logo
NTDB id 1426 RS RS05575 WP 011001070.1 MRALLIGFVAGCMALQTRPALAPVWQPALLGLACLLAMALLRRGARLPRLVPVMAVLAAACAGFGWSDWRAQQRLSVVLA 80
NTDB id 373724 FFH90 RS08535 WP 138866992.1 MRTGMMALALGLLALRFLPALPPAGLWLLLPVVGLMLLP.....F...R....TFPLAFWLFGFSWACANAQWALDDRLA 68
NTDB id 1199 PSJM300 12650 AFN78592.1 MRAGMLALAAGLLLLRFLPQLPSFGWLLALAPVGLLLLP.....F...R....WRPLAFFLFGLVWACLNAQWALDDRLP 68
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NTDB id 1426 RS RS05575 WP 011001070.1 PAWEGKDIRATGVVAELPEVTEDATRFLFRIESSDAGDAVPPRVRLSWYGLRSWHERQREAEAATDRRAGIPDLQPGQRW 160
NTDB id 373724 FFH90 RS08535 WP 138866992.1 LKLDGETRWVEGRVTGLPQQAEGVVRFELTDS.QSRRTQLPRHLRLAWYGG.....PPVNSGERW 127
NTDB id 1199 PSJM300 12650 AFN78592.1 VDLDGRTFWLEGQVTGLPDRRGDVVRFELEDI.HSRHAGLPSRIRLAWYGG.....PEIRSGERW 127
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NTDB id 1426 RS RS05575 WP 011001070.1 QLTLRLKRPHSLMNPGGFDGEYAMLADDVRAAGYVYTGKRARNALMGEADAGSGVGLRVERWRAAVRRHLLAALPEARYA 240
NTDB id 373724 FFH90 RS08535 WP 138866992.1 RLAVKLKRPAGLLNPHGFDYDAWLLAQRIGATGTVKDGQRLGQA.....QWAWRDGIRQRLQAVDAH.GRT 192
NTDB id 1199 PSJM300 12650 AFN78592.1 RLAARLKRPSGMVNPSAFDYEAWLLARRIGATGTIKAGERIAEAA.....SSGAWRDRLRQRLLTVEAH.GRA 194
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NTDB id 373724 FFH90 RS08535 WP 138866992.1 GALTALVLGDGAGLSREDWQVLQDTGTVHLLVISGQHIGLLAGLVYLLIAG.....LARYGLW..PRRLPWLPWACGL 263
NTDB id 1199 PSJM300 12650 AFN78592.1 GAIAALVVGDGSGLSTADWRVLQDTGTVHLMVISGQHVGMLAGLLYAVVLL.....LARWGIW..PQRLPWLPWACGL 265
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NTDB id 1426 RS RS05575 WP 011001070.1 AMAAAFAYCLLAGMGVPAQRTLLMLATVAVARLADRSVPVSLSLCWAAALVALVDPWAVMSAGGWLSFGAVAVIFLAARI 400
NTDB id 373724 FFH90 RS08535 WP 138866992.1 AFAAALGYGLLAGFEVPVRRACLMIGLLLLWRLRFRHLGAWWPLLLALDGVLLLDPLASLQPGFWLSFAAVAVLIFTFGG 343
NTDB id 1199 PSJM300 12650 AFN78592.1 AFAGALGYGWLAGFEVPVRRACLMVAIVLLWRLRFRHLGVWLPLLIALVGVLMVDPLASLQPGFWLSFVAVALLIWIFRG 345
consensus
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NTDB id 1426 RS RS05575 WP 011001070.1 VAADARDDVRRPAWPVRAWRGLRGAARVQLAVTFGLLPLTLLLFQQVSVVSAVANALAIPVVSFVTTPLALMGTA.LPEP 479
NTDB id 373724 FFH90 RS08535 WP 138866992.1 ....RL.......GPWRWWQTWTRAQWLIAIGLCPLLLVLGLPISLSGPLANLLAVPWISLLVLPPALLGTLLLPVP 409
NTDB id 1199 PSJM300 12650 AFN78592.1 ....RL.......GVPSWWRALLYAQWAMALGLLPAMLVLGLPVSLSGPLANLLAVPWISVLVVPLALLGSFLLWVP 411
consensus

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NTDB id 1426 RS RS05575 WP 011001070.1 WSQPVLIWAEASFAWLMPWLDALALPGGSVWVAPAAPAWAWGLAVVGVPLLLVPGGYRAWAWRAQGAVLLLPMLLARAPA 559
NTDB id 373724 FFH90 RS08535 WP 138866992.1 YVGEGLLWLAGGLIDLLFKGLALVAGQLPAWVPVAVPLWVWAFGALGAFLLLLPPGVP...LRPLGWPMLLLLVFAPRHT 486
NTDB id 1199 PSJM300 12650 AFN78592.1 WLGEALLWLAGGLLAVLFELLALIAGWQPAWLASALPLWAWSLVALGTLVLLLPAGVP...LRVLGTLLLAPLLFAPDSR 488
consensus
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NTDB id 1426 RS RS05575 WP 011001070.1 PPPGEFRMVAFDIGQGAATLVETAGHRLLFDTGPRQGDLADAADRVIVPYLRGHGVQAIDTLVVSHEDSDHAGGTETVMA 639
NTDB id 373724 FFH90 RS08535 WP 138866992.1 LPEGVAEIWQLDVGQGLAILVRTRHHTLLYDAGPRFGDI.DLGERVVLPTLRKLGVNGLDLMLISHADADHAGGARAIAS 565
NTDB id 1199 PSJM300 12650 AFN78592.1 PEQGRAEVWVFDVGQGLSVLVRTRDHALLYDAGPRYGDF.DIGERVVFPSLRQLGLERLDLMMLSHADSDHAGGALAIQR 567
consensus
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NTDB id 1426 RS RS05575 WP 011001070.1 GVPVRTMLASLPPGHGLERQARALGIEPAGCVAGQRWAWDGVAFEILYPVQVPSDRAAVSSNARSCVLRIANARYAAMLA 719
NTDB id 373724 FFH90 RS08535 WP 138866992.1 GVPVTRVLSGDPLAL.....PDELHAEGCESGQQWTWDDVRFQLWQWS.....SASDSNQKSCVLQIEANGERLLLT 632
NTDB id 1199 PSJM300 12650 AFN78592.1 AMPVGAVLSGEPERL.....PAALDARSCRTGQRWTWNQVNFSVWRWP.....QATNGNQASCVLMVEAAGERLLLT 634
consensus
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NTDB id 1426 RS RS05575 WP 011001070.1 GDIGRAQEAALIAAEAPERLMADILLVPHHGSRTSSSGAFLDAVSAQVAVFQVGYRNRYGHPHPQVWQRYGARGIERLRT 799
NTDB id 373724 FFH90 RS08535 WP 138866992.1 GDIDTAAERVLLD..SPLAVPTDWLQAPHHGSRSSSSMALLSALLPKAVLISRGNGNSFGHPHPTVVARYQKRGMAIYDS 710
NTDB id 1199 PSJM300 12650 AFN78592.1 GDIDAQAERALVD..GGMEVNARWLLAPHHGSNSSSSAVFLAAVAADGALISRSSHNAFGHPHPAVLKRLQNAGAELHDT 712
consensus
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NTDB id 1426 RS RS05575 WP 011001070.1 DATGAVVIETRGDALAVRTARSMRPRYWSSALEVAALAETTEPTGAQP 847
NTDB id 373724 FFH90 RS08535 WP 138866992.1 AEQGAIRLQLGRF..KPPWTMLQERRFWRDPPALRQ..... 744
NTDB id 1199 PSJM300 12650 AFN78592.1 AEHGALSLQLGAF..GGARRMRDEPRFWREK..... 741
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X non conserved
X similar
X ≥ 50% conserved