



logo

NTDB id 1306 DR RS10060 WP 027479822.1	REALQVQARECKVKPLGCEVITELGCFASPD	400
NTDB id 1168 A1552VC RS11070 WP 000957200.1	REALQVQAREGKVKPLGEVITELGFASPD	69
NTDB id 1293 VP RS12240 WP 005479695.1	EQAVVTQVSASGTSVPEALLSIFHAQELTEQLSHIFG	69
NTDB id 1401 DSB67 RS12665 WP 010643259.1	EESLIEQVKASGISMPEALLSSGFFTSSELAEHLSSIF	69
NTDB id 1110 NGFG RS09235 WP 003689817.1	EESLIEQVKASGISMPEALLSSGFFQSHLTHSSVFC	68
NTDB id 1251 GC085 RS07725 WP 011946523.1	AEHYYNESQA.CKEVLPLMFLSDGVI.SPKSLAAL	77
NTDB id 1101946 Q9279 RS07730 WP 011408072.1	AIELHKLAAAEKISLLQYIVKNKILSAEQIALTAA	79
NTDB id 1058 ABD1 RS01615 WP 001274986.1	ARTAMEQAAA.AKVPLPQWFAENRLVSASQLAAAN	77
NTDB id 1018 ACIAD RS01685 WP 004920473.1	MQTALDAAKRAKQDVTAYLIEEVHLSPSLAEET	77
consensus	MQALEKAKKFQKD.VPYLIDNFS.SPLTTAEILS	

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NTDB id 1306 DR RS10060 WP 027479822.1	DPKVALMIPATARRVYVVPVRLQGNLSLVAMKD	480
NTDB id 1168 A1552VC RS11070 WP 000957200.1	ANLCQQQLGLRELITRYDALPIAKQCNLLLLAVS	149
NTDB id 1293 VP RS12240 WP 005479695.1	ASLCQQQLGLRELITRHNALPLHRTPTSTLLLAV	149
NTDB id 1401 DSB67 RS12665 WP 010643259.1	ASLCQTLGLRDLITRHNALPLNRTSSTLLI	149
NTDB id 1110 NGFG RS09235 WP 003689817.1	HRVLMGVLTETHEQMVETHCVPVFRGDKVFF	157
NTDB id 1251 GC085 RS07725 WP 011946523.1	CTIPVNLVNEKLIKRRHAMVPLFSRGTNLYLAT	159
NTDB id 1101946 Q9279 RS07730 WP 011408072.1	NQNAVKLVSLELLQKYHVLPLFKRCNRLFVGV	157
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SQIPKDLVDQKLLKHLRLPLIQRGQILYVATS	157
NTDB id 1018 ACIAD RS01685 WP 004920473.1	ALFLKDKIDEKLLIQYRIMPLVHRGHVLYVAT	157
consensus	DPKVALMIPATARRVYVVPVRLQGNLSLVAMKD	

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NTDB id 1306 DR RS10060 WP 027479822.1	ELA.....ERNKTQSSQEA.DLSVADESALVQV	547
NTDB id 1168 A1552VC RS11070 WP 000957200.1	GK.EISQDELANLVK...VSDDELQSI.DLSQD	225
NTDB id 1293 VP RS12240 WP 005479695.1	LKEINQEELASLVD...VGADENIDEDLSQD	225
NTDB id 1401 DSB67 RS12665 WP 010643259.1	LKEINQDELAGLVD...IGDEIENIEDLSQD	225
NTDB id 1110 NGFG RS09235 WP 003689817.1	GEGQ.E.....EESHTLYIDNEEAEDGPVPR	215
NTDB id 1251 GC085 RS07725 WP 011946523.1	YVEDSGDLEGLETSADD...EDQDIDTATSVT	234
NTDB id 1101946 Q9279 RS07730 WP 011408072.1	GNDD.EEMGDLDVSTGDEDMGAGG.DSGVD	238
NTDB id 1058 ABD1 RS01615 WP 001274986.1	EE.FDLDVN..LDGPTAQED.EEAPQG	229
NTDB id 1018 ACIAD RS01685 WP 004920473.1	EE.FDLDVN..LDGPTAQED.EEAPQG	229
consensus	ELA.....ERNKTQSSQEA.DLSVADESALVQV	

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NTDB id 1306 DR RS10060 WP 027479822.1	FPKGAAQQIMARLKIMGHLDIAERRVPQDGRV	625
NTDB id 1168 A1552VC RS11070 WP 000957200.1	PASHLSRRRLAARIKILSKLDIAERRVPQDGR	304
NTDB id 1293 VP RS12240 WP 005479695.1	PPNHLSRRLSARIKILSKLDIAERRVPQDGR	304
NTDB id 1401 DSB67 RS12665 WP 010643259.1	PPSHLSRRLSARIKILSKLDIAERRVPQDGR	304
NTDB id 1110 NGFG RS09235 WP 003689817.1	PPIAVRGQLASRIKIVMSRLDISEKRIPQDGR	295
NTDB id 1251 GC085 RS07725 WP 011946523.1	PPASLSSRITARIKIVMSNLDISERRIPQDGR	313
NTDB id 1101946 Q9279 RS07730 WP 011408072.1	APVKLSQRILARIKIVMSQLDIAEKRVQDGR	317
NTDB id 1058 ABD1 RS01615 WP 001274986.1	PPLQLANRLASRIKIVMSQMDISEKRVQDGR	308
NTDB id 1018 ACIAD RS01685 WP 004920473.1	PPLQLATRLASRIKIVMSQMDISEKRVQDGR	308
consensus	FPKGAAQQIMARLKIMGHLDIAERRVPQDGRV	

 non conserved
 similar
 $\geq 50\%$ conserved