

logo

		MSXLSIGDRRLGAALLDQGYLDNBLQRALERHSEVGGRLAELVLYDSGGYSEERRIAEAEIESEFGLPLVLEHSEVETSPKXK	
NTDB id 1410 SGL RS13950 WP 010873519.1			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 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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		SLTPAEKAKELRAIPFALDEEAGVVRVAFLNPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLEIVVPSGA	
NTDB id 1410 SGL RS13950 WP 010873519.1	RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA		0
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NTDB id 1018 ACIAD RS01685 WP 004920473.1			0
consensus			

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		SGSGSGLKLGEELELYGQGYIIBSLLEYALQERQGTGSLGSLTYSSRLYFEEELYEXLAEQEGTEFLSEFEGHYPSEEXK	
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NTDB id 1028 TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA		237
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consensus			

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consensus

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[illegible]

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consensus

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[illegible]

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[illegible]

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NTDB id 1058 ABD1 RS01615 WP 001274986.1
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consensus

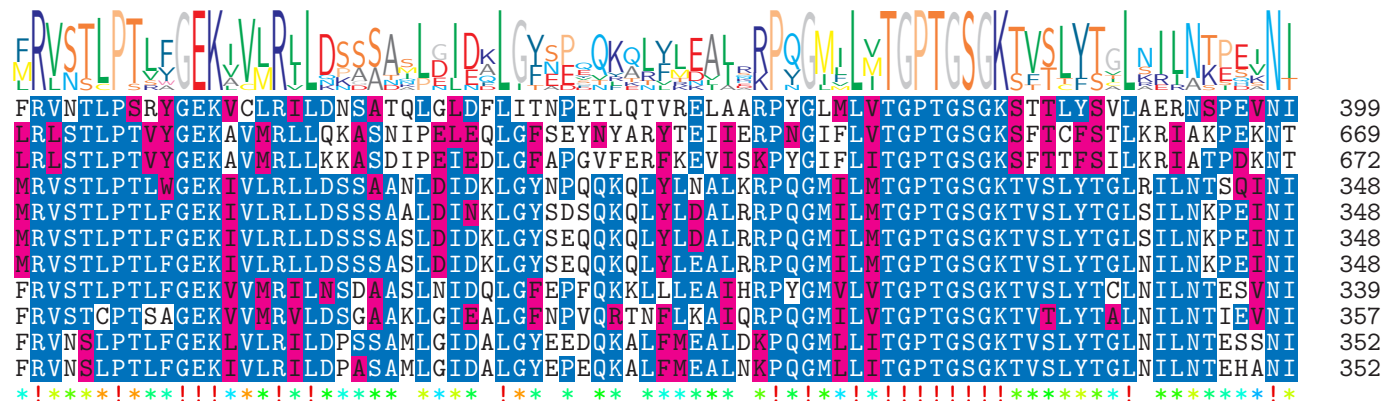
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consensus

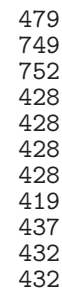
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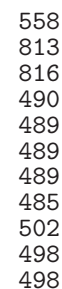
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

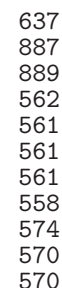
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NTDB id 1110 NGFG RS09235 WP 003689817.1
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NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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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