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logo

NTDB id 1094244 ACMUKF RS01315 WP 002437988.1  MKFAYIQSIRNEISIILIILLFFALIFYVFSLPFDAYVLAISIILLLMCVRWWIKYLSFKKNEHLKDKVAYLEHELAHVK   80
NTDB id 8 SA RS13860 WP 000143425.1            ..MTFLKSITQEIAIVIVIFALFGLMFYLYHLPLEAYLLALGVILLLLLIFIGIKYLSFVKTISQQQQIENLETALYQLK   78
consensus                                      **  ** !! *!! !***!   !*!*!!***!! *!!*!!* *!!!!***   !!!!!! !  *    *  !!* ! **!

logo

NTDB id 1094244 ACMUKF RS01315 WP 002437988.1  NQQIEYRNDVESYFLTWVHQIKTPITASQLLLERNEENVVNRVRQEIVHIDNYTSLALSYLKLLNEESDMTITKVTVDDL  160
NTDB id 8 SA RS13860 WP 000143425.1            NEQIEYKNDVESYFLTWVHQMKTPITAAQLLLERDEPNVVNRVRQEVIQIDNYTSLALSYLKLLNETSDISVTKISINNI  158
consensus                                      ! !!!!*!!!!!!!!!!!!!!*!!!!!! !!!!!! !*!!!!!!!!!***!!!!!!!!!!!!!!!!! !!***!!***  *

logo

NTDB id 1094244 ACMUKF RS01315 WP 002437988.1  IRPLLLKYRIQFIEQKTQIHYKKSEDIILTDAQWASIMIEQLLNNALKYAKGKDIWIDFDVANQTLQIKDNGIGISKADI  240
NTDB id 8 SA RS13860 WP 000143425.1            IRPIIMKYSIQFIDQKTKIHYEPCHHEVLTDVRWTSLMIEQLINNALKYARGKDIWIEFDEQSNQLHVKDNGIGISEADL  238
consensus                                      !!!***!! !!!!*!!! !!! **** *!!!  ! !*!!!!!*!!!!!!!*!!!!!!*!!   * !**!!!!!!!! !!*

logo

NTDB id 1094244 ACMUKF RS01315 WP 002437988.1  PKIFDKGYSGFNGRLNEQSTGIGLFIVQHIANHLNIQVTVQSELNHGTVFFIHFTKEK   298
NTDB id 8 SA RS13860 WP 000143425.1            PKIFDKGYSGYNGQRQSNSSGIGLFIVKQISTHTNHPVSVVSKQNEGTTFTIQFPDE.   295
consensus                                      !!!!!!!!!!*!!  * *!*!!!!!!! *!  ! !**!*! !  !*!! ! !*!* !*
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X non conserved

X similar

X ≥ 50% conserved