logo
NTDB id 1426 RS RS05575 WP 011001070.1 MRALLIGFVAGCMALQTRPALAPVWQPALLGLACLLAMALLRRGARLPRLVPVMAVLAAACAGFGWSDWRAQQRLSVVLA 80
NTDB id 221596 B5U27 RS19240 WP 005743843.1 MRTGMIALASGLVALRFLPALPPTWLLLLMPILALMLLPLRT.....YPLALFLLGFTWACVSAQWALNDRLV 68
NTDB id 1199 PSJM300 12650 AFN78592.1 MRAGMLALAAGLLLLRFLPQLPSFGWLLALAPVGLLLLPFRW.....RPLAFFLFGLVWACLNAQWALDDRLP 68
consensus
logo
NTDB id 1426 RS RS05575 WP 011001070.1 PAWEGKDIRATGVVAELPEVTEDATRFLFRIESSDAGDAVPPRVRLSWYGLRSWHERQREAEAATDRRAGIPDLQPGQRW 160
NTDB id 221596 B5U27 RS19240 WP 005743843.1 HRLDGQTLWLQGKVVGLPSVAEGVVRFELEGARS.RRAILPARIRVAWYGG.....PQVNSGERW 127
NTDB id 1199 PSJM300 12650 AFN78592.1 VDLDGRTFWLEGQVTGLPDRRGDVVRFELEDIHS.RHAGLPSRIRLAWYGG.....PEIRSGERW 127
consensus
logo
NTDB id 1426 RS RS05575 WP 011001070.1 QLTLRLKRPHSLMNPGGFDGEYAMLADDVRAAGYVYTGKRARNALMGEADAGSGVGLRVERWRAAVRRHLLAALPEARYA 240
NTDB id 221596 B5U27 RS19240 WP 005743843.1 RMAVKLKRPAGLVNPDAFDYEAWLLAQRIGATGTVVDGQLLT..P.....ARGAWRDLIRQRLLAVDAQG.RE 192
NTDB id 1199 PSJM300 12650 AFN78592.1 RLAARLKRPSGMVNPSAFDYEAWLLARRIGATGTIKAGERIAEAA.....SSGAWRDRLRQRLLTVEAHG.RA 194
consensus
logo
NTDB id 1426 RS RS05575 WP 011001070.1 PVIVALVVGDQSGIAQADWERFRRTGISHLVSISGLHITMIAGLFGALWMALWRRSFGLARWLRTPLPLRMPAQRAGAVA 320
NTDB id 221596 B5U27 RS19240 WP 005743843.1 GGLAALVLGDGSGLSSTDWQVLQDTGTVHLLVISGQHIGLLAGVIYALV.....AGLARWGLWP..RALPWLPCACAL 263
NTDB id 1199 PSJM300 12650 AFN78592.1 GAIAALVVGDGSGLSTADWRVLQDTGTVHLMVISGQHVGMLAGLLYAVV.....LLLARWGIWP..QRLPWLPWACGL 265
consensus
logo
NTDB id 1426 RS RS05575 WP 011001070.1 AMAAAFAYCLLAGMGVPAQRTLLMLATVAVARLADRSVPVSLSLCWAAALVALVDPWAVMSAGGWLSFGAVAVIFLAARI 400
NTDB id 221596 B5U27 RS19240 WP 005743843.1 AFIAALGYGLLAGFEVPVRRACVMVAMVLLWRLRFRHLGVVWPLLLSFNAVLIFEPLVTLQPGFWLSFAAVGILVLIFSG 343
NTDB id 1199 PSJM300 12650 AFN78592.1 AFAGALGYGWLAGFEVPVRRACLMVAIVLLWRLRFRHLGVWLPLLIALVGVLMVDPLASLQPGFWLSFVAVALLIWIFRG 345
consensus
logo
NTDB id 1426 RS RS05575 WP 011001070.1 VAADARDDVRRPAWPVRAWRGLRGAARVQLAVTFGLLPLTLLLFQQVSVVSAVANALAIPVVSFVTTPLALMGTA.LPEP 479
NTDB id 221596 B5U27 RS19240 WP 005743843.1 ....RLG.......AWNWWQSWTRAQWLIAIGLLPILLALNLPISLSGPFANLLAVPWVSLIVLPPALLGTLLLPVP 409
NTDB id 1199 PSJM300 12650 AFN78592.1 ....RLG.......VPSWWRALLYAQWAMALGLLPAMLVLGLPVSLSGPLANLLAVPWISVLVVPLALLGSFLLWVP 411
consensus

X non conserved
X similar
X ≥ 50% conserved