

logo

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NTDB id 1028 TT RS08215 WP 011173991.1		MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
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NTDB id 1293 VP RS12240 WP 005479695.1			0
NTDB id 1110 NGFG RS09235 WP 003689817.1			0
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NTDB id 1018 ACIAD RS01685 WP 004920473.1			0
consensus			

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consensus			

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consensus			

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consensus		

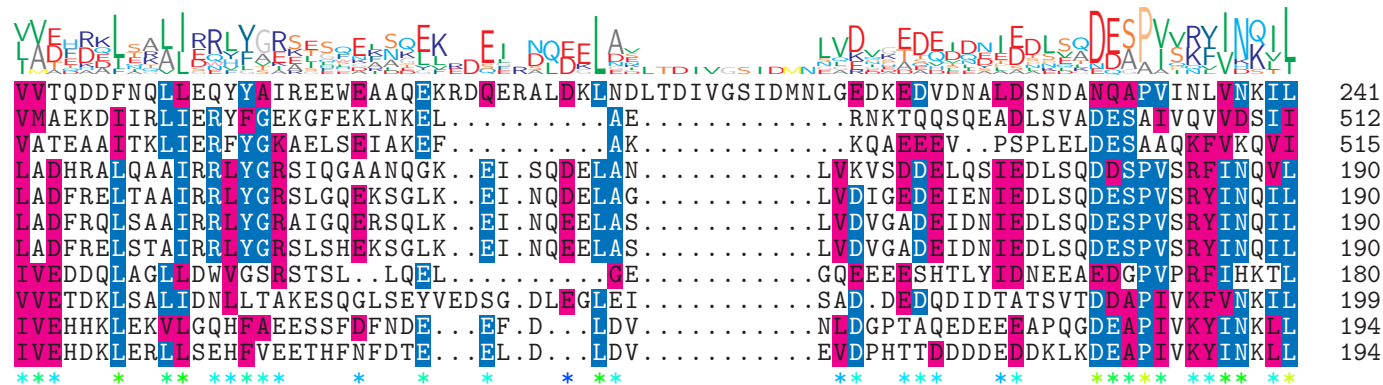
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NTDB	id	1251	GC085 RS07725 WP 011946523.1	MALATEEYRLQGIGQLLV	LEKLLDKTKAIELHLK	AAAEEKISLLQYIVK..NKIL	SAEQIALT.....	AAQNF	65	
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NTDB	id	1018	ACIAD RS01685 WP 004920473.1	MSAFTTPPKFSGFIRRLV	EEGYVNAQMMALEKA	KKKFKQDTIPVYLID..NFS	ISPLTI AEI.....	ISLEF	65	
consensus				* * *	* * *	* * *	* * *	* * *	* * *	* * *

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NTDB id 1028 TT RS08215 WP 011173991.1	GYPYVDEEDPPD....PGAPLLLPEDLCRRYGVFPHRLE...GNRLVLLMKDPRNILALDDVRLALKRKGLNIEVAPA	460
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	GLSCTSLNQYEYA.....SLCQTGLGLRLITRHNAFLPNRT...SSTLLIADAPTNLQAEDDFRFATG.....LQVELV	124
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	GEPLLDLGVEDPA....LFLKDKIDEKLIQKYRIMPLVHR...GHVLYVATSNPTNIEAMDIAIRFNSK.....LKVEPI	132
consensus	* * * * *	

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consensus

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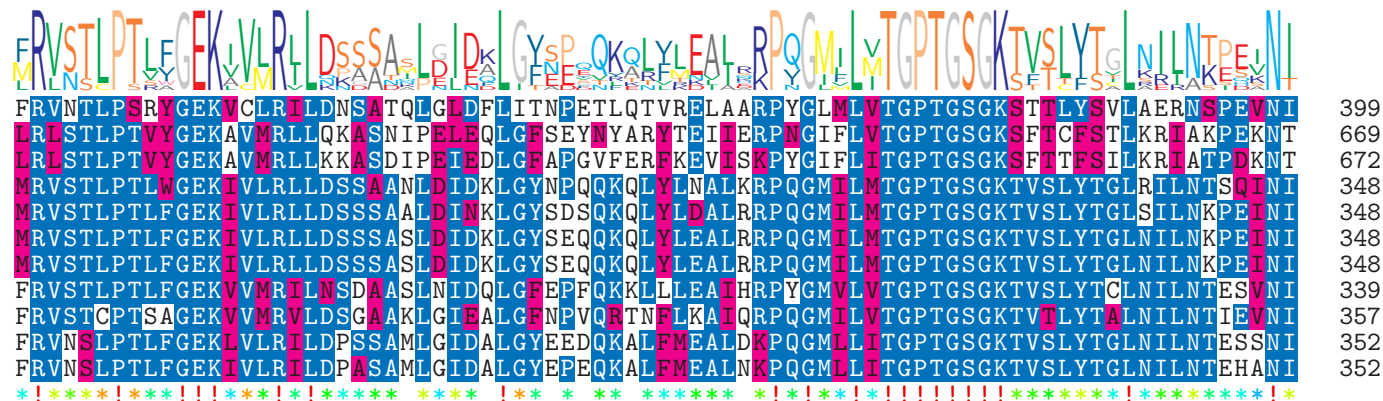
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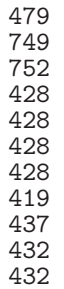
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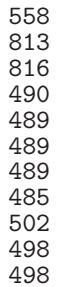
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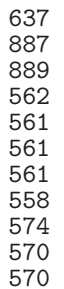
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consensus

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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