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		MSXLSIGDRLGALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVK	
NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVK	80
NTDB id 1306	DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLDIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	79
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
NTDB id 1110	NGFG RS09235 WP 003689817.1	.....	0
NTDB id 1145490	BT141 RS07800 WP 011946523.1	.....	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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		RSTPRAKALRLRAIPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLEIVVPSGA	
NTDB id 1028	TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLEIVVPSGA	158
NTDB id 1306	DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	157
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
NTDB id 1110	NGFG RS09235 WP 003689817.1	.....	0
NTDB id 1145490	BT141 RS07800 WP 011946523.1	.....	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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		SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGFLESTEGIVPDPSSAA	
NTDB id 1028	TT RS08215 WP 011173991.1	SDAGRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
NTDB id 1306	DR RS10060 WP 027479822.1	.....	237
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
NTDB id 1110	NGFG RS09235 WP 003689817.1	.....	0
NTDB id 1145490	BT141 RS07800 WP 011946523.1	.....	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 1028 TT RS08215 WP 011173991.1	LSMLRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSR	317
NTDB id 1306 DR RS10060 WP 027479822.1	LLLLRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSR	317
NTDB id 1168 A1552VC RS11070 WP 000957200.1	.....	0
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NTDB id 1110 NGFG RS09235 WP 003689817.1	.....	0
NTDB id 1145490 BT141 RS07800 WP 011946523.1	.....	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 1028 TT RS08215 WP 011173991.1	MSYDTEERLITGLRYIVQECILTAIQEELLEAKASGISLPEYLYSSGILSPEEIAEILTSFGYPLDLSG	397
NTDB id 1306 DR RS10060 WP 027479822.1	5SLREALGAPAELEKYNLFTERREAKYRSEKQAEADRYGAEKLDYDQITLITSTLTAQAEELAESLQETAE	397
NTDB id 1168 A1552VC RS11070 WP 000957200.1	.....MLTNLVAILRQABLISATTEQAVVTQVSASCTSVPEALLESIFHAQELTEQLSHIFGLPETDLR	66
NTDB id 1293 VP RS12240 WP 005479695.1	.....MHSNLSLTLRQKGLLTFSGEESLIEQVKASGISMPEALSSCFFTSSELAPHLSSIFGLSQPELSQ	66
NTDB id 1401 DSB67 RS12665 WP 010643259.1	.....MLTNLSTVLRQAGILTFSGEESLIEQVKASGISMPEALSSCFFQSHLTELSSVFGLSCTSLNQ	66
NTDB id 1110 NGFG RS09235 WP 003689817.1	.....MSVGLLRLLVQNVVTVVEQAEHYY.NESQACKEVLPMLFSDQVTSPKSLAALARVFSYSILDRH	65
NTDB id 1145490 BT141 RS07800 WP 011946523.1	.....MALATEEYRLQGTGQLLVLEKLLDKTKATEIHKLAAAEEKISLLQYIVKNKILSAEQIALTAQNFVGPMLDINC	74
NTDB id 1251 GCO85 RS07725 WP 011946523.1	.....MALATEEYRLQGTGQLLVLEKLLDKTKATEIHKLAAAEEKISLLQYIVKNKILSAEQIALTAQNFVGPMLDINC	74
NTDB id 1058 ABD1 RS01615 WP 001274986.1	.....MSALHTSPKFTGFFRRLVVEEKHVSAAATMQTALDAAKRAKQDTVAYDIEEVHLSPSLLETSAEFAEPYFDLDV	74
NTDB id 1018 ACIAD RS01685 WP 004920473.1	.....MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQALEKAKKFKQDLPYLLIDNFSISPLTIAEISLEFCEPLLDLIGV	74
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NTDB id 1028 TT RS08215 WP 011173991.1	YDYALPQSLYLEKLLRHAYDLEERRNLYLAYSDDTNGALDDPRRFATG	477
NTDB id 1306 DR RS10060 WP 027479822.1	LEVDLQSLYRREKRYSLYSELESLYLLMANRSLSTRAYSLEKRRKLNPTLRAATADEKATKRALESLYK	472
NTDB id 1168 A1552VC RS11070 WP 000957200.1	.....DPPDPGAPLLIPEDLCRRYGVFPHRLEGNRLVLLMKDPRNLLALDDVRLALKRKGNYEVAPAVATEAAITKLERFYGA	141
NTDB id 1293 VP RS12240 WP 005479695.1	.....NPPDPKVALMIPEATARRVYVVPVRLQGNLSLVAMKDPRNVFALDDDLKLTG.....KEILPAVMAEKDIRLERFYGE	141
NTDB id 1401 DSB67 RS12665 WP 010643259.1	YDYANLCQQLGLRELITRYDALPIAKQCNMLLLAVSDPTLLQAEDEFRFATG.....LQVELALADHRALQAIRRLYGR	141
NTDB id 1110 NGFG RS09235 WP 003689817.1	YDYANLCQQLGLRELITRHNAIPLHRTPSTLLAVADPTNQAEEDDFRFATG.....LQVELVLADEFRELTATIRRLYGR	141
NTDB id 1145490 BT141 RS07800 WP 011946523.1	YDYANLCQQLGLRELITRHNAIPLHRTPSTLLAVADPTNQAEEDDFRFATG.....LQVELVLADEFRELTATIRRLYGR	141
NTDB id 1251 GCO85 RS07725 WP 011946523.1	YPRHRVLMGVLTEEQMVEFHCVVFRRGDKVFFAVSDPTQMPQIQKTVSAAG.....LAVELVIVEDDQLACLLDWGSR	140
NTDB id 1058 ABD1 RS01615 WP 001274986.1	IDVGTIPVNLVNEKLIKRRHAMVPLFSRGTNLYLATDDPSKQASLKEIQFHTG.....LNTHAIVVETDKLSALIDNLLTA	149
NTDB id 1018 ACIAD RS01685 WP 004920473.1	IDVGTIPVNLVNEKLIKRRHAMVPLFSRGTNLYLATDDPSKQASLKEIQFHTG.....LNTHAIVVETDKLSALIDNLLTA	149
consensus		



