```
logo

NTDB id 46671 SLUG RS01650 WP 002478513.1  MTMFFDTIKRELIITAMALVVFLVIFFVFNLPYYAYFLGSGIMLFIMVIYWFIALSEFKKQQRLTSQLKVLEQEVQQLKSKQTQY   85
NTDB id 8 SA RS13860 WP 000143425.1        .MTFLKSITQEIAIVIVIFALFGLMFYLYHLPLEAYLLALGVILLLLLIFIGIKYLSFVKTISQQQQIENLETALYQLKNEQIEY   84
consensus                                  *  !  *!  !* !  *   *! **!****!!  !! !* !**! ***!*  !    ! !      !*  !!  * !!!  !  !

logo

NTDB id 46671 SLUG RS01650 WP 002478513.1  QNDIESYFLTWVHQIKTPITASKLLLERNEDNVVNRVRQEIIQIDNYTSLALSYLKLMNHETDMSFTKVTVNELVKPLIMKYSIQ   170
NTDB id 8 SA RS13860 WP 000143425.1        KNDVESYFLTWVHQMKTPITAAQLLLERDEPNVVNRVRQEVIQIDNYTSLALSYLKLLNETSDISVTKISINNIIRPIIMKYSIQ   169
consensus                                   !!*!!!!!!!!!!*!!!!!!  !!!!! !*!!!!!!!!!*!!!!!!!!!!!!!!!!!*!* *!*! !!***! ***!*!!!!!!!

logo

NTDB id 46671 SLUG RS01650 WP 002478513.1  FIDQHTRIHYDRSDEEVLTDVQWASIMIEQILNNALKYARGKDIWLTFDKDINVLSIKDNGVGISTADLPKIFDKGYSGYNGRLN   255
NTDB id 8 SA RS13860 WP 000143425.1        FIDQKTKIHYEPCHHEVLTDVRWTSLMIEQLINNALKYARGKDIWIEFDEQSNQLHVKDNGIGISEADLPKIFDKGYSGYNGQRQ   254
consensus                                  !!!!*!*!!!*****!!!!!! ! !*!!!!**!!!!!!!!!!!!!* !!    ! !**!!!!*!!! !!!!!!!!!!!!!!!!  *

logo

NTDB id 46671 SLUG RS01650 WP 002478513.1  EQSTGIGLFIVKHISNHIHHQVEVDSKLGQGTTFKIYFPKVDKMNEL   302
NTDB id 8 SA RS13860 WP 000143425.1        SNSSGIGLFIVKQISTHTNHPVSVVSKQNEGTTFTIQFPDE......   295
consensus                                   *!*!!!!!!!!*!! ! *!*! ! !!   !!!! ! !!  ******
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X non conserved

X similar

X ≥ 50% conserved