

logo

NTDB id 976742 MGAS38345 RS00695 WP 010921802.1	MTFEKIEEAYQLLLENCQLIENDLKTHIYDAIVEQNSFYLGAEGASPPVAQNSDKLKALCLTKEEWRKAYQFLFIKAAQT	80
NTDB id 401 FSA28 RS09370 WP 002291646.1	MDFEKIEEAYNLLLDNCQRLETQLHTHLYDALIEQNACYLGAEGADDDFIKENNKKLRQLNLSKEEWRRSFQFIFIKATQS	80
NTDB id 392 SMU RS09000 WP 002263442.1	MDFEKIEEAYNLLLDNCQRLETQLHTHLYDALIEQNACYLGAEGADDDFIKENNKKLRQLNLSKEEWRRSFQFIFIKATQS	80
consensus	!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!	

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NTDB id 976742 MGAS38345 RS00695 WP 010921802.1	EQQLQSNHQFTPDsIGFILLYLLEGLTNKSQLDLEIGSGTGNLAETIYNNSKSLDYMGIEVDDLLIDLSASIAEIMDSS	160
NTDB id 401 FSA28 RS09370 WP 002291646.1	QQQLQSNHQFTPDSSIGFIILYLLEGLTNKSQLDIIIEIGSGTGNLAETIVNNSKSIDYIGIEVDDLLIDLSASIAADVLDSS	160
NTDB id 392 SMU RS09000 WP 002263442.1	QQQLQSNHQFTPDSSIGFIILYLLEGLTNKSQLDIIIEIGSGTGNLAETIVNNSKSIDYMGIEVDDLLIDLSASIAADVLDSS	160
consensus	*!!!!!!!!*!!!!!!!!*!!!!!!!!*!!!!!!!!*!!!!*!*!*!*!*!!!!!!!!*!*!*!*!*!*!!!!*!!!!*!!!!!!!!*!!!!*	

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NTDB id 976742 MGAS38345 RS00695 WP 010921802.1	AHFVQEDAVRPQILKESDIVISDLPVGYYPNDDTAKRYKVASSDKHITYAHLLMEQSLKYLKKDGF AIFLAPVNLLTSPQ	240
NTDB id 401 FSA28 RS09370 WP 002291646.1	VHFVQEDAVRPQILKESDVIISDLPVGFYPNDKIAQRYQVAAKLGHTYAHLLMEQSLKYLKKEGLAIFLAPVDLLTSEQ	240
NTDB id 392 SMU RS09000 WP 002263442.1	VHFVQEDAVRPQILKESDIIISDLPVGFYPNDKIAQRYQVAAKLGHTYAHLLMEQSLKYLKKEGLAIFLAPVDLLTSEQ	240
consensus	*!*!!!!!!!!*!!!!!!!!*!!!!!!!!*!!!!!!!!*!!!!*!*!*!*!*!!!!!!!!*!*!*!*!*!*!!!!*!!!!*!!!!!!!!*!!!!*	

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NTDB id 976742 MGAS38345 RS00695 WP 010921802.1	SPLLKKWLQNNVTILAVISLPEAIFKQKSNVKSIFVLKKQSQQSLESFVYPLTDLQDPEVLRFRFMKNFKKWKEDNVF	317
NTDB id 401 FSA28 RS09370 WP 002291646.1	SPLLKKWLQNNVTILAVISLPEAIFKQKSNVKSIFVLKKQSQQSLESFVYPLTDLQDPEVLRFRFMKNFKKWKEDNVF	317
NTDB id 392 SMU RS09000 WP 002263442.1	SPLLKKWLQNNVTILAVISLPEAIFKQKSNVKSIFVLKKQSQQSLESFVYPLTDLQDPEVLRFRFMKNFKKWKEDNVF	317
consensus	!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!	

-  non conserved
-  similar
-  ≥ 50% conserved