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NTDB id 1410 SGL RS13950 WP 010873519.1	MSXLTIGDrRLGAALLDQGYLDNQLQRALERHhEVGGhLAhVLYDhGhEhRRIABhIEhGhIPLVhLhVhVhPxxk	0
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1306 DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLIIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	79
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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 1018 ACIAD RS01685 WP 004920473.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	hEhGhEhKAKELRAhPFALhEhEhRVAhYhPLhGhStEhYEDhGhYhEPYQhTtRtYhAhAhHYPELhGLEhYhPshk	0
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consensus		

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consensus		

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consensus		

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NTDB id 1018 ACIAD RS01685 WP 004920473.1	MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQAIEKAKKFKQDIVPYLID..NFSISPLTIAEI.....ISLEF	65
NTDB id 1058 ABD1 RS01615 WP 001274986.1	MSALHTSPKFTGFFRRLVEEKHVSAATMQTALDAAKRAKQDTVAYLIE..EVHLSPSLLAET.....ISAEF	65
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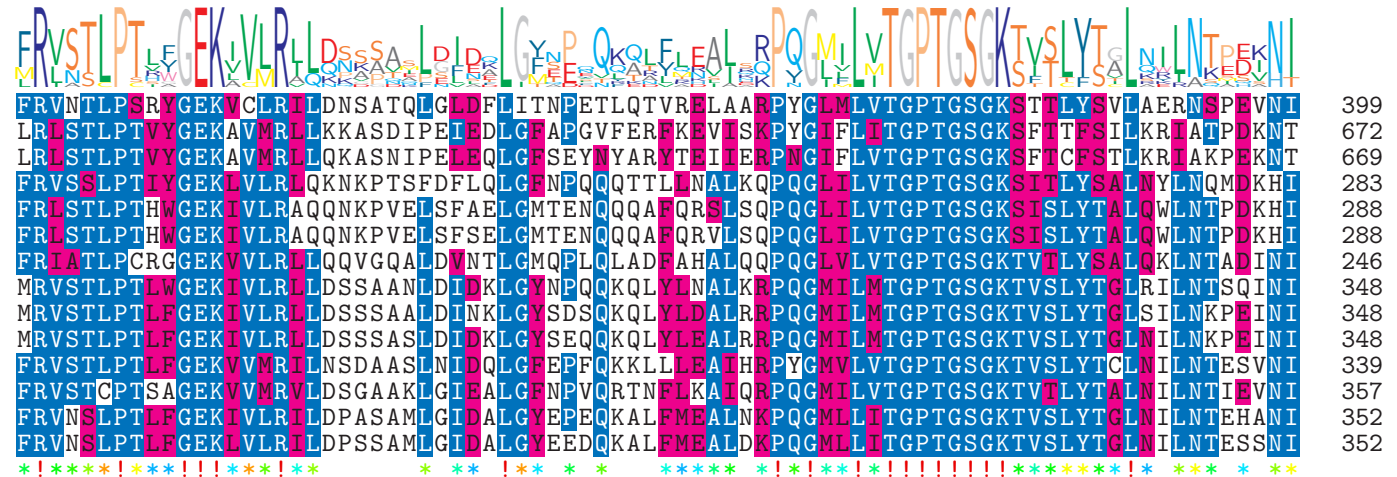
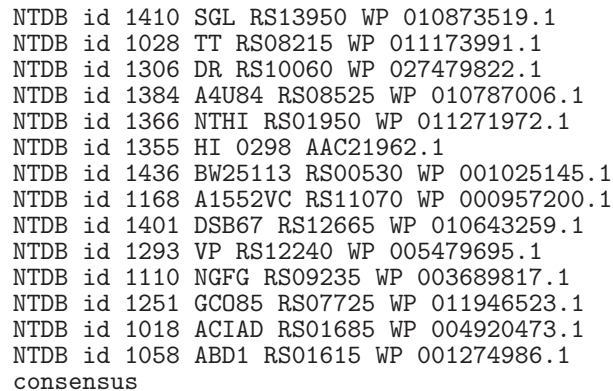
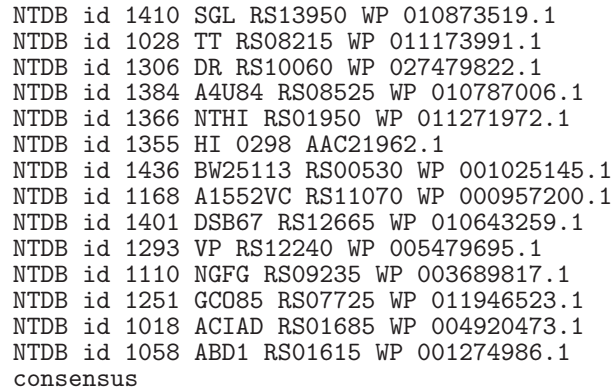
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	GEPLLDLGVFDPAL....FLKDKIDEKLIQKYRIMPLVHRG...HVLYVATSNPTNIEAMDAIRFNSK....LKVEPI	132
NTDB id 1058 ABD1 RS01615 WP 001274986.1	AEPYFDLDVYDTSQ....IPKDLVDQKLILKHRILPLIQRG...QILYVATSNPSNIEAIDAIRFNSK....LLVEPV	132
consensus	** * * * *	

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consensus

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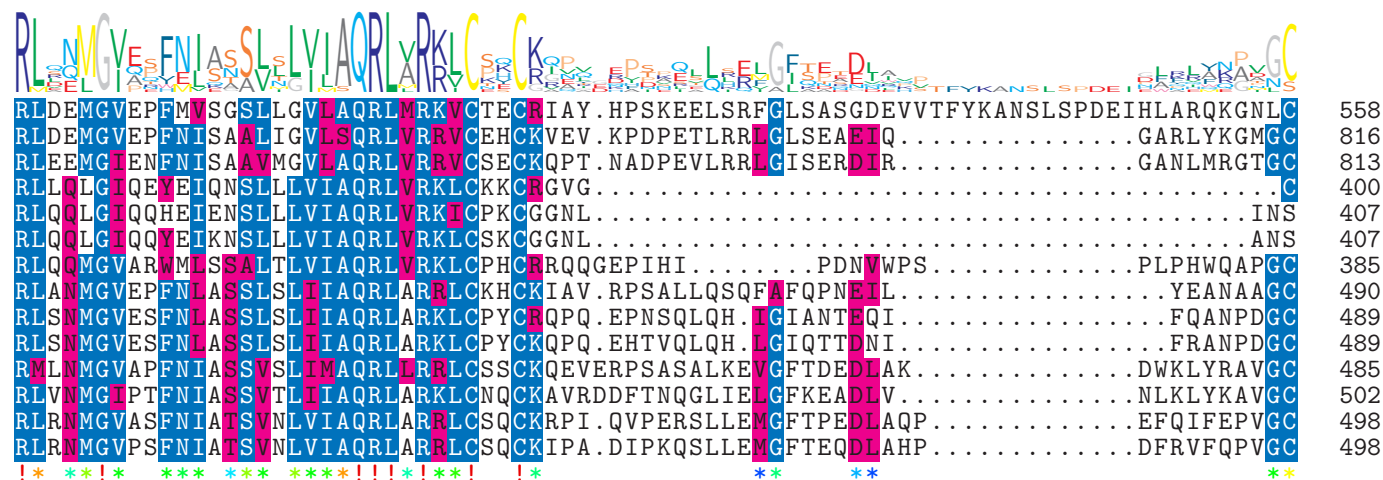
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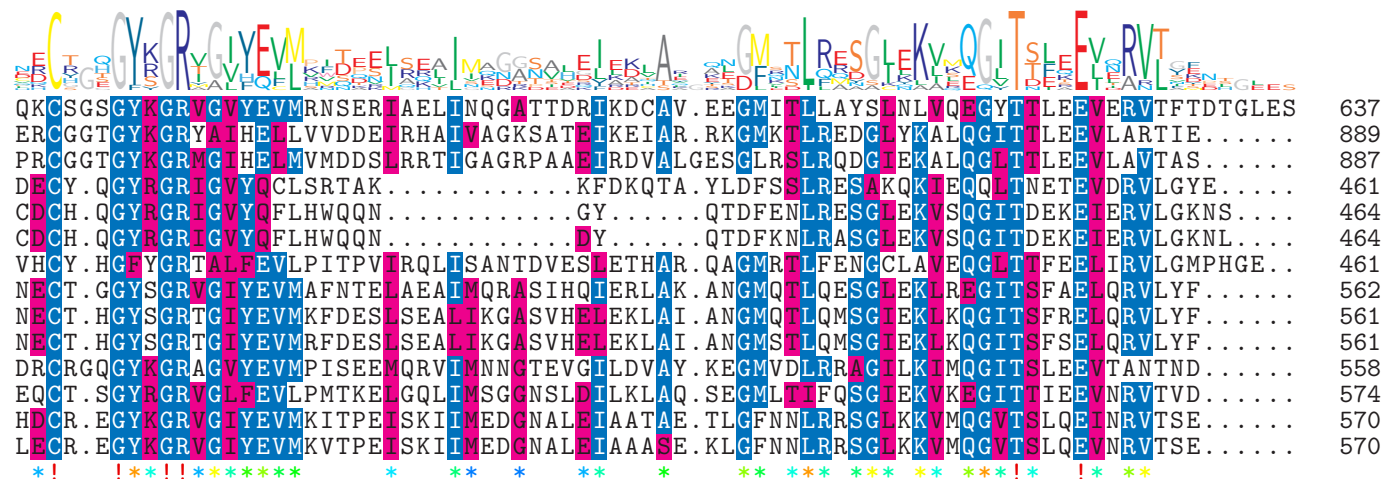
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consensus

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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved