

NTDB id	Accession	Sequence	Count
NTDB id 1306	DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAALR	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 1102155	Q9280 RS07700 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	Length	Consensus
1306	DR RS10060 WP 027479822.1	160	SIRPQTALNLQAFPPFALEGGDRLRVALVDPLSSFSIETLEDDSGGFDIEPYQALREEVMMWAIATHYPELGLEIVVPSGASDA
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	
1102155	Q9280 RS07700 WP 011408072.1	0	
1058	ABD1 RS01615 WP 001274986.1	0	
1018	ACIAD RS01685 WP 004920473.1	0	
consensus			

NTDB id	Accession	Sequence	Count
1306	DR RS10060 WP 027479822.1	GRTGGKLG ^{ER} LITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGGFHPGEEVLGSM	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
1102155	Q9280 RS07700 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	MHSNLSTILRQKGLLTF SQ.		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 1102155	Q9280 RS07700 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	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	EEELLHQVKASCISMPEALLSSGFFQSHLTHSSVFC	69
NTDB id 1110 NGFG RS09235 WP 003689817.1	EEELLHQVKASCISMPEALLSSGFFQSHLTHSSVFC	68
NTDB id 1251 GC085 RS07725 WP 011946523.1	AEHYYNESQA.CKEVLPMLFSDGVI.SPKSLAALIA	77
NTDB id 1102155 Q9280 RS07700 WP 011408072.1	AEHYYNESQA.CKEVLPMLFSDGVI.SPKSLAALIA	79
NTDB id 1058 ABD1 RS01615 WP 001274986.1	AEHYYNESQA.CKEVLPMLFSDGVI.SPKSLAALIA	77
NTDB id 1018 ACIAD RS01685 WP 004920473.1	AEHYYNESQA.CKEVLPMLFSDGVI.SPKSLAALIA	77
consensus	*****	

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NTDB id 1306 DR RS10060 WP 027479822.1	DPKVALMIPATARRVYVVPVRLQGNLSLVAMKDPRN	480
NTDB id 1168 A1552VC RS11070 WP 000957200.1	ANLCQQQLGLRELITRYDALPIAKQCNLLLLAVSDPT	149
NTDB id 1293 VP RS12240 WP 005479695.1	ANLCQQQLGLRELITRYDALPIAKQCNLLLLAVSDPT	149
NTDB id 1401 DSB67 RS12665 WP 010643259.1	ANLCQQQLGLRELITRYDALPIAKQCNLLLLAVSDPT	149
NTDB id 1110 NGFG RS09235 WP 003689817.1	ANLCQQQLGLRELITRYDALPIAKQCNLLLLAVSDPT	148
NTDB id 1251 GC085 RS07725 WP 011946523.1	ANLCQQQLGLRELITRYDALPIAKQCNLLLLAVSDPT	157
NTDB id 1102155 Q9280 RS07700 WP 011408072.1	ANLCQQQLGLRELITRYDALPIAKQCNLLLLAVSDPT	159
NTDB id 1058 ABD1 RS01615 WP 001274986.1	ANLCQQQLGLRELITRYDALPIAKQCNLLLLAVSDPT	157
NTDB id 1018 ACIAD RS01685 WP 004920473.1	ANLCQQQLGLRELITRYDALPIAKQCNLLLLAVSDPT	157
consensus	*****	

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NTDB id 1306 DR RS10060 WP 027479822.1	ELA.....ERNKTTQSSQEA.DLSVADESALVQV	547
NTDB id 1168 A1552VC RS11070 WP 000957200.1	GKELISQDELANLVK...VSDDELQSIEDLSQDDSP	225
NTDB id 1293 VP RS12240 WP 005479695.1	LKEINQEEELASLVD...VGADENIDEDLSQDDSP	225
NTDB id 1401 DSB67 RS12665 WP 010643259.1	LKEINQEEELASLVD...VGADENIDEDLSQDDSP	225
NTDB id 1110 NGFG RS09235 WP 003689817.1	GEGQ.E.....EESHTLYIDNEEAEDGPVPRFI	215
NTDB id 1251 GC085 RS07725 WP 011946523.1	YVEDSGDLEGLETSADD...EDQDIDTATSVTD	234
NTDB id 1102155 Q9280 RS07700 WP 011408072.1	GND.D.EEMGDLDVSTGDEDMGAGGDSGVDAKGD	238
NTDB id 1058 ABD1 RS01615 WP 001274986.1	EE.FDLVDN..LDGPTAQEDEEAPQGD	229
NTDB id 1018 ACIAD RS01685 WP 004920473.1	EE.FDLVDN..LDGPTAQEDEEAPQGD	229
consensus	*****	

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NTDB id 1306 DR RS10060 WP 027479822.1	FPKGAAQQIMARLKIMGHLDIAERRVPQDGRVRF..	625
NTDB id 1168 A1552VC RS11070 WP 000957200.1	PASHLSRRRLAARIKILSKLDIAERRVPQDGRVRF..	304
NTDB id 1293 VP RS12240 WP 005479695.1	PPNHLSSRLSARIKILSKLDIAERRVPQDGRVRF..	304
NTDB id 1401 DSB67 RS12665 WP 010643259.1	PPNHLSSRLSARIKILSKLDIAERRVPQDGRVRF..	304
NTDB id 1110 NGFG RS09235 WP 003689817.1	PPNHLSSRLSARIKILSKLDIAERRVPQDGRVRF..	295
NTDB id 1251 GC085 RS07725 WP 011946523.1	PPNHLSSRLSARIKILSKLDIAERRVPQDGRVRF..	313
NTDB id 1102155 Q9280 RS07700 WP 011408072.1	PPNHLSSRLSARIKILSKLDIAERRVPQDGRVRF..	317
NTDB id 1058 ABD1 RS01615 WP 001274986.1	PPNHLSSRLSARIKILSKLDIAERRVPQDGRVRF..	308
NTDB id 1018 ACIAD RS01685 WP 004920473.1	PPNHLSSRLSARIKILSKLDIAERRVPQDGRVRF..	308
consensus	*****	

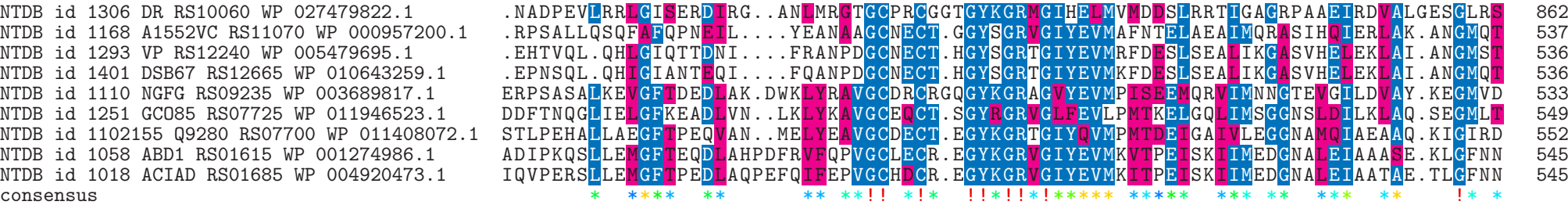
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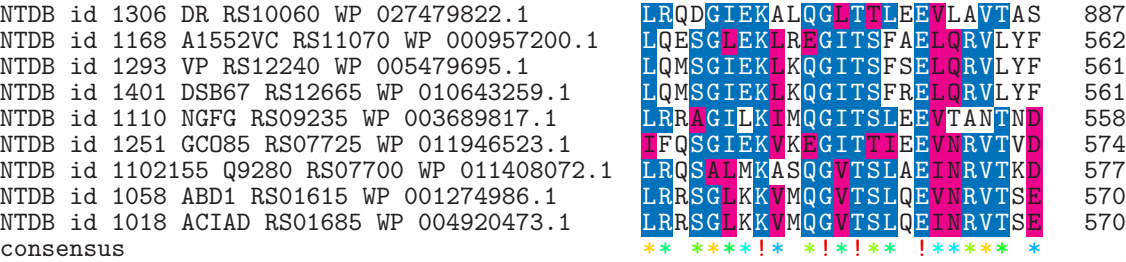
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