RAI IE EMRSVAKYFGKEFLREVDE



 $H F \vee V A D S G I A A S E T K A V V G D L R Q R W L A D$
$E \square P L L L V N T K$
QSRSTAEEVAK
VR S L LESH D



## $\square \mathrm{VIADSG} \square \mathrm{KG}$

$L T$ L V I G D T G V R S
L I I A D T G V FV I A D S GVVS G G
$\begin{array}{llllll}E & L \\ D & \mathrm{~A} & \mathrm{~L} & L & L & V \\ \mathrm{~L} & \mathrm{D} & \mathrm{A} & \mathrm{D} & \mathrm{T} & \mathrm{G}\end{array}$
$D L F L V$ I A D S G $\square K G$ $L \vee \vee A D S G G K$
 $V F L V$ VD T G $\square R H S T H P K R Q Q E \square D$

## L V I A D S G VT G N T

ATHRPVCD
H T R E A
T R EAVE LLRRGGFERE
QSRSTMTEVAKVLKKAHPV VTE Q T GQTLKAI Q SVKELLK

K Q A V H H V S T

T A VEVAK VALLKGKHP QEAVDAVKKVHPEFVLE

VGE $\square R E L$ K E

## DFHFlVADSGGR R $\quad$ G T R S AVESVA H L L K K A



पT LVVGDTGVRS
P T R
LP VGAVRERWQ
VRKNREQNP
$E \square P L L L V N T G R Q Q R S T A V E A K D G R D R E V T E$
$L \vee A D$
Y G Q TREADQKV EALGQKALPDLQE

$T A H E V A K \square A R \square R$
T DEV
V S K V R K F K E
NFH V I A N S N $\square$ E H S
$R Q S R S T A V E V A K V Q N L R$
ELPLLLVNT
K F S S T R Q L V S N V A Q
Y G H T R E A $\square Q V$ V E S K G K E A L


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UniRef90 J2ZQT7 UniRef90－E8N5A6 UniRef90－S9V9N5
UniRef90 B8MK77 UniRef90－UPI00034643B0 UniRef90＿W8BHD5 UniRet90－B9DU64 UniRef90－W3WVO2 UniRef90－L9W6Y2 UniRef90－F3KL19
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DAKNWDEEL KKPEKYKALIFDR NEFYEHIIPMAR TENDAYLAKTALREGDYREVG

 TYPAVLEPLYHAAGRLVVEAARR
DARSALGDRAVLRALHYFEETRRVQAQQHALEAGOFAEFLD QLGELSQKR

YNRQFVAVG NVVRQMQTALAGNDAELFG
 $\square$ GLLTEQAEKALAQKDAPT G
 KYPEKYDK $\square F N E A G$ G RRTVHATLENDLARQSADDLEQGGLDSLG KVTYQTDSLSDPKQA KKELGQLTKLAKEALLNNYILELG K SSQPHL HEDSEKUG PHLDAVLGDL SQLDTDYEDTWQK\｜VHLGQDTNGVQKADQDKDDQKIGG HETGQM

TQK LGYपTKQAKQAI DENSPETLA NTQNVLIFKLYQRAKHVYSEAARVLQFKK CEDNTVQLLG SILEADDQVTQKTGQK｜QQLGYLTKQAKKAI IDNSPEKLA

 LDAGHQLGVL
 ACTSQLGEDSRETGKETTTEMRSTERALAKHGLHTVG KLVGS LDDADDKVTQ TSAELIAEDDFSTEKEDSLRRVG VHDNLLDQPEFAKPLDDQLGQUAKDARQAGQUSDEQRLG KLVNSLDDAMDRVTLSASEVIDHREDFDEKKEESLQKVG

 KAPEKIGQMLEQLGQH TEDTVIALKNNDLVKLG AE

PATYEARFDAV
AAVVTQARAADAAGDLAALG TNDPY信AGGR

LTEAVEOALTOKDNQTLG SILNADDQVTCSAQRFVGQVSDLTNEAVRDLARGRWDGFG QEKNRFNAQLSIGK LGGESR GAKEALLNNQRQELG PLIDQLGTIADS位

LQER【IDPTTNETGSDAVQS IGIVQDTKTADETKDPMKVG AVSMKRADHIVNEAIKAQKMKQICSQQKWEQIF ATVDSILNADDMVGQSAAAMDSDPEYDSENLKC VESLG ESCDLVPEEFQGYWYATUYNYHITLEAKKELLKPNPNISG NNPTWFNALLGRY NACVGEAKAAMEVGNLFRMG PKKDQSKVRRATDLTDQGRHALADGKAEELG पETDPTAHERIKHLGEDAGTKAAKQAVDDNNDEMLGG KEFSKLCE DESLLIDENVELLKENNTTELG ADPARYEALFDQV ATLVVAARTALAEGDVVT G GYERKRपEQFLIE SLVLVPAGARLDTAADPGVG EKYNADFEKD GQIVLYARKLIEHGNVVDLG AMDDADDNPKKFAQLVTKYNTVVEEAMLALQDGDLHAVG AMDDADDKVTMSASE MDKENDVTKFEADDDEDLEQFG NTRHMTQFKDRQRADHVIQESNLTKIVERKAICSUKDLITLG TNDPYLKQLLA THPKLVGPILDADDKCVHTMNDLIEDDFDKDEVESLSKVG LREEYEFAADTVESDGEIVRNGEEADAADDIEELG KNEKTFSDMCQK NAHPG AAEQLLSAMNDVTESAHKLISAFDAKDPAALLHHLG LKKNYPKDIDPDLSTDGRILTGEKMVAKKEYESUG RFNHLVSKDYQNHWYAA！HNYQLTKRAKKLLPNSNLEELA ERFGGRAGEVDAAVAALAE

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UniRef90 - x0x
Us UniRef90- $\mathrm{S3A}$ AG0 UniRef90-B5GPS7 UniRef90_UPIOOO3808FE5 UniRef90-M7TEC6 UniRef90-U2TQMO UniRef90-S3CSA8 UniRef90_K1RVT5 UniRef90-A1RYL2
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K|LIMDASHLLS CSRLYDCSCEEA DRFVSV|AVDTGGAAARMTG ALTNASGASSGMYLQVSTSGSYQPAMLALGLGGSAVRMHG ELLNQSHQLLKFDLTVSNPKLDQLVVAAQDAGACGAKLTG NLDAASGRSDQNYEGGSELIQLYEILLRAPVHGARFSG NLVNKSGDS WELLONVYSPKEOGVSLALALDSGACRVHG SLMNDSCASSHAEYESGQDETIALHALVSASTVYGSRESG PVDTEGHVSTRDDFRVSCPEDDLVVSAALAAGALGARMTG SVMNRSHLLQ ARRVSDERLDLVEVAKNQALGAKLTG TOVKRSGNSSFEYLQNVYSSKDTCMSEKTLKDKGVVRVHG DLMRASGNSSWKWLQNTVSAQDQSDPLVLADTELYCRVHG KLMTDNH GLLVALGVSHPRLDRDRELVDHNGIGWTKLTG QK MYETHHGLSKEY
RLMNVNH GFLSATG/VSTLEDEKLVYTARRAGALGSKLTG $R L V R A S G A S S A Q F L Q N V S P R G D G S G A D Q P A A V V L A C R I H G$ QLMSQAHNHLKSLGVSCDLSDLLVATALEHGALGAKMSG QLLSQNHQLLQT LGVSAEKLDYLVQTALAAGAWGAKMSG KLFDASHASMRDTYDCTHPLVDSLQELCLKSGADGSRMTG ELIRDNHGGLVSLGVSHPKLERLRELVDFAGDGWTKLTG RLVERSQRNARDNLGNQVRETDRMQALARDLGAQAASSFG RILTDCHH YLQAVGVSCPEADQLVELALASGALGAKMSG KAVTDSGLSSWRFLQNCDPKHQAMNEIYFQNHKGVCRVHG KLMTQAHR KLAALGVSCPEADQLVEAALSAGAFGAKMSG KCMDENHK QLQAI EVSCKEDDFLVDDARDQGAFGAKMTG RCMNDAHNLLDALRVSHPVVNALVRTARSNGALGAKMTG EVMNQAHQLLA RHMNQSHLS
RHMNQSHES
$H F M N E A Q K H L S$
$K L M T Q S Q R D L A$ HMMTKAHAHLQABGVSNHTLDTFCQVACDNHALGAKLTGG LLFTAAHH ELLQSLGGSHPKLDNLVNAAIHAGALGAKMSG ELLMNVNQ GLLDAGGVSUPELSSLQVNAARSCGAMGAKKVTG DLMNQSHVSCRDMYECSCPELDQLVDDCR GALGAKLTG EAMNDAQR LLKSLTDSNDFDTLIATARAAGALGAKLTG TLI RDNH GFLVSLGVSHPRLERTRELVDYADIGWTKLTG RCLNEAHR DLSALQVSSVELDHLVEVARSAGALGAKLTG HIFNACH SDLRELTVSHDKIEQLLNIGMANGAIAGKITG ESMNNAHQ ALQQLGVSNQQLDQLVSSALAHGAYGAKLTG RLMTYNHY YLNQLGVSNATIDHLVNQAWMADALGAKLTG AEMLLQHRLLSELYDVSLPELDRLVEEAVNAGAYGAKLSG DLFNQAHQLLRK LSVSSPEIENLITCALDNQALGAKLTG ELMNTNH GLLVSLGVSHPRLERVRELVDHQGIGWTKLTG RLMTQSQY NLSKLGVSTRKLDDFCRI AIQNGALGAKLTG ELMNDNH GLLVSLGVSHPRLERVRELDDHEGDGWTKLTG DNVKESGDSSWKYLQNCYEEQG CIGLAKRHKRGACRVHG ELMGMNH GLLVSLGVSHPRLERVRELVDHEGIGWTKLTG AHMTENHRLLR DLGISTEPI DAMVEAALAAGGLGAKITG TLVRVNH GFLVSLGVSHPRLERGRELVDFANDGWTKLTG TLMNHHTILRDVLKITRPKIDRMIEGALQAGAYGAKIVG Q $\mathrm{F} T \mathrm{DAN}$ S LSKLNLAVPK TDKLNKAAISAGALGSKI TG MSSSHSSLQSLYECSHPRLDVLVEAANACGALGARLTG
 RRMTENHRLLR DI GISTDRMDALVEAALAAGGLGAKLSG TLIRDNH GFLVSLGVSHPRLERIRELVDYADMGWTKLTG EI MNEAHQLLA SLTVSNAALDRLSQVARQAGALGAKLTG MLMNQSQHHLSK LGVSTDTLDDFCRIAVANNALGAKLTG SLDTANHE ELKKLGVSAPDMDALQEVALQQGALGAKLTG RVMTYQHRLLSEYYDVSLPITDELVNFLVNEGVYGAKLSG ELMT INH GLLVSLGVSHPRLERVRELVDHEGIGWTKLTG QLMNSNHDSRKKLGVSHDDIEALVSTAHENHALGAKVTG ELMTVNH GLLVALGVSHPRLERIRELVDHEGIGWTKLTG DLMNQHQLLLQNQIKNTPSEMVEMMDVAREMGALGTKIIG ELMNENHKLCQELTVSCAELDTVVNFCCENGALGAKMSG ARMTEYH GLLREAGLSTHRIDALVEAAHGAGSLGAKTTG QMLNEAQK ELVALGVSDEGLDRLINFARKEGALGAKLTG QTLD NAQDDLQTLKVSSSELNHLI QVANRAGALGAKLTG QLFDANQALLEESLGVSSPTLERLIGGARRAGALGAKLSG ELMSINH GLLVSLGVSHPRLEKVRELIDHEGVGWTKITG TLLNTNHELLR DIGVSSAELDCLTTAARSAGALGAKLAG DVLQRSQELADQGLRNQVPQTRLMVSLAREMGAVGASSFG SMLTENHCLLK EMGVSCKELDILVETADHAGAYGAKLSG RLMNECHKLCVVELTVSCDELESVVNCCRENGALGAKLSG
 QLMRQSHESLKKLYECTHVDGERLIALSDDAGVGAKLSG YSMTQAHESAKKAIGVSHAKANQLVELADKEGAGGAKMSG EMMTDNH GLLNALGVSHPRLERVRELVDHEGLGWTKLTG RLMNNG GLLSALGVSSRSLDSMVWAARNAGAHGAKL G
 VLMNINQ GLLDAI GVSSPELADLIYAARNAGAYSAKITG ELTKAH GNLKEI GVSS DEADALVETADQHGALGAKMSG TVMNAHQKLLYERIQNTPEEMHSMMQAALEAGAIGTKTTG A QLT GY H R L L R T A G L L T T DLLID A MVEA A L D A G S L G A K I T G

|  |  |
| :---: | :---: |
| UniRef90 B6KJ89 |  |
| iRef90－K9EBR | GGRGGCMIALAAQSNDASNDAQKLEKAGADETW■ |
|  |  |
| UniRef90－13T0V8 | AGFRGCCIAFVEAPLATEAASFVRREY㑑AQPEDASVVIC |
| UniRef90－F5YDN4 | G G |
| UniRef90－X0XS58 UniRef90 S3A7G0 | AGHG |
| UniRef90－B5GP | GGFGGSAIVLTRAAAAGAVAAAVRRAFAEASLTEPVVrat |
| UniRet90－UPIOOO38 | GGRGGCMI ALTNQKAVANIAQAL |
| UniRef90－M7 | GGLGGNMIALTPKELQEKVANA |
| UniRef90 R7GMZ UniRef90－U2TQMC | PGFEGTI QVFVENDYARKYKNE！ |
| UniRef90－s3 | AGGGGCALTLLKPEKLDQLAEtLeemgyekfott |
| Ree | AGGGGFTVALVTAEKEKEVVEALRQ LSPRDFSV |
| A1RY |  |
| IRef9－FUVK8 | GGFGGSVLAFV |
| UniRef90 E4 UniRef90-A9A | GGLGGGCI ALTSKDQART｜AKKLQEKGAV |
| UniRee90－Q5KLF | GGWGGSVVSLVESSQVPEFLEKVRKGYEKYGDLEDEEFAT |
| UniRef90＿U4L9L3 | A |
|  | A |
| Re990＿UP5C5 |  |
| iRef90－D4KJ | G |
|  | AGGGGCCI TL LKPGKLDKLEEQL |
| UniRet90－UP1000360BF | GGGGGCMVA TV |
| UniRef90－UPI00046328 | GGL |
| UniRef90－M2FJ23 | GGRGGCLI ALAKKESAEK |
| UniRef90－H1X |  |
| iRet90＿C5RBP2 | G GVGG |
| UniRet90－F4F | SGLGGCVI ALAKNQRAQQI STALRKA |
| UniRef90－FOVY82 | GGLGGCLIALASKSDAEKISKALKEGGA |
| UniRef90－Q0W8D2 | AGGGGCMVALCEPDRCKE：ATAIGKS |
| UniRet90＿C9C4G5 | GGRGGCMபALAQKTKAQE S S ALEDAGAAETW■Q |
| UniRef90－V5XL |  |
|  |  |
| UniRef90＿UP10003658 | GGRGGCMV边 |
| 仡 | A |
| UniRef90－cgABC8 | GGRGGCMALKKEQAEN SRLCS |
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| UniRet90＿Q82 |  |
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| UniRef90－${ }^{\text {B2B64 }}$ | AGGGGCSIMTL |
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| UniRef90－G5GH17 | GGFGGVIMAIIPEEITSQY |
| iRe990＿－7HXY8 | AGGGGC |
| Q76 | GGLGGGMIALGDPGOA |
| UniRef90－A2C | AGGGGCAI T LRPDASVDALRELERKLIAAENFSKYE |
| UniRef90－UP1000249319F | GGRGGCTISLAKNEDAKKIQQALRENGAEQTWTQPSTYE |
| UniRef90－K71W48 | AGWGGCVVAMT TNDRVEDFVENLKKKFYKGLDLMRYVFPT |
| 11CF2 |  |
| iRef90－E8Q9Q3 | GGLGGCMIALADKEMAE |
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| UniRef90－H11 | SGLGGCMI SLTDNQRDAQIISKL |
| UniRef90－UP10003B55E64 | GGMGGCMI |
| 10PS4 | AGLIGGSVIALVRDEAARELNKVLGR |
| Re990＿E9F1Y0 | AGGGGCS TLLRPDKLDELEERLD |
| C8NHR4 | GGLGGCM｜ALV ATEND |
| UniRef90－UP100006CFEBC | DGGGGSVI GlVEESKADKIIEFPMDGYYSNKDNKLYVFKS |
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| UniRef90－K4E5Y8 | TGRGGLVVALAADEAQDGI ADAVRQRCAAKFVWKYTV |
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| P1000225B6CB | AGNGGCl |
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| UniRef90－M1VUE6 |  |
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| Rei90－J2ZQT7 | AGWGGSVYALVPTDDAEDFASQWLQAYRDREQEAESTIVT |
| UniRet90－E8N5A6 6 | GGKGGNMIALASSEKLIPQTTQA |
|  | TGRGGUAVALVSSQQRDRLVEALQKNCKEAKFI |
| UniRef90－UP100034643B0 | GGLGGCVI |
| UniRef90 W8BHD5 | A GWG |
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| UniRef90－L9W6Y2 | AGGGGClVALDPTDE |
| 3KL19 | AGGGGCIISLTDESN［EKT\｜LKRERNYDCFSVK】DF |
|  | A |
| UniRef90＿E8J | AGGGGCMV A |
| UniRef90－UPIOOO2 | S |
| 90 UPIO0037D1A59 |  |

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E $\square$ S E G V R
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$L G G A G V G V$
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UniRef90-J4KM26
UniRef90- G5GHI

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KVDPGARI
EPKNGAS $\square$
$\square \mathrm{G}$ G R G $\square \mathrm{G}$ V
UniRef90 Q767H2

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| UniRef90-A0A023BXF5 | KVG P G A R |
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| $L$ |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: |
| $L$ | $P$ | $S$ | $R$ | $G$ |
| $E$ | $A$ | $C$ | $C$ |  |

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$\square G G D G \nabla G \square$
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1
20 QTS QAlI Ql VVKKK LIESPQ
 NFHLVI ANSNVEHST LPLLLVDTK QPRRTAEQLANV RNMLNAHPE
$V F V V L D S G \square K H S T H P R R Q A E \square N M G \square R L L L Q M N D R E \square R E$
SLVI AD T GVA GSTG
DAWI VI ADSGVEGSTRET V G S V A R L R A LVI A DS GI K GQTGKARRKTTDPTETDLQLGTLGRLS G LLLANTHVERDAKAVISRVHALK $M V V L D T G I K G N T \quad K E A V S D V H E$ KLVNSGFADRDRESNEGEQ\|LKKRFTMEMLE DMHLVIGYSGVKKADTTAGVKNL

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    A T MVV I A D T G|IK G
    ACLLVI A D TGGIHG
                HTREAVAKV
                                    ATKEA\I
```



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EL PLLLLVNS R
                            QPRS T A T EVA KVGGKCK A H P A V S E
```

            LVIADSGVKDEAVSRVRLKVSQNAEALQ
    DYA IVALDTGLDKSTYKVRRAQCDGFAKELFTSNFDQVRA
DAY LVICDSGDK $G Q T S Q A D A T V$

$Y R L C V \square D T G G S H A D Y A A \square P A E M F E A A R S C G Q N Q A K G L T G$

NACLLICDSGVK GKTSAAVAEVKERDNDQPVSTTQVLNN $D A V I I V A D T G K K G$ LVIADS G $\square R G Q T G P A V R T V R E R M A D F P Q E T S V$

A QLVIADSGVHG
D GVVVVADSGVAG

RTREA
VGGLRERYE
KTKDADELLRQRFER

ATMVI ADTGIKG
NASLIIADSGVHGS
EAFLVIADTGIQG TREA
YWLVVADTGLQG HTREAVNKVAQY VGGLRQRYE


| SF V $\square$ A D S G V S A S T | A VAVATVRARRE |
| :--- | ---: |
| T LL $\square G D T G V R S$ | $A T R \square P V G E V R R R W E$ |

DAVLVVADTG ESGQTRLAVAEVNAMLKDN PEYVE VKLLLTNTKVPKST AQLVGGVGKLNN DAYUVIAAESGl|ADVLQNAREYSQNADASVRKINPDFDVQQ


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DAYLLISSDSGIKG
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QTSEA!K IVK
AEKRKKKTAELVNE

LT LVIGDSGVRS PTR
NTSEA $\square$ QNVKNLGSNADSS VEKVNKFKENDE LPVGAVRERWQ KD $\square \mathrm{EE}$
EMK LVVGFTG $\square$ PH

NAF LVIADSGVHGS
TREAVSGLRQRYEND
DLWLCVGSTGVSSST

RSMVELVARLFRKPT LDATS $\square$ RDTKRLVAGVAQMK $M Q C$ KQSVEDV

DYSFVVAVSGVARELYNRAS LATTE $\quad$ VVK $\square W N Q A T G R Q S S P$

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60
GOTAYksprio
80
EYEKYUPSDLKPYFFAALKNHSITLEALQELKKPDMKYLG LCK DESHLTENVLELLKENNTRELG DTAEAHDVIDSDDFTSFPSLSRLG AHLTGQGARALTAGTLEQLG

$\qquad$ ANQPGLFDECMRRDQDVVDEAVRCFQDTDLPRESMLEGGLG LVIKNNQENMAFI ER\|GRLVHKANQAINTHNFESLA EVKGEMSEDVYKRCFHMIRETDRTLRYVDLILKADWESIG EFRDHNSESKYRKDFEKIGKDR VEEAKISLVNSDWRKTG

VRASTRHPIYENFRVKAFKALLSDEQLTALG AVKHELDTNHSTALSHRNHLGKLVNLAKDYLSNNEAMSLG
 STLEASDQVTESASDFIVSSGGFDSSSIDEAIDHFG LVEEKEKTQDSINHLGQLATDSKDFLMTDKLKEFG RDKHLSS VTYEVKAADGKKNAGLL$G$ AYASGAFQQHFKCFDYIYRAQRRFDAMMDAWRRGDVATVG KEKLQATPTATKNMTTATG A LS WQARKQLAQNDVVGLG AYANELPEPLRSKAIAHIHDRDXCREAYAMLSRGEVEHLG EMPEIIDPDLNSMGRLVGEKAREVILAEYDKEGKFKRLG ELRANHPHVTGLIQLTSSALADLENGMSDMD
$\frac{\square}{D}$
NPTVVNPTMDRDE
FMNQLAEL

 PhLVNSmFEATQSTSDEARRALGDPELSRHQLLSALS RRWQEDPARYEALFDLV ADVVHQARHALARGDVATLG ADPDGVGPLVRAPTRHPNYENFRAQAGWAALDDGDAQALG APESRMDFMGRSADLTYAAVKELTEGRARDFG DVQALIRKDPASANQHDDHLGALVKDVRGFLEHNEVGKLG NDPDNUGPRDNRLGGHGSGALAQNADTALEQADAPTLG VE $\square \quad D S A K M A N D Q W F T A \square C R T \quad F K E A H K H L R$

NKIVAEAEQTDNAGDLER口G AAEPEVVDKVLDATQSTTDEARBALLDPELSRPALLSAGG KHPA\INPVLDSDE A $\square F L K K L G E L T D E$ MVKILTENNES KEKKVDK\G T DMDRPF REDADDLKDFG DTDFVAL AAPQVVDAVIEEL SALVDPVAAALGSGDMTGLG RDP SRYEMLFDA

$$
S_{\mathrm{K}}^{\mathrm{s}}
$$ SLMNRLADLTNETRIH EHARIVE GKTKNEAGG (LVQEAYRVMTHPD FGSEETAILR RYLEAVPELYKNHWYAAI HNYDLTLKAKKADS SVDLVQVG KKLGDLTDEMAK ALTENSKS AAAADLTAGMGAKAGG ARAQLATAAGLIE VAK UENKDE】FKEDDKVIDE

KKLGKLTDEMAK TLTENDQS
$K$ HEKLRG A
 HKKGKपGDENS $\square$ NAMHA $\square K Q D$ A $\square D M K E A \square F R A D F E K T A$
 K DIHQLGDDSRETKTHDEGDADTVK
QDRFLRRAAELVAEAEQALAEGEQEEFGARLTGYH

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PEN\squareGPR\squareNH\squareGT
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| UniRef90_V7HVB1 UniRef90-A0A023BRX3 | NVD S G A RV |
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| UniRef90-D8LWR0 | E A G G A RV |
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| UniRef90-C6A3T5 |  |
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| UniRef90_Q04EX3 |  |
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| UniRef90-K5VW31 | VGG S G M G $\square$ |
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| UniRef90-C0BIK8 |  |
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| UniRef90-W7HNB5 | LGG D GVGV |
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| UniRef90-U1PD01 | R P G P S A |
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