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NTDB id 1306 DR RS10060 WP 027479822.1	GSMLRADALRLLAVPVDETEQQVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPQGRGLGELVQEGTLSSR	317
NTDB id 1028 TT RS08215 WP 011173991.1	LLLLRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYLALPQAWHEELFRRAYPQKNRLLGEVLVQEGKLSR	317
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 1121090 EL143 RS04870 WP 085416671.1	0
NTDB id 1110 NGFG RS09235 WP 003689817.1	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	DQLREALIQQVQAREGKVKPLGEVITELGFASPDVDSALQQKQNVGCGRIEDTLVQSCKLSPEMLARSLLAAQLGYEFLDPIQ	397
NTDB id 1028 TT RS08215 WP 011173991.1	EALKEALEVQKGLPRAKPLGETLVLEGLARPEDVEALQQKQRGCGRIEDTLVQSCKLRPEALAGAVATQIGYPYVDPEE	397
NTDB id 1168 A1552VC RS11070 WP 000957200.1	66
NTDB id 1293 VP RS12240 WP 005479695.1	66
NTDB id 1401 DSB67 RS12665 WP 010643259.1	66
NTDB id 1121090 EL143 RS04870 WP 085416671.1	65
NTDB id 1110 NGFG RS09235 WP 003689817.1	65
NTDB id 1251 GCO85 RS07725 WP 011946523.1	74
NTDB id 1058 ABD1 RS01615 WP 001274986.1	74
NTDB id 1018 ACIAD RS01685 WP 004920473.1	74
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NTDB id 1306 DR RS10060 WP 027479822.1	NPPDPKVALMIP EATARRYVVV PVR LQGN SLV VAMK DPRNV FALDD LK LITG K E L P A V M A E K D I R L I E R Y F G E	472
NTDB id 1028 TT RS08215 WP 011173991.1	DPPDPGAPLLLPEDLCRRYGVFP HRLEGNRLVLLMKDPRNILALDDVRLALKRKGLNVEVAPAVATEAATKLERFYVK	477
NTDB id 1168 A1552VC RS11070 WP 000957200.1	YDYANLCQQLGLRELITRYDALPTAKQCNLLLLAVSDPTLLQAE EEFRFATG LQVELALADHRA LQAATRRLYGR	141
NTDB id 1293 VP RS12240 WP 005479695.1	YEYASLCQQLGLRELITRHNA LPLHRTPSTLLLAVADPTNQQAE DDFRFATG LQVELVLADFRELSTAIRRLYGR	141
NTDB id 1401 DSB67 RS12665 WP 010643259.1	YEYASLCQQLGLRDLITRHNA LPLNRSTSLILAVADPTNLQAE DDFRFATG LQVELVLADFRELTAATRRLYGR	141
NTDB id 1121090 EL143 RS04870 WP 085416671.1	YPRN NVLTDV LSEHAMVQRCLP IFRGNKVFLAVADPTQIQTYQKLAFTAG VTFDLVVVRS DQLDSLLEWQNR	140
NTDB id 1110 NGFG RS09235 WP 003689817.1	YPRHRVLMGVLT E EQMVEHCV PFVRGDKVFFAVSDPTQMPQIQKT VSAAG IAVELVVE D DQLAGLLDWVGSR	140
NTDB id 1251 GCO85 RS07725 WP 011946523.1	IDVGTIPVNLVNEKLIKRRHAMVPLFSRGTNLYLATDDPSKQASIKETQFHTG LNTHAVVETDKLSALT DNLLTA	149
NTDB id 1058 ABD1 RS01615 WP 001274986.1	YDTSQIPKDLVDQK LILKHRILPLIQRGQILYVATSNPSNIEA DAI RFNSK LLVEPVIVEHHKLEKVLGQHFAE	149
NTDB id 1018 ACIAD RS01685 WP 004920473.1	FDPALFLKDKID EKL IQKYRIMPLVHRGHVLYVATSNPTNIEA DAI RFNSK LKVEPIIVEHDKLERLISEHFVE	149
consensus		

