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NTDB id 1426 RS RS05575 WP 011001070.1	MRALLTGFVAGCMATQTRPALAPVWQPAALGACLAMALLRRGARLPRLVPVMAVLAAACAGFCWSDWRAQQRLSVVLAPAWEG	85
NTDB id 34592 A458 RS35730 WP 080583069.1	MRTWMLALAAGLVLPRLPALPSFWITCVLGVLGG...LLSRR.....RWASSGLLLLGFAWACWQCQSAIDDLRPVSQDG	74
NTDB id 1199 PSJM300 12650 AFN78592.1	MRAGMLALAAGLLLRFLPQLPSFGWLLALAPVGL...LLLPF.....RWRPLAFFLFGLVWACLNAQWALDDRLPVDLDG	73
consensus	!!* ****!!* ****!!* **** * *!* **** *!* ****	

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NTDB id 1426 RS RS05575 WP 011001070.1	RTYWTLYTIGLPVTVTGDWREFLEDLSRHHAGLPSSRVRLSNVYGG	
NTDB id 34592 A458 RS35730 WP 080583069.1	KDFRAETQAEESDRRELATILFRTEESDAGDAVPRVRLSNVYGG	170
NTDB id 1199 PSJM300 12650 AFN78592.1	KDIRATGVVAELPEVTE DATRFLFRIESSDAGDAVPRVRLSNVYGLRSWHERQREAEAAATDRRAGIPDIQPGQRWQLTLRLKRPH	138
consensus	RTYWIQGTIVSGLPTS TGDVVRFELEDIE.SRHAGLPSKVRLSNVYGG.....PQVRAGERWRLLAARLKRP	137

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NTDB id 1426 RS RS05575 WP 011001070.1	GLVNPDAFDYEAWLLARRIGATGYKAGERLRNA	
NTDB id 34592 A458 RS35730 WP 080583069.1	SLMNPGEFDGEYAMLADDVRAAGYVYTGRARNALMGEADAGSGVGLRVERWRAAVRRHLLAALPEARYAPVIVALVVGDDQSGIA	255
NTDB id 1199 PSJM300 12650 AFN78592.1	GLVNPDAFDYEAWLLARRIGATGSVKAGERIFAP.....HSVNSWRDRLRLRMEVSASGRQGAVALVVGDDSGLS	210
consensus	GMVNPSAFDYEAWLLARRIGATGTKAGERIAEA.....ASSGAWRDRLRQLLTVEAHG.RAGATAALVVGDDSGLS	209

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NTDB id 1426 RS RS05575 WP 011001070.1	TADWRVQLQDTGTVHLMVSGQHGLMAGLLYALVLLARNGFWPSRW	
NTDB id 34592 A458 RS35730 WP 080583069.1	QADWERFRRTGISHTVSTGLHTMTAGIFGALWATWRSEFGLARWLRTPLPLRMPAQRACAVAAAMAAFAAYCLLAGMGVPAQR	340
NTDB id 1199 PSJM300 12650 AFN78592.1	REDWQVLQDTGTVHLMVISGQHIGMLAGLLYGLVATLAAWGFWPSRW.....PWLPACALALAGALSYGWLAGFDVPVRR	286
consensus	TADWRVQLQDTGTVHLMVISGQHVGMLAGLLYAVVLLARWGTPQRL.....PWLPWACGLAFAGALGYGWLAGEVVPVRR	285

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NTDB id 1426 RS RS05575 WP 011001070.1	ACLMVAAYVLLWRIRFRHLGVWLPPLLWALVGVLLVDPLASLQPGFVLSFGAVAVLAFR	
NTDB id 34592 A458 RS35730 WP 080583069.1	TLMLATVAVARLADRSVPVSLCWAALVALVDPWAVMSAGGWLSFGAVAVIFLAARIVAADARDDVRRPAWPVRAWRGLRGA	425
NTDB id 1199 PSJM300 12650 AFN78592.1	ACLMVAAYVLLWRIRFRHLGVWLPPLLALVGVLLVDPLASLQPGFWLSFGAVAVLALAFS.....GRLGRWKW.....WATL	357
consensus	ACLMVAAYVLLWRIRFRHLGVWLPPLLALVGVLLVDPLASLQPGFWLSFVAVALLIWIIFR.....GRLGVPSW.....WRAL	356

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NTDB id 1426 RS RS05575 WP 011001070.1	TRAQWAVTIGLLPLTMTYGLPVSYSGLPLANLLAVPWVSYYTVPALLGTLLPYPWSEALLVLYGTSALLFELLALAAWQPA	
NTDB id 34592 A458 RS35730 WP 080583069.1	ARVQLAVTIFGLLPITLLFQQVSVSAVANALAPVVSFVTTPLALMGTA.LPEPWSQPVLIVAEASFALMPWLDALALPGGSV	509
NTDB id 1199 PSJM300 12650 AFN78592.1	TRAQWAAVSLGLLPITLLGLPVSASGPLANLLAVPWVS.LTVPFALLGTLLLPGGIGEALLWLVTISLALLFELLASAMATWQPA	441
consensus	LYAQWAAVSLGLLPITLLGLPVSLSGPLANLLAVPWVS.LVLPVPLALLGSFLLWVLPWGEALLWLAGLLAVLFELLALIAWQPA	441

