```
logo
NTDB id 302516 EGCR1 RS05805 WP 205242169.1  MKNNKLVKRLLLIFLA.LIVVACIIVPIPYYIEQPGETINLKELITVNDKKDEHKGSFSLTSVGIRRATVFTALRSKMEP   79
NTDB id 360 SMU RS02495 WP 002262039.1       MKTNKKFKWWLISGISLILLLMVFFFPLPYYLEMPGGAYDIRSVVTVNHKEDKEKGSYNFVAVSLSKATPIQVLYAWLTP   80
consensus                                    !! !!   !  !*  * ***** *   !*!!!*! !!    ** **!!!*! ! *!!!*    ! * *!!*   !   * !

logo
NTDB id 302516 EGCR1 RS05805 WP 205242169.1  FNEITSADELTGGASDDEYMQIQKFNMETSQNFAIEQALKLANKPYKMEYKGVYVLGVEKNSNFAGKINVGDTVTKADGK  159
NTDB id 360 SMU RS02495 WP 002262039.1       FTEITSAKETTGGVSDADYMRINQFYMETSQNGAIYKALSLAHKKASFNYMGVYVLQVSRNSTFKGVLNIADTVTGVNDK  160
consensus                                    ! !! !! ! !!! !! *!! !* ! !!!!!! !!  !! !!*!*    ! !!!!! ! *!! ! ! *!**!!!!    !

logo
NTDB id 302516 EGCR1 RS05805 WP 205242169.1  TFTSSEDFMKYVKSQKVGQEMKVTYVHNGETKEATEKLIKLPTDKKPGIGITLTDHTEIDSQEKVKVDAGDIGGPSAGLM  239
NTDB id 360 SMU RS02495 WP 002262039.1       TFHSSKELIKYVSGLKLGSKVSVQYTSENKKKTAKGKVIKLSN.GKNGIGIGLVDHTKVTSDDKIEFSTNGIGGPSAGLM  239
consensus                                    !!*!! * *!!!   !*!  * ! ! *    ! !  !*!!!* * !*!!!! ! !!! * ! *!*      !!!!!!!!!

logo
NTDB id 302516 EGCR1 RS05805 WP 205242169.1  FTLEIYEQLIDKDLRHGEKIAGTGTINANGDVGRIGGIDKKVASAAADKNKVFFAPDDEITKEMKEAQPGIKSNYQEAKA  319
NTDB id 360 SMU RS02495 WP 002262039.1       FTLDIYDQLVKEDLRKGRIIAGTGTIGEDGSVGDIGGADMKVAAADRINADIFFVPNNPVSKATLKKNPKALTNYQEAKQ  319
consensus                                    !!!*!!*!!*  !!!*!  !!!!!!!   ! !! !!! ! !!! !      *!! !  ***!     *!   *!!!!!!

logo
NTDB id 302516 EGCR1 RS05805 WP 205242169.1  AAKKIDTEMKIVPVKTVKDALDYLETMK   347
NTDB id 360 SMU RS02495 WP 002262039.1       AAKKLGTKMKIVPVKNVQEAIDYLRKH.   346
consensus                                    !!!!* ! !!!!!!! ! *!*!!!  **
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X non conserved

X similar

X $\geq$ 50% conserved