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NTDB id 1306 DR RS10060 WP 027479822.1	MALSI GDRRLGAILLDQGYLGDN DLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
NTDB id 1384 A4U84 RS08525 WP 010787006.1		0
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NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPPFALEGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	160
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consensus		

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consensus		

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NTDB id 1306	DR RS10060 WP 027479822.1	LRADALRLLAVPVDETEQGVTVLTSDPRKRPDI DAL I GRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGTLSRDQL	320
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consensus			

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NTDB id 1306	DR RS10060 WP 027479822.1	REALQVQAREGKVKP LGEVITELGFASPDDEVDSALQKQNVGGGRLEDTLVQSGKLSPEMLARSLAAQLGYEFLDPIQNPP	400
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NTDB id 1293	VP RS12240 WP 005479695.1	MHSNLSTILRQKGLLTFSQEESEIEQVKASGISMPEALLSSGFFTSSELAEHLSSIFGLSQPELSQYEE	69
NTDB id 1401	DSB67 RS12665 WP 010643259.1	MLTNLSTVLRQAGILTFSQEESEIEQVKASGISMPEALLSSGFFQSHLTELHSSVFGLSCTSLNQYEE	69
NTDB id 1110	NGFG RS09235 WP 003689817.1	MSVGLLRILVQNQVVTVEQAEHYY.NESQAGKEVLPMLFSDGVISPKSLAAL IARVFSYSILDLRHYPR	68
NTDB id 1251	GC085 RS07725 WP 011946523.1	...MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVKNKILSAEQIALTAQNFGVPMMLDINCIDV	77
NTDB id 1018	ACIAD RS01685 WP 004920473.1	...MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQAIEKAKKFKQDIVPYLIDNFSISPLTIAEITISLEFGEPLLDLGVFDP	77
NTDB id 1058	ABD1 RS01615 WP 001274986.1	...MSALHTSPKFTGFFRRLVEEKHVSAAATMQTALDAAKRAKQD TVAYLIEEVHLSPSLLAETISAFAEFPYFDLDVYDT	77
consensus		* ** ** * ** **** ** * *** ** **	

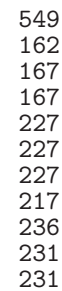
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NTDB id 1058	ABD1 RS01615 WP 001274986.1	SQTPKDLVDQKLILKHRILPLIQRGQILYVATSNPSNIEAIDAIRFNSKLLVEPVIVEHHKLEKVLGQHFAEESSFDND	157
consensus		* ** ** ** **!* * ***** ***** * ***** ***** ***** * ** **	

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consensus

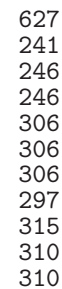
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consensus

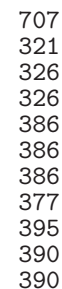
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
consensus

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-  non conserved
-  similar
-  >50% conserved