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NTDB id 1306 DR RS10060 WP 027479822.1	MSXLIIGDRRLGAALLDQGYLGNDLQRALERHSEVGGRLAEVLYDSQMYERRIABTIEBFGIPLVQLHAEVETSPKXKRRTBR	84
NTDB id 1028 TT RS08215 WP 011173991.1	.MALSIGDRRLGAILLDQGYLGNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALRSIRP	85
NTDB id 1384 A4U84 RS08525 WP 010787006.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSRLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKVALLPA	0
NTDB id 1355 HI 0298 AAC21962.1		0
NTDB id 1366 NTHI RS01950 WP 011271972.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		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 1306 DR RS10060 WP 027479822.1	ETAKELRAIPFALBEEKRYRVAHYBPLBTFSEYEYEDLSGFYEPYQITREEMVAIAHYPELGLIEMYPSCASBEGSTSLKL	167
NTDB id 1028 TT RS08215 WP 011173991.1	QTALNLQAFPPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLIIVVPSGASDAGRTGGKL	168
NTDB id 1384 A4U84 RS08525 WP 010787006.1	EKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..PSGEGQKDLKL	0
NTDB id 1355 HI 0298 AAC21962.1		0
NTDB id 1366 NTHI RS01950 WP 011271972.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	CEELIYHGYIIBERLEYALYKQERTGSLCRTLYSBRATIEEALYEXLAIEQEGTEFLESTEGAMPBEEEXLMLRLDALRYLAVP	252
NTDB id 1028 TT RS08215 WP 011173991.1	GERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVLGSMRLRADALRLLAVP	252
NTDB id 1384 A4U84 RS08525 WP 010787006.1	GELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLLLRSDALRYGAVP	0
NTDB id 1355 HI 0298 AAC21962.1		0
NTDB id 1366 NTHI RS01950 WP 011271972.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	YGECPSEVYVTSDPRKRRERYRLLNRPXBLITPBRWEELABEYVQRNRLGEXLVQEGLSRBRLEALVQQLMKNLIG	337
NTDB id 1028 TT RS08215 WP 011173991.1	VDETEQGVTVLTS DPRKRDPIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRGLGEQLVQEGTSLRDQLREALQVQAREGKVKPLG	337
NTDB id 1384 A4U84 RS08525 WP 010787006.1	IGFQNGEVEVVLSDPRHKEAVAQLNRPARFYLALPQAWHEELFRRAYPKQNRLLGEVLVQEGKLSREALKEALEVQKGLPRAKPLG	0
NTDB id 1355 HI 0298 AAC21962.1		0
NTDB id 1366 NTHI RS01950 WP 011271972.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		6
NTDB id 1251 GC085 RS07725 WP 011946523.1	MALATEEYRLQGTLG	14
NTDB id 1293 VP RS12240 WP 005479695.1	MHSNLS	6
NTDB id 1401 DSB67 RS12665 WP 010643259.1	MLTNLS	6
consensus		*



