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		MSXLSLTCDFRLGAAALLDQGYLQDHSLQRALERHSEVGGRLAEVLYDSQYSEERRIASXIESEHGIPLVQLHxVensPxxk	
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NTDB id 1306	DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAL	79
NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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consensus			

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NTDB id 1028	TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSEEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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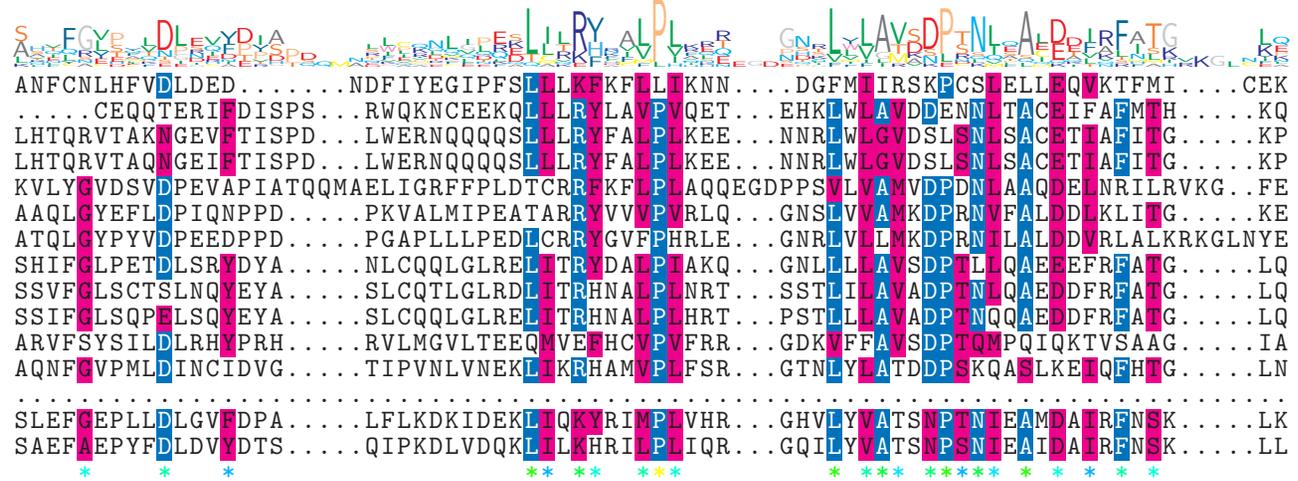
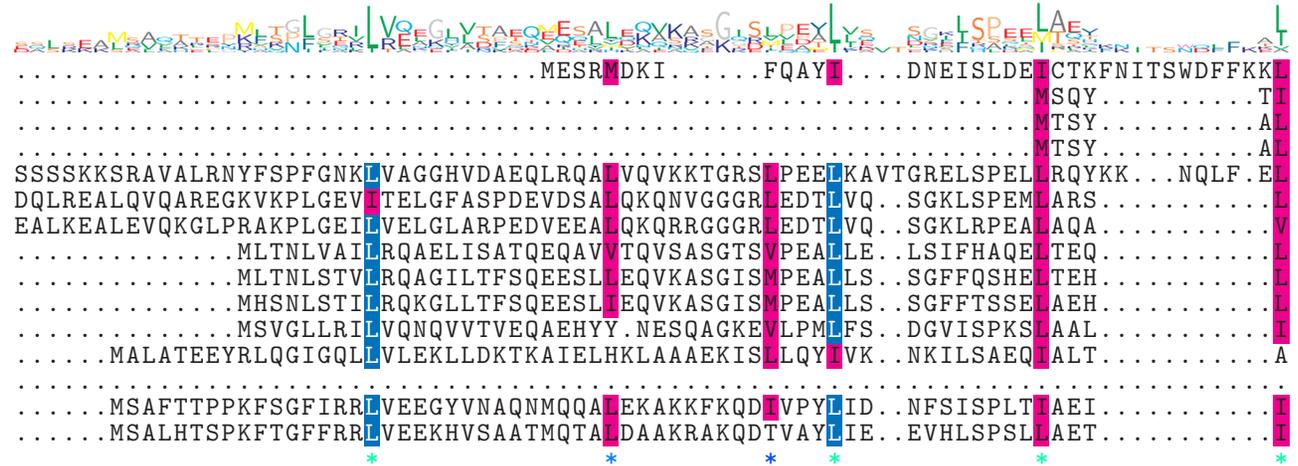
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NTDB id 1227	CJJ81176	RS07050	WP 011812800.1	.....	0																																																																				
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NTDB id 1410	SGL	RS13950	WP 010873519.1	.....	0																																																																				
NTDB id 1306	DR	RS10060	WP 027479822.1	SDAGRTGGKLGGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237																																																																				
NTDB id 1028	TT	RS08215	WP 011173991.1	SGEGQKDLKLGELLQKQGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA	237																																																																				
NTDB id 1168	A1552VC	RS11070	WP 000957200.1	.....	0																																																																				
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NTDB id 1058	ABD1	RS01615	WP 001274986.1	.....	0																																																																				
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NTDB id 1227 CJJ81176 RS07050 WP 011812800.1  
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1  
 consensus



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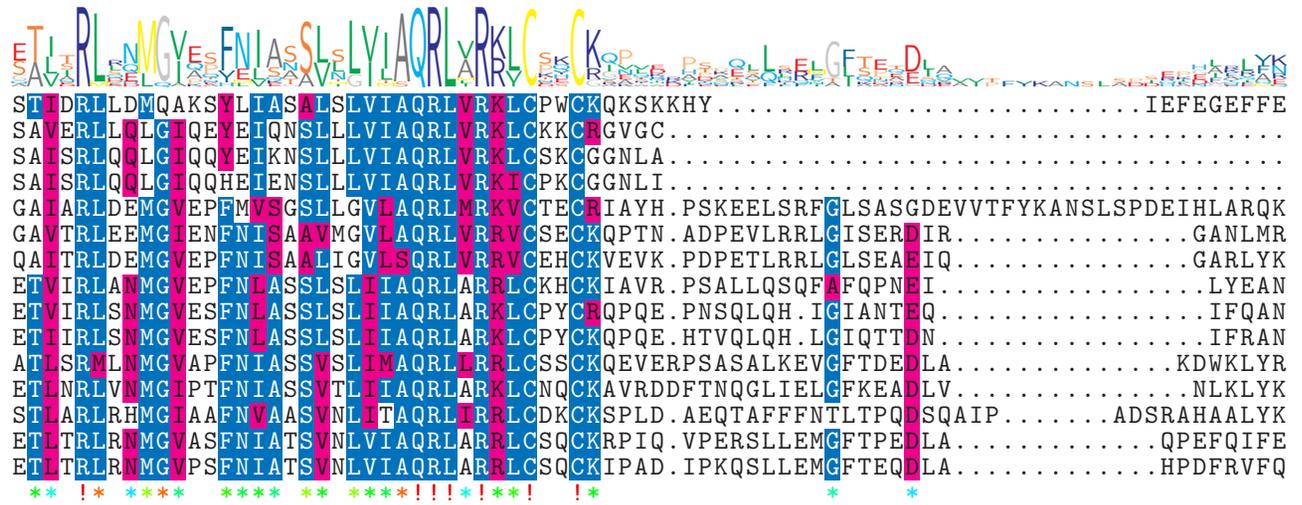
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 consensus



443  
400  
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NTDB id 1227 CJJ81176 RS07050 WP 011812800.1  
 NTDB id 1384 A4U84 RS08525 WP 010787006.1  
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 consensus



519  
461  
464  
464  
633  
887  
889  
562  
561  
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558  
574  
406  
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570

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	GLESEMRAKRKKSALTCRTCAAELQQEWLDCPYCMTSRFS	
NTDB id 1227 CJJ81176 RS07050 WP 011812800.1	.....	519
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	.....	570
consensus		

 non conserved  
 similar  
 ≥ 50% conserved