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NTDB id 1306 DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQDPDPAALR	80
NTDB id 1355 HI 0298 AAC21962.1	0
NTDB id 1366 NTHI RS01950 WP 011271972.1	0
NTDB id 1118696 ELZ64 RS01995 WP 126374476.1	0
NTDB id 1384 A4U84 RS08525 WP 010787006.1	0
NTDB id 1168 A1552VC RS11070 WP 000957200.1	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	0
NTDB id 1293 VP RS12240 WP 005479695.1	0
NTDB id 1251 GCO85 RS07725 WP 011946523.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
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NTDB id 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPPFALEGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIIVVPSGASDA	160
NTDB id 1355 HI 0298 AAC21962.1	0
NTDB id 1366 NTHI RS01950 WP 011271972.1	0
NTDB id 1118696 ELZ64 RS01995 WP 126374476.1	0
NTDB id 1384 A4U84 RS08525 WP 010787006.1	0
NTDB id 1168 A1552VC RS11070 WP 000957200.1	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	0
NTDB id 1293 VP RS12240 WP 005479695.1	0
NTDB id 1251 GCO85 RS07725 WP 011946523.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
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NTDB id 1306 DR RS10060 WP 027479822.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITDQLYEVLAEQEGTTFLPNPSGFHPGEEVLGSM	240
NTDB id 1355 HI 0298 AAC21962.1	0
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NTDB id 1118696 ELZ64 RS01995 WP 126374476.1	0
NTDB id 1384 A4U84 RS08525 WP 010787006.1	0
NTDB id 1168 A1552VC RS11070 WP 000957200.1	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	0
NTDB id 1293 VP RS12240 WP 005479695.1	0
NTDB id 1251 GCO85 RS07725 WP 011946523.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
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NTDB id 1306 DR RS10060 WP 027479822.1	LRADALRLLAVPVDETEQGVTVLTSDFRKRDPIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGLGEQLVQEGTLSRDQL	320
NTDB id 1355 HI 0298 AAC21962.1	0
NTDB id 1366 NTHI RS01950 WP 011271972.1	0
NTDB id 1118696 ELZ64 RS01995 WP 126374476.1	0
NTDB id 1384 A4U84 RS08525 WP 010787006.1	0
NTDB id 1168 A1552VC RS11070 WP 000957200.1	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1	0
NTDB id 1293 VP RS12240 WP 005479695.1	0
NTDB id 1251 GCO85 RS07725 WP 011946523.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
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NTDB id 1306 DR RS10060 WP 027479822.1	REALQVQAREGKVKPLGEVTELGFASPDEVDSAQKQNVGGGRLEDTLVQSGKSPEMARSAAQICYEFLDPIQNPP	400
NTDB id 1355 HI 0298 AAC21962.1	24
NTDB id 1366 NTHI RS01950 WP 011271972.1	24
NTDB id 1118696 ELZ64 RS01995 WP 126374476.1	18
NTDB id 1384 A4U84 RS08525 WP 010787006.1	19
NTDB id 1168 A1552VC RS11070 WP 000957200.1	69
NTDB id 1401 DSB67 RS12665 WP 010643259.1	69
NTDB id 1293 VP RS12240 WP 005479695.1	69
NTDB id 1251 GCO85 RS07725 WP 011946523.1	...MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEKISLLQYLVKNKIISAEQIALTAQAQNFVPMPLDINCI	77
NTDB id 1058 ABD1 RS01615 WP 001274986.1	...MSALHTSPKFTGFFRRLVEEKHVSAAATMQTALDAAKRAKQDVTAVYLIEEVHLSPLLAETISAEFAEPYFDLDVYDT	77
NTDB id 1018 ACIAD RS01685 WP 004920473.1	...MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQAIEKAKKFKQDVPYLDNFSISPLTIAEISLEFGEPLLDIGVFDIP	77
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NTDB id 1306 DR RS10060 WP 027479822.1	DPKVALMPEATARRYVVVRLQGNLIVVANKDPRNVFALDDLKLIITGKEILPAVMAEKDIIIRLIERYEGEKGFEKLNK	480
NTDB id 1355 HI 0298 AAC21962.1	DLWERNQQQQSLILRYFALPLKEENRRLWLGVDLSNLSACETIAFITGKPVPEPILLESSQKELLQQLTPNQMQVEEQV	104
NTDB id 1366 NTHI RS01950 WP 011271972.1	DLWERNQQQQSLILRYFALPLKEENRRLWLGVDLSNLSACETIAFITGKPVPEPILLESSQKELLQQLTPHQMQVEEQV	104
NTDB id 1118696 ELZ64 RS01995 WP 126374476.1	EQWQKNYDERHILRLRYLTVPLREDEGTLWLAIDDMNLSACEIFAFITHKRTEPVLVSSEELKYLLGQLMPEQQSVYDET	98
NTDB id 1384 A4U84 RS08525 WP 010787006.1	SRWQKNCEEKQLILRYLAVPVQETEHKLLWAVDDENNLTAACEIFAFMTHKQIEPVVIAADELKYLLNALSPEQQPIYEE	99
NTDB id 1168 A1552VC RS11070 WP 000957200.1	ANLCQQIGLRELIITRYDALPIAKQGNLILAVSDPTLLQAEDEFRFATGLQVELALADHRAAQAAIRRLYGRSIQGAANQ	149
NTDB id 1401 DSB67 RS12665 WP 010643259.1	ASLCQTLGLRDLITRHNALPLNRTSSTLILAVADPTNLQAEDDFRFATGLQVELVLADEFRELTAAIRRLYGRSLGQEKSG	149
NTDB id 1293 VP RS12240 WP 005479695.1	ASLCQQIGLRELIITRHNALPLHRTPTLILAVADPTNLQAEDDFRFATGLQVELVLADEFRELTAAIRRLYGRSLSHEKSG	149
NTDB id 1251 GCO85 RS07725 WP 011946523.1	GTIPVNLVNEKLIKRRHAMVPLFSRGTNLYLATDDPSKQASLKEIQFHTGLNTHAVVETDKLSALIDNLTAKESQGLSE	157
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SQIPKDLVDQKLLIKHRILPLIQRGQILYVATSNPSNIEAIDAIRFNSKLLVEPVIVEHHKLEKVLGQHFAEESSFDFND	157
NTDB id 1018 ACIAD RS01685 WP 004920473.1	ALFLKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTNIEAMDAIRFNSKLLVEPVIVEHDKLERLLSEHVEETHFNFD	157
consensus		

