```
logo
NTDB id 422526 G3G87 RS01840 WP 002478513.1  MTMFFDTIKRELIITAMALVVFLVIFFVFNLPYYAYFLGSGIMLFIMVIYWFIALSEFKKQQRLTSQLKVLEQEVQQLKS   80
NTDB id 8 SA RS13860 WP 000143425.1          .MTFLKSITQEIAIVIVIFALFGLMFYLYHLPLEAYLLALGVILLLLLIFIGIKYLSFVKTISQQQQIENLETALYQLKN   79
consensus                                    *  !  *!  !* !  *  *! **!****!!  !! !* !**! ***!*  !    ! !      !*  !!  * !!!

logo

NTDB id 422526 G3G87 RS01840 WP 002478513.1  KQTQYQNDIESYFLTWVHQIKTPITASKLLLERNEDNVVNRVRQEIIQIDNYTSLALSYLKLMNHETDMSFTKVTVNELV  160
NTDB id 8 SA RS13860 WP 000143425.1          EQIEYKNDVESYFLTWVHQMKTPITAAQLLLERDEPNVVNRVRQEVIQIDNYTSLALSYLKLLNETSDISVTKISINNII  159
consensus                                     !  ! !!*!!!!!!!!!!*!!!!!!  !!!!! !*!!!!!!!!!!*!!!!!!!!!!!!!!!!!!*!* *!*! !!***! **

logo

NTDB id 422526 G3G87 RS01840 WP 002478513.1  KPLIMKYSIQFIDQHTRIHYDRSDEEVLTDVQWASIMIEQILNNALKYARGKDIWLTFDKDINVLSIKDNGVGISTADLP  240
NTDB id 8 SA RS13860 WP 000143425.1          RPIIMKYSIQFIDQKTKIHYEPCHHEVLTDVRWTSLMIEQLINNALKYARGKDIWIEFDEQSNQLHVKDNGIGISEADLP  239
consensus                                    *!*!!!!!!!!!!!!*!*!!!*****!!!!!! ! !*!!!!**!!!!!!!!!!!!!* !!   ! !**!!!!*!!! !!!!

logo

NTDB id 422526 G3G87 RS01840 WP 002478513.1  KIFDKGYSGYNGRLNEQSTGIGLFIVKHISNHIHHQVEVDSKLGQGTTFKIYFPKVDKMNEL   302
NTDB id 8 SA RS13860 WP 000143425.1          KIFDKGYSGYNGQRQSNSSGIGLFIVKQISTHTNHPVSVVSKQNEGTTFTIQFPDE......   295
consensus                                    !!!!!!!!!!!!  * *!*!!!!!!!!*!! ! *!*! ! !!   !!!! ! !!  ******
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X non conserved

X similar

X ≥ 50% conserved