## QUERY

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 FSEVCRVGVEGFPAYAI IVKNPPEFKSI ELRDEAIRFFA
 LFSEP E KVENMPAYALVVVKNPGEFKS EARAEAARYA
 $\square \mathrm{LF} \mathrm{S} E \mathrm{P}$ LFSEPVEVS

 Y A | $F$ | $K$ | $S$ | 1 |
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KNNAV I KKAAVNTGLEQRDYAIVMYG CAVLVKGMAAHNGNEQRDEAVYSGA GKYNYAILLKKSK I GNGMQQRDYAIILYGA GHYNCAVQVPRAAVKYGCEERDSADKSGA GKFNHAI LLKKYATKSGTEQRDFA I MYGA GQS SVAVQVKGMGDKSGMEQRDQAIMAGA GK IDIIA TIVRGAAKKRGIEQRDEAIVKVA GRFNYAVLVRGFAVRSGI EQRDAAIKMGA GVILQCKFNHAILI KNYSKMTGLEQRDYA:IVGS GKHNCAIILRKTGIGNGMQQRDYAILYGA GKYNHAVLLKNDAVKSGDEQRDAAVVMGA VHGRKGVAV QVVGVEDKS GMEQRDRAMIAGA CAVLVKGMAGRTGYEQRDDAARAGA GA DNCAVVVVRAANKVGCEERDNADKSGA SVVGDFTQGVIVYGASSKNGMQQRDAGIKAGA VI GEHSQGILVSGASEFNGDDQRDAGIRAGG $\square F D \square N \square G E S V V G S H Q Q G V V V F G V S S K N G M Q Q R D A G I K A G A$ GESSHAVMLRGCAGRSGVEQRDYAVMYGA
 N\|A TVVVRGAAAMRGIEQRDDAIKVGA EAVYVLLHGYPPAENVVK LRDEAVREGA CSVGLSVREGAR $\square R G L E E R D E A V R H G A$ EDII IKSEGYNPCFCLVRNESNKNGI EQRDAAI K IGGG SADGTYQVALLLKGKAKMKGVEQRNAALKSGG YVGDPNQCCCGIIVKQSSKDQGIRQRDDCDKSGA YCLI KNAVHKNGVSQRDAAIKIS G NYGLVVK GLAERTGLEERDEAVRHGA QVRGAAEFPAWLYRD[IAI REGA $G R R C V A H V V R G A L G G D V K I R D S L V R \square G A$ LENEK $\square G R L V L S K K N Y C U V L K N L K V K S G V E E R D D A L K F G S$ NAVAMRDELVRLGA RDEVVRRGG
 VRRLMKHNGIVARDAAIK NG EAREVLC GKRCDAYVVDEY VNDVLGLRDDLVRRGA GKRCVAHVVEGE VGDVLKTRDSLLI RLGGS
 KGY LMDSD KKYRDGA MMGA $R \square E S \square N W D S \square R \square V L K N K A K Y K \square T E L R D \| I I R E G A$


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QUERY
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