

NTDB id	Accession	Protein	Sequence	Score
NTDB id 1306	DR RS10060 WP 027479822.1	MALSIIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80	
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 1251	GC085 RS07725 WP 011946523.1		0	
NTDB id 404616	IX0842 RS07650 WP 011408072.1		0	
NTDB id 1058	ABD1 RS01615 WP 001274986.1		0	
NTDB id 1018	ACIAD RS01685 WP 004920473.1		0	
consensus				

NTDB id	Accession	Sequence	Count
NTDB id 1306	DR RS10060 WP 027479822.1	SIRPQTALNLQAFPPFALEGGDRLRVALVDPLSSFSIETLEDDSGGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	160
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 1251	GC085 RS07725 WP 011946523.1		0
NTDB id 404616	IX0842 RS07650 WP 011408072.1		0
NTDB id 1058	ABD1 RS01615 WP 001274986.1		0
NTDB id 1018	ACIAD RS01685 WP 004920473.1		0
consensus			

NTDB id	Accession	Sequence	Count
1306	DR RS10060 WP 027479822.1	GRTGGKLG ^{ER} LITHGYITDAQLQVALDAQQQTGEALGATLISQRAIT ^{ED} QLYEVLA ^{EQ} EGTTFLPNPSGGFHPGEEVLGSM	240
1168	A1552VC RS11070 WP 000957200.1		0
1293	VP RS12240 WP 005479695.1		0
1401	DSB67 RS12665 WP 010643259.1		0
1110	NGFG RS09235 WP 003689817.1		0
1251	GC085 RS07725 WP 011946523.1		0
404616	IX0842 RS07650 WP 011408072.1		0
1058	ABD1 RS01615 WP 001274986.1		0
1018	ACIAD RS01685 WP 004920473.1		0
consensus			

NTDB id	Accession	Protein	Sequence	Score
NTDB id 1306	DR RS10060 WP 027479822.1	LRADALRLLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGR	LGEQLVQEGLSRDQL	320
NTDB id 1168	A1552VC RS11070 WP 000957200.1	MLTNLVAIILRQAELISATQ.		19
NTDB id 1293	VP RS12240 WP 005479695.1	MHSNLSITILRQKGLLTFSQ.		19
NTDB id 1401	DSB67 RS12665 WP 010643259.1	MLTNLSTVLRQAGILTFSQ.		19
NTDB id 1110	NGFG RS09235 WP 003689817.1	MSVGLLRILVQNQVVTVEQ.		19
NTDB id 1251	GC085 RS07725 WP 011946523.1	MALATEEYRLQGIGQLLVLEKLLDKTK.		27
NTDB id 404616	IX0842 RS07650 WP 011408072.1	MNSVSTTNLVGITGIARRLVQDGALEETA.		29
NTDB id 1058	ABD1 RS01615 WP 001274986.1	MS..ALHTSPKFTGFFRRLVEEKHVSAAAT.		27
NTDB id 1018	ACIAD RS01685 WP 004920473.1	MS..AFTTPPKFSGFIRRLVEEGYVNAQN.		27
consensus			**** *!***** *	

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NTDB id 1306 DR RS10060 WP 027479822.1	REALQVQAREGKVKPLGEVITELGFASPD	400
NTDB id 1168 A1552VC RS11070 WP 000957200.1	EQAVVTQVSASGTSVPEALLSIFHAQELTEQLSHIFGPE	69
NTDB id 1293 VP RS12240 WP 005479695.1	EESLIEQVKASGISMPEALLSSGFFTSSELAEHLSSIFG	69
NTDB id 1401 DSB67 RS12665 WP 010643259.1	EESLIEQVKASGISMPEALLSSGFFQSHLTHSSVFCIS	69
NTDB id 1110 NGFG RS09235 WP 003689817.1	AEHYYNESQA.CKEVLPMLFSDGVI.SPKSLAALIA	68
NTDB id 1251 GC085 RS07725 WP 011946523.1	AIELHKLAAAEKISLLQYIVKNKILSAEQIALTA	77
NTDB id 404616 IX0842 RS07650 WP 011408072.1	ARTAMEQAAAQKVP.LPQWFAENRLVSASQLAA	79
NTDB id 1058 ABD1 RS01615 WP 001274986.1	MQTALDAAKRAKQDVTAYLIEEVHLSPSLAE	77
NTDB id 1018 ACIAD RS01685 WP 004920473.1	MQQALEKAKKFQKD.VPPLYIDNFS.SPLTTAB	77
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NTDB id 1306 DR RS10060 WP 027479822.1	DPKVALMIPATARRVYVVPVRLQGNLSLVAMKD	480
NTDB id 1168 A1552VC RS11070 WP 000957200.1	ANLCQQQLGLRELITRYDALPIAKQCNLLLLAV	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	HRVLMGVLTETHEQMVETHCVPVFRGDKV	148
NTDB id 1251 GC085 RS07725 WP 011946523.1	CTIPVNLVNEKLIKRRHAMVPLFSRGTNLYLAT	157
NTDB id 404616 IX0842 RS07650 WP 011408072.1	NQNAVKLVSLELLQKYHVLPLFKRCNRLFV	159
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SQIPKDLVDQKLLKHLRILPLIQRGQILY	157
NTDB id 1018 ACIAD RS01685 WP 004920473.1	ALFLKDKIDKLLIQYRIMPLVHRGHVLY	157
consensus	*****	

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NTDB id 1306 DR RS10060 WP 027479822.1	ELA.....ERNKTQSSQEA.DLSVADESALV	547
NTDB id 1168 A1552VC RS11070 WP 000957200.1	GK.EISQDELANLVK...VSDDELQSI.DLSQ	225
NTDB id 1293 VP RS12240 WP 005479695.1	LKEINQEELASLVD...VGADEINIEDLSQ	225
NTDB id 1401 DSB67 RS12665 WP 010643259.1	LKEINQDELAGLVD...IGDEIENIEDLSQ	225
NTDB id 1110 NGFG RS09235 WP 003689817.1	GEGQ.E.....EESHTLYIDNEEAEDG	215
NTDB id 1251 GC085 RS07725 WP 011946523.1	YVEDSGDLEGLETSADD...EDQDIDTATS	234
NTDB id 404616 IX0842 RS07650 WP 011408072.1	GNDD.EEMGDLDVSTGDEDMGAGG.DSGV	238
NTDB id 1058 ABD1 RS01615 WP 001274986.1	EE.FDLVDN..LDGPTAQEDEE	229
NTDB id 1018 ACIAD RS01685 WP 004920473.1	EE.LDLDVD..VDPHTTDDDED	229
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NTDB id 1306 DR RS10060 WP 027479822.1	FPKGAAQQIMARLIKIMGHLDIAERRVPQDGR	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 404616 IX0842 RS07650 WP 011408072.1	APVKLSQRIARAARIKIVMSQLDIAEKRPQDGR	317
NTDB id 1058 ABD1 RS01615 WP 001274986.1	PPLQLANRLASRIKIVMSQMDISEKRVPQDGR	308
NTDB id 1018 ACIAD RS01685 WP 004920473.1	PPLQLATRLASRIKIVMSQMDISEKRVPQDGR	308
consensus	*****	

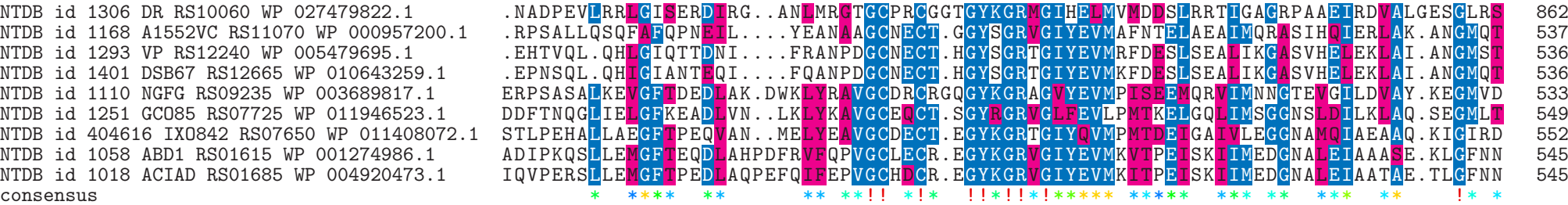
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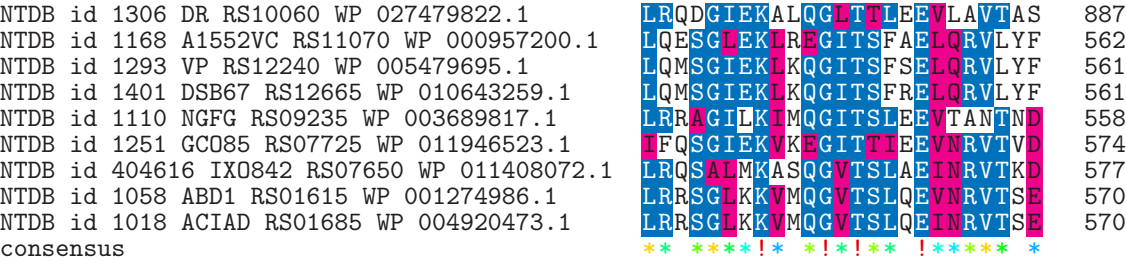
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