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logo

NTDB id 186787 A0031 RS06015 WP 071677086.1  MKPLVW..VLIWALGSLSLFALAGSLPDEPRFAARVDLKVSESQVRAGATLPLDVVLEALAKSVGLQPLIYRAYDASGDP   78
NTDB id 1027 TT RS05145 WP 011173436.1       .MKSAWIRAAVIALAGLGAFALAGSFPEEPRFQAPVNLKVSESQVKAGQTLPLDVVLEALARSVGLQPLIYRAYDPSGDP   79
consensus                                    * *  !**  * !!* !  !!!!!! !*!!!! !*! !!!!!!!!*!! !!!!!!!!!!!!*!!!!!!!!!!!!!*!!!!

logo

NTDB id 186787 A0031 RS06015 WP 071677086.1  AKAQPPLPNVKLDFQGKPFREVWDLLFATYGAQFNLDYLFLPPDVVVVAPTQVITALVDAPSRAGAAERRPYVVGIPEAA  158
NTDB id 1027 TT RS05145 WP 011173436.1       ATAQPPLPNVKLDFQGKPFREVWDLLFATYGNQYSLDYLFLPPDVVVVAPTQVITALVDAPSRTGAMERRPYIVGVPEIA  159
consensus                                    ! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !* !!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !! !!!!!*!!*!! !

logo

NTDB id 186787 A0031 RS06015 WP 071677086.1  YRQAIQSQGGQVQVVSNVESAKAWVQNDLLPFLSREAAGLNVNWIVEEAGRLKAILSVLATPEQHARFSDILQRAGIDF  238
NTDB id 1027 TT RS05145 WP 011173436.1       YKRTETDAQGQPRTVVNIEGAKAWVQNDLLPFLSREAAGLNVNWIVEEGGRLKAVLSVLATPEQHARFSDILQRAGIDF  239
consensus                                    !*        !!*   !  !*! !!!!!!!!!!!!!!!!!!!!
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logo

NTDB id 186787 A0031 RS06015 WP 071677086.1  TISGGNVVASILSSGLSLIFDSTRSLASLNIIATLDALQKQGLSRALRDVNQLVLNNQTARLQSGETFFIRRVVNDREER  638
NTDB id 1027 TT RS05145 WP 011173436.1       SIAGGNVAASILDSGLSLIFDSTRSLAALNIMATLDALQQQGLSRALRDVNQTVLNNQTARLQSGETFFIRRVVNDQVER  636
consensus                                    *! !!!! !!!! !!!!!!!!!!!!!! !!!*!!!!!!! !!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!  !!

logo

NTDB id 186787 A0031 RS06015 WP 071677086.1  VPFDIGIIVEATPQITADGQILLNIKAEVSGNVQRNPVNGDVDRFTKQVVTTTLRVRDGQTVVLGGLTSQETNQTQQGVP  718
NTDB id 1027 TT RS05145 WP 011173436.1       VPFDVGLIVEVTPQITADGQILLNIKAEVSGNVQRNPVDGDVDRFTKQVVTTTLRVKDGETVVLGGLTSQESNQSQQGVP  716
consensus                                    !!!!*!*!!! !!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!*!! !!!!!!!!!!!*!!*!!!!!

logo

NTDB id 186787 A0031 RS06015 WP 071677086.1  FLMDIPLIGELFKQRSQDTTEKELLVVITANIVRETANRP.  758
NTDB id 1027 TT RS05145 WP 011173436.1       LLMDIPLIGELFKQRTNESTDKELLVVITADILKEAASANP  757
consensus                                     !!!!!!!!!!!!!!****!*!!!!!!!!!! !**! !  **
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X non conserved

X similar

X $\geq$ 50% conserved