```
logo
NTDB id 27584 MEMAR RS09875 WP 011844836.1  MGFKDLFNDFLNRRPSDQERIPE...DTPDADRFVEQDQEIFGRIENWQKYQEGFYRFLKHPFRVMMEEPYTVLFVSVPVALLLF   82
NTDB id 63 F7686 RS01500 WP 066956433.1     MGFKEIIDDFLNRLPGQDRTTGGAAADALVSESFEEREEEIFGRIENWRKYQEGFYRFLKHPLRVMAEEPLTVLFISIPAALLLF   85
consensus                                   !!!!**  !!!!! !      * ***! *  * ! ! * !!!!!!!! !!!!!!!!!!!!! !!! !!! !!!!*!*! !!!!!

logo
NTDB id 27584 MEMAR RS09875 WP 011844836.1  IGGFVSMVTSYGIGALFTTTMIDDVFVFALLIAIVPLGLLDFKEAWRVSSVEASLPNFFRDVAGMNDSGMTLPHAIHIVSEGEYG  167
NTDB id 63 F7686 RS01500 WP 066956433.1     AGGFMSLVLSYGVGVLFSTTMIDDVFVFSLLIAIVPLAILDLKESLRVSSIEASLPNFFRDVAGMNDSGMTLPHAIHIVSEGEYG  170
consensus                                    !!!*!*! !!!*! !!*!!!!!!!!!! !!!!!!!!**!! !!  !!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 27584 MEMAR RS09875 WP 011844836.1  ALTPHIRKLDTEMSWGVPFVEAIRRFGKTVNTPLAERSVDLIAHASSAGGDVSEVLRAAAHDAYEFFNLKTERKNGMMIYMIIIL  252
NTDB id 63 F7686 RS01500 WP 066956433.1     ALTPHIRKLDTEMSWGVPFVEAIRRFGKAVNTPLTERSVDLIAHASSAGGDVSEVLRAAAHDAYEFFNLKMERRNGMMIYMIIIV  255
consensus                                   !!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*

logo
NTDB id 27584 MEMAR RS09875 WP 011844836.1  MSFFVFLFVIGVLSGTFLTTMAEAGEAVSASGSSSAQSFMGNVDLFLYNRIFCHSALIQGFFSGLAAGIMGEGRVVGGLKYSALM  337
NTDB id 63 F7686 RS01500 WP 066956433.1     MSFFVFLFVIGVLSSTFLTTMAEAGTAVAASGS..TQTFMGTVDLFLYNRLFCHAALIQGFFSGLGAGVMGEGRVLAGLKYSAIM  338
consensus                                   !!!!!!!!!!!!!! !!!!!!!!!!! !! !!!!** !*!!! !!!!!!!!*!!! !!!!!!!!!!*!!*!!!!!!**!!!!!!*!

logo
NTDB id 27584 MEMAR RS09875 WP 011844836.1  VMIAWIAFRFII   349
NTDB id 63 F7686 RS01500 WP 066956433.1     VLIAWIAFRFFI   350
consensus                                   !*!!!!!!!!! !
```

X non conserved
X similar
X ≥ 50% conserved