

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	MSXLSIGDRRLGAALLDQGYLDNBLQRALERHSEVGGRLAELVLYDSGGYSEERRIAEAEIESEFGLPLVNLHSEVETSPKXK	0
NTDB id 1306 DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	79
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
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NTDB id 408159 GP469 RS10570 WP 005493983.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	SLTPAEKAKELRAIPFALDEEAGVVRVAFLNPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVP	0
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NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVP	158
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	SDAGRTGGKLGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	0
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consensus		

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consensus		

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NTDB	id	1168	A1552VC	RS11070	WP	000957200.1	MLTNLVAAILRQAELISATQEQA VVTQVSASGTSVPEALLE..LSIFHAQELTEQ.....LSHF	57
NTDB	id	1401	DSB67	RS12665	WP	010643259.1	MLTNLSTVLRRAGILTFSQEESLLEQVKASGISMP EALLS..SGFFQSHELTEH.....LSSVF	57
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NTDB	id	1293	VP	RS12240	WP	005479695.1	MHSNLTSTILRQKGLLTFSQEESLIEQVKASGISMPEALLS..SGFFTSSSELAEH.....LSSIF	57
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NTDB	id	1251	GC085	RS07725	WP	011946523.1	MALATEEYRLQGIGQLLVLEKLIDKTKAIELHLKAAA EKISLLQYIVK..NKILSAEQIALT.....AAQNF	65
NTDB	id	1058	ABD1	RS01615	WP	001274986.1	MSALHTSPKFTGFFRRLVEEKHVS AATMQTALDAAKRAKQDTVAYLIE..EVHLSPSL LAET.....ISA EF	65
NTDB	id	1018	ACIAD	RS01685	WP	004920473.1	MSAFTTPPKFSGFIRRLVEEGYVNAQMMALEKAKKF KQDITVPYLID..NFSISPLTIAEI.....ISLE F	65
consensus							* * * * *	

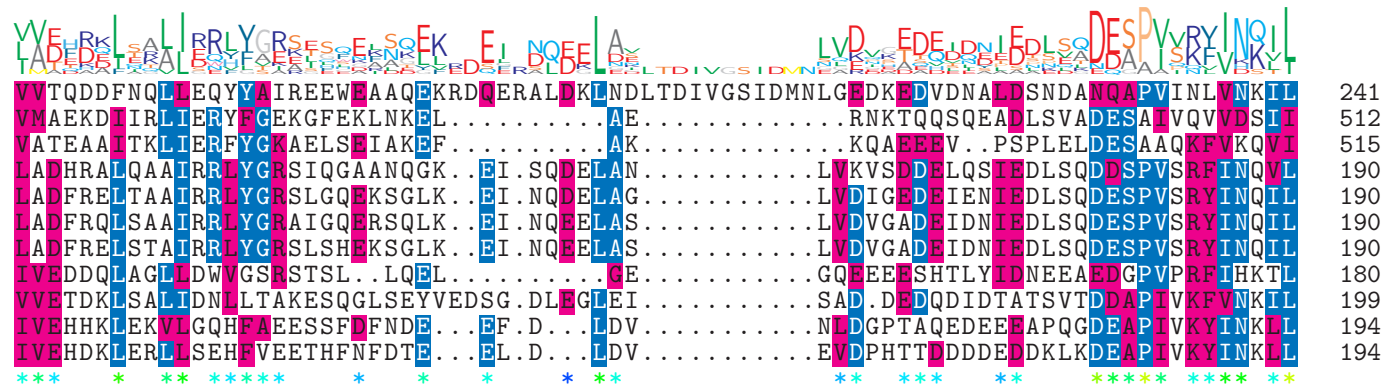
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	GLSCTSLNQYEYA.....SLCQTGLGLRLITRHNAFLPNRT...SSTLLI LAVADPTNLQAEDDFRFATG.....LQVELV	124
NTDB id 408159 GP469 RS10570 WP 005493983.1	GLSQPESLSQYEYA.....SLCQQLGLRELITRHNAFLPHRT...PSTLLLAVADPTNQQAEDDFRFATG.....LQVELV	124
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NTDB id 1110 NGFG RS09235 WP 003689817.1	SYSILDLRHYPRH....RVLMGVLTTEEQMVEFHCVPFRR...GDKVFFAVSDPTQMPQIQKTIVSAAG.....IAVELV	123
NTDB id 1251 GC085 RS07725 WP 011946523.1	GVPMLDINCIDVG....TIPVNLVNEKLIKRRHAMVPLFSR...GTNLYLATDPSKQASLKETIQFHTG.....LNTHAI	132
NTDB id 1058 ABD1 RS01615 WP 001274986.1	AEPYFDLDVYDTS....QIPKDLVDQKLLKKHRILLPLIQR...GQILYVATSNPSTNTEAIDAIRFNISK....LLVEPV	132
NTDB id 1018 ACIAD RS01685 WP 004920473.1	GEPLLDLGVEDPA....LFLKDKIDEKLIQKYRIMPLVHR...GHVLYVATSNPNTNTEAMDADAIRFNISK....LKVEPI	132
consensus	* * * * *	

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consensus

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consensus

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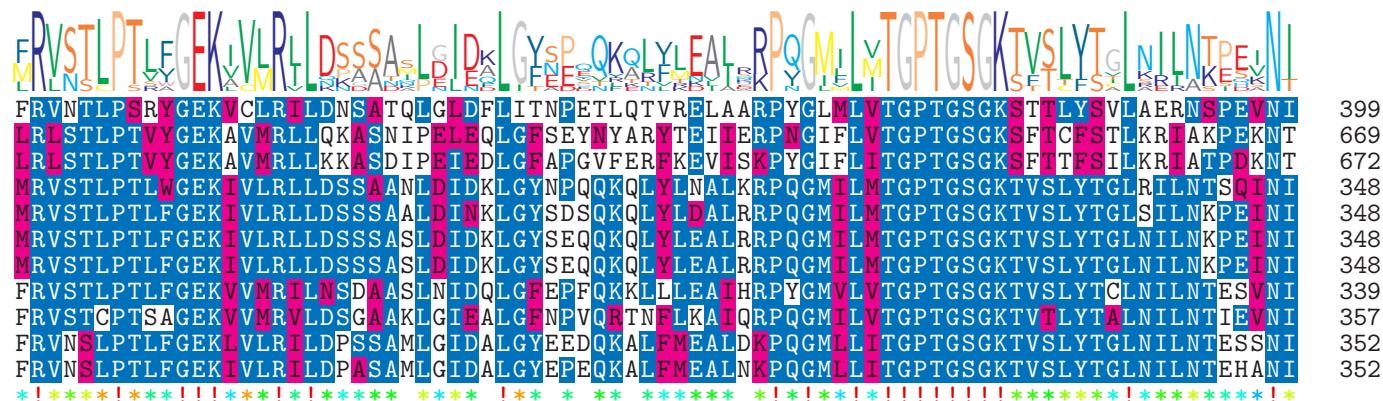


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consensus

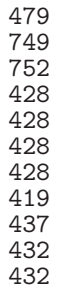
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consensus

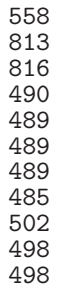
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consensus

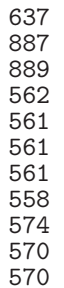
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NTDB id 1251 GC085 RS07725 WP 011946523.1
NTDB id 1058 ABD1 RS01615 WP 001274986.1
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consensus

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NTDB id 408159 GP469 RS10570 WP 005493983.1		561
NTDB id 1293 VP RS12240 WP 005479695.1		561
NTDB id 1110 NGFG RS09235 WP 003689817.1		558
NTDB id 1251 GC085 RS07725 WP 011946523.1		574
NTDB id 1058 ABD1 RS01615 WP 001274986.1		570
NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved