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NTDB id 1313 DR RS12030 WP 010888966.1	IYGPESGGKTTLALAIIVAAQAQKAGGTC	AFIDAEHALDPVYARALGVNTDELLVSQPDN	GEQALEIMELLVRS	GAVD	IVVV	155				
NTDB id 230830 CDIF104451 RS06915 WP 003428218.1	VYGPESGGKTTVALSCV	ASAKKDGIIAAAFIDAEHALDPVYAKALGV	DVNDLTIISQPD	TGEQALEIAEAL	IRSGAIDITVT	144				
NTDB id 600 KW2 RS01765 WP 011675413.1	IYGPESGGKTTVALHAVA	AAVQKEGGIAAFIDAEHALDPEYAKALGVN	IDELLLSQPDYGEQ	QLIAEKLIT	SGAVDLVVI	157				
NTDB id 269 KZH43 RS08665 WP 001085462.1	IYGPESGGKTTVALHAVA	AQAQKEGGIAAFIDAEHALDPAYAAALGVN	IDELLLSQPDSGEQ	GLEIAGKLI	DSGAVDLVVV	156				
NTDB id 228 SPD RS09265 WP 001085462.1	IYGPESGGKTTVALHAVA	AQAQKEGGIAAFIDAEHALDPAYAAALGVN	IDELLLSQPDSGEQ	GLEIAGKLI	DSGAVDLVVV	156				
NTDB id 194 SPR RS08825 WP 001085462.1	IYGPESGGKTTVALHAVA	AQAQKEGGIAAFIDAEHALDPAYAAALGVN	IDELLLSQPDSGEQ	GLEIAGKLI	DSGAVDLVVV	156				
NTDB id 159 SP RS09750 WP 001085462.1	IYGPESGGKTTVALHAVA	AQAQKEGGIAAFIDAEHALDPAYAAALGVN	IDELLLSQPDSGEQ	GLEIAGKLI	DSGAVDLVVV	156				
NTDB id 527 SMSK321 RS11065 WP 001085454.1	IYGPESGGKTTVALHAVA	AQAQKEGGIAAFIDAEHALDPAYAAALGVN	IDELLLSQPDSGEQ	GLEIAGKLI	DSGAVDLVVV	156				
NTDB id 499 SM12261 RS08315 WP 078228268.1	IYGPESGGKTTVALHAVA	AQAQKEGGIAAFIDAEHALDPAYAAALGVN	IDELLLSQPDSGEQ	GLEIAGKLI	DSGAVDLVVV	156				
NTDB id 408 SMU RS09530 WP 002262392.1	IYGPESGGKTTVALHAVA	AQAQKDGIIAAAFIDAEHALDPAYAAALGVN	IDELLLSQPDSGEQ	GLEIAGKLI	DSGAVDLVVV	156				
NTDB id 453 Spy49 1753c ACI62003.1	IYGPESGGKTTVALHAVA	AQAQKEGGIAAFIDAEHALDPAYAAALGVN	IDELLLSQPDSGEQ	GLEIAGKLI	DSGAVDLVVV	156				
NTDB id 115 BSU 16940 NP 389576.2	VYGPESGGKTTVALHAI	AEVQQQGGQAAAFIDAEHALDPVYAKLGVN	TELLLSQPD	TGEQALEIAEAL	VRS	GAVD	IVVV	141		
NTDB id 625 LCA RS02525 WP 011374196.1	IYGPESGGKTTVALHAVA	AEVQKQGGTAA	YIDAENAMDPKYAT	ALGVN	IDDLLLSQPD	TGEQ	GLEIADAL	VSSGAVD	ILVV	143
NTDB id 1245 Cj1673c YP 002345041.1	IYGPESGGKTTLTLHI	IAECCQKAGGVCAFIDAEHALDVHYAKNLGVN	TDDLIVSQPDF	GEQALEIVET	IARSGAVD	LI	VV	142		
NTDB id 1211 C694 RS00795 WP 000963128.1	IYGPESGGKTTLSLHI	IAECCQKNGGVC	AFIDAEHALDVHYAKRLGVN	TENLLVSQPD	TGEQALEILET	ITRSG	GID	LVVV	144	
NTDB id 1214 NCTC11637 00359 SQJ03206.1	IYGPESGGKTTLSLHI	IAECCQKNGGVC	AFIDAEHALDVHYAKRLGVN	TENLLVSQPD	TGEQALEILET	ITRSG	GID	LVVV	144	
NTDB id 1334 RAOC RS04870 WP 004918236.1	IYGPESGGKTTLTLHAI	IAEAQKVGGTAAAFIDAEHAFDMHYAEKLG	INLEELIVSQPD	NGEQALEIAD	SLIRSG	GVD	IVVV	151		
NTDB id 1389 A4U84 RS07560 WP 021112712.1	IYGPESGGKTTLTLSLI	AQAQKGNKTC	AFIDAEHALDPVYARNLGVN	TDELLISQPD	NGEQALEICD	ALVRS	GAVD	VITV	159	
NTDB id 1130 NGFG RS03960 WP 003688695.1	IFGPESGGKTTLCLEAVA	QCCQKNGGVC	AFVDAEHAFDPVYARKLGVK	VEELYLSQPD	TGEQALEICDTL	VRS	GGID	MVVV	143	
NTDB id 1124 NGFG RS03960 WP 003688695.1	IFGPESGGKTTLCLEAVA	QCCQKNGGVC	AFVDAEHAFDPVYARKLGVK	VEELYLSQPD	TGEQALEICDTL	VRS	GGID	MVVV	143	
NTDB id 1123 OK783 RS03845 WP 003688695.1	IFGPESGGKTTLCLEAVA	QCCQKNGGVC	AFVDAEHAFDPVYARKLGVK	VEELYLSQPD	TGEQALEICDTL	VRS	GGID	MVVV	143	
NTDB id 1427 RS RS02760 WP 011000510.1	IYGPESGGKTTLTLQV	VAEMQKLGCTCAFIDAEHALDVTYADKLG	VKVPDLLISQPD	TGEQALEIAD	ALVRS	GS	VD	LIV	I	151
NTDB id 1165 A1552VC RS01565 WP 000344154.1	IFGPESGGKTTLLELI	IAAAQREGKTC	AFIDAEHALDPVYAKKLGVN	IDELLVSQPD	TGEQALEICD	ALAR	SGAVD	VIVV	142	
NTDB id 1142 GTF74 RS11550 WP 000344154.1	IFGPESGGKTTLLELI	IAAAQREGKTC	AFIDAEHALDPVYAKKLGVN	IDELLVSQPD	TGEQALEICD	ALAR	SGAVD	VIVV	142	
NTDB id 1022 ACIAD RS06390 WP 004925639.1	IYGPESGGKTTMTLQAI	AQCCQKAGGTC	AFIDAEHALDPYARKLGVN	IDNLLVSQPD	HGEQALEIAD	M	VRS	GAT	DLIVV	142
NTDB id 1074 ABD1 RS10195 WP 000344167.1	IYGPESGGKTTMTLQAI	AQCCQKSGGTC	AFIDAEHALDPYARKLGVN	IDNLLVSQPD	NGEQALEIAD	M	VRS	GAT	DLIVV	142
NTDB id 1205 PSJM300 05545 AFN77183.1	IYGPESGGKTTLTLSVI	IAAAQRMGSTC	AFVDAEHALDPDYAGKLGVN	VDDLVSQPD	TGEQALEITD	M	VRS	NAV	VDVIV	142
consensus	**!!!!*!!!!* ! **!	!* **	!*****!***!!	!!	!!*****!***!!!*!!	*	***!***!***			

