

NTDB id	Accession	Protein	Sequence	Score
NTDB id 1306	DR RS10060 WP 027479822.1	MALSI	CDRRRLGAILLDQGYLGDN	0
NTDB id 1168	A1552VC RS11070 WP 000957200.1	MALSIGDRRLGAILLDQGYLGDN	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 1102002	Q9282 RS15215 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 1102002	Q9282 RS15215 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} QLYEVLAEQEGTTFLPNPSGGFHPGEEVLGSM	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
1102002	Q9282 RS15215 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	LRADALRLLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRIGEQLVQEGLSRDQL	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 1102002	Q9282	RS15215 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 1168 A1552VC RS11070 WP 000957200.1
NTDB id 1293 VP RS12240 WP 005479695.1
NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 1110 NGFG RS09235 WP 003689817.1
NTDB id 1251 GCO85 RS07725 WP 011946523.1
NTDB id 1102002 Q9282 RS15215 WP 011408072.1
NTDB id 1058 ABD1 RS01615 WP 001274986.1
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consensus

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consensus
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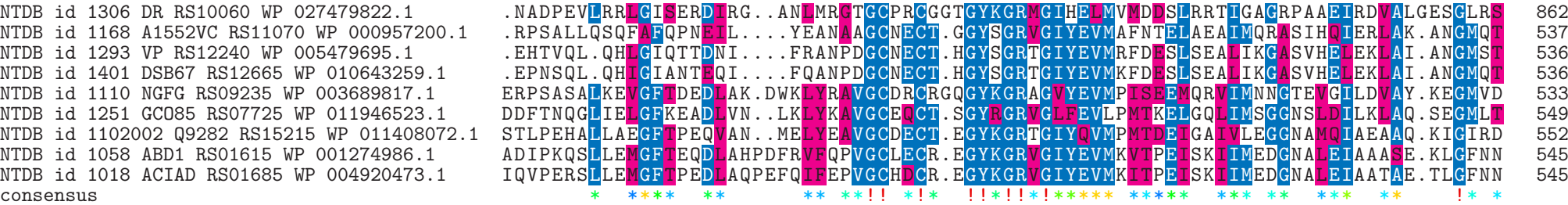
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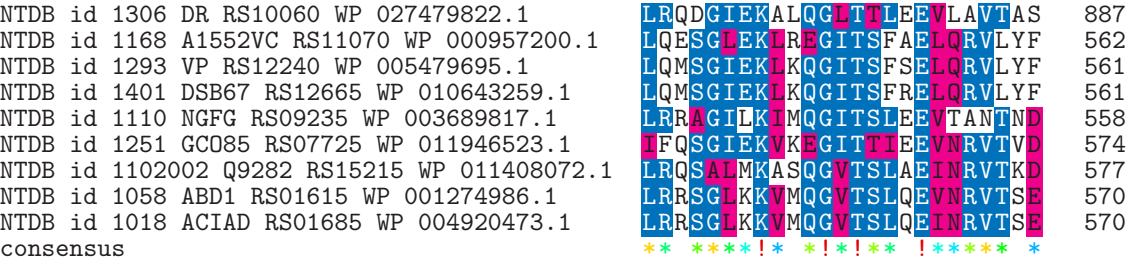
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 non conserved
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 $\geq 50\%$ conserved