

logo

NTDB id 1306 DR RS10060 WP 027479822.1	MALSI GDRRLGAILLDQGYLGDN DLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
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NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
consensus		

logo

NTDB id 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPPFALEGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEI WVPSGASDA	160
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
consensus		

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NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	LRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLGEQLVQEGT LSRDQL	320
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NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
consensus		

NTDB id 1306	DR RS10060 WP 027479822.1	REALQVQAREGKVKP	LGEVITELGFASPD	EVDSALQKQNVGG	RLEDTLVQSGKLSPEM	LARS	LAAQLGYEFLDPIQNP	400
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NTDB id 1293	VP RS12240 WP 005479695.1	MHSNLS	TILRQKGLLTF	SQEESSLIEQVKAS	GISMPEALLSSGFF	TSSELAEHLSSIFGLSQPELSQY	69	
NTDB id 1401	DSB67 RS12665 WP 010643259.1	MLTNLS	TVLRQAGILTF	SQEESSLLEQVKAS	GISMPEALLSSGFF	QSHELTEHLSVFGLSCTSLNQY	69	
NTDB id 50150	HT081 RS08765 WP 003689817.1	MSVGLLR	LRLVQNVVTV	EQAEHYY.NESQAGKEV	LPMFLSDGV	ISPKSLAALIA	RVFSYSIDLRLHYPR	68
NTDB id 1110	NGFG RS09235 WP 003689817.1	MSVGLLR	LRLVQNVVTV	EQAEHYY.NESQAGKEV	LPMFLSDGV	ISPKSLAALIA	RVFSYSIDLRLHYPR	68
NTDB id 1251	GC085 RS07725 WP 011946523.1	...MALATEEYRL	QCTGQLLVLEKLL	DKTKAIELHKLAAAEKIS	LLQYIVKNKILSAEQ	IALTA	AQNFVGVMPLDINCIDV	77
NTDB id 1018	ACTAD RS01685 WP 004920473.1	...MSAFTTPPKF	SGFIRRLVEEGYV	NAQNMQQALEKAKKFKQD	IVPYLIDNFSISPLT	IAEITISLEFGEPLDLGV	FDP	77
NTDB id 1058	ABD1 RS01615 WP 001274986.1	...MSALHTSPKF	TGFFRRLVBEKHVS	AATMQTALDAAKRAKQDT	VAYLIEEVHLSPLSL	LAETISAEFAEPYFLD	VDVYDT	77
consensus		*****						

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NTDB id	1058	ABD1 RS01615 WP	001274986.1
consensus			

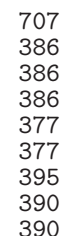
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NTDB id 1251	GC085 RS07725 WP 011946523.1	YVEDSGDLEGLAISAD.DEDQDIDTATSVT	DDAPIVKFVNKILLDAIRQGASDIHFEPYEREY	YRIRYRQDGIILHEVATPP	236
NTDB id 1018	ACTAD RS01685 WP 004920473.1	E...ELD...LDVEVDPHTTDDDDDEDK	KLKDEAPIVKYINKLLIDAIRMSGASDLHFEPY	EKSYRVRYVDGVLRLIATPP	231
NTDB id 1058	ABD1 RS01615 WP 001274986.1	E...EFD...LDVNLDPGTAQEDDEEAPQ	QDEAPIVKYINKLLIDAIRMSGASDLHFEPY	EKSYRVRYVDGVLRLQIANPP	231
consensus		* * * * *	* * * * *	* * * * *	

[illegible]

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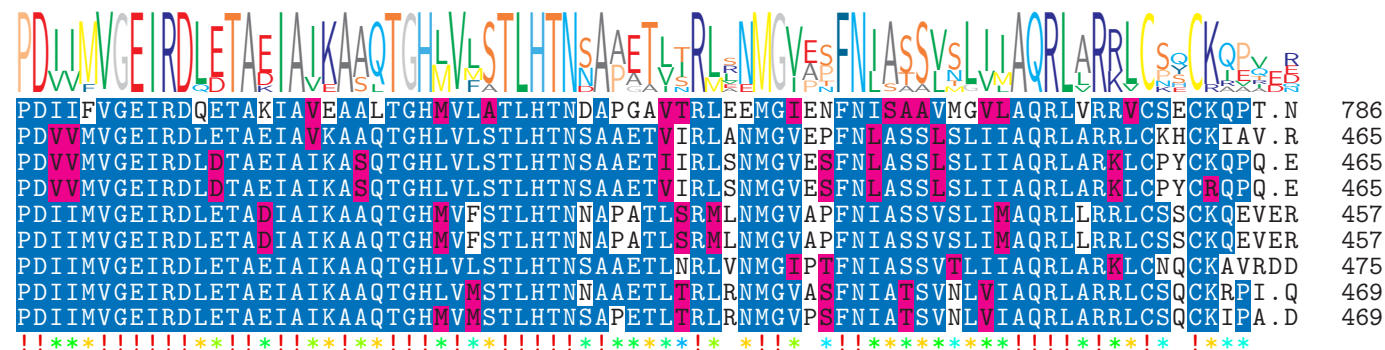
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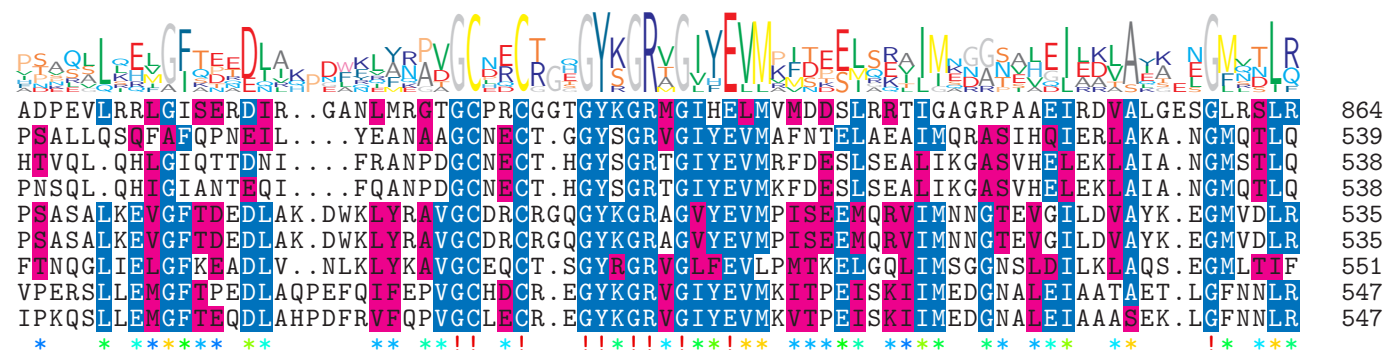
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
NTDB id 1058 ABD1 RS01615 WP 001274986.1
consensus

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NTDB id 1306 DR RS10060 WP 027479822.1
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NTDB id 1110 NGFG RS09235 WP 003689817.1
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
NTDB id 1058 ABD1 RS01615 WP 001274986.1
consensus

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 non conserved
 similar
 $\geq 50\%$ conserved