

logo

		MSXLSIGDRRLGALLDQGYLDRDLQRALERHSEVGGRLAEVLYDSQYSEERRIABXIEBSEFGIPLVQLHREVERBPSXK	
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NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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consensus			

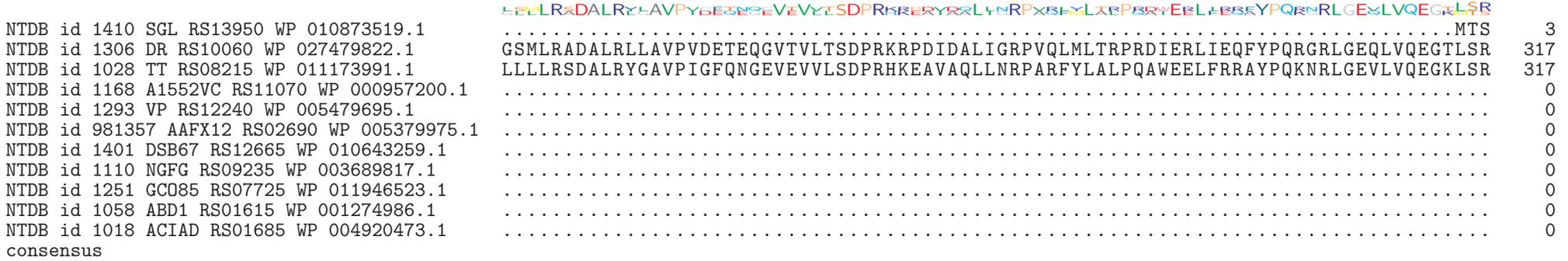
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		RSIRPQTAALNLQAFPFALE.GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIVVPSGA	
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NTDB id 1028	TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDPLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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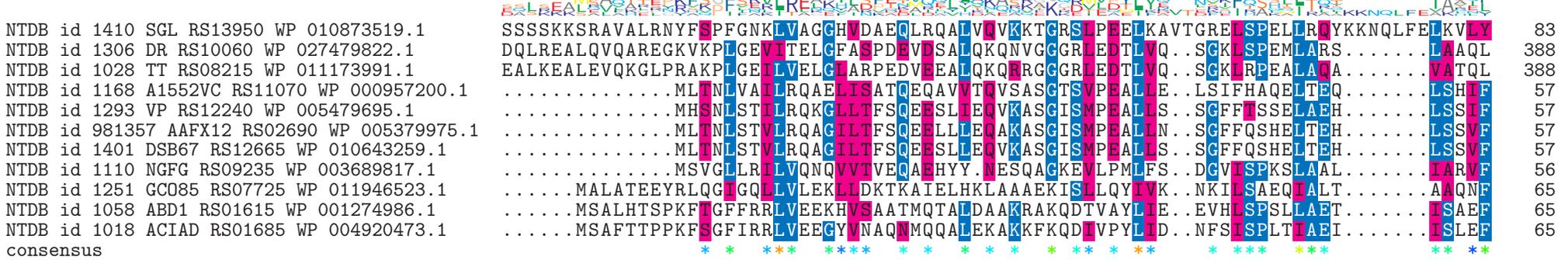
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NTDB id 1028	TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKQGWISREALEEALVEQEKTDLLGRILV.RKGLPEEALYRALAEQKGLFLESTEGIVDPDPSAA	237
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consensus			

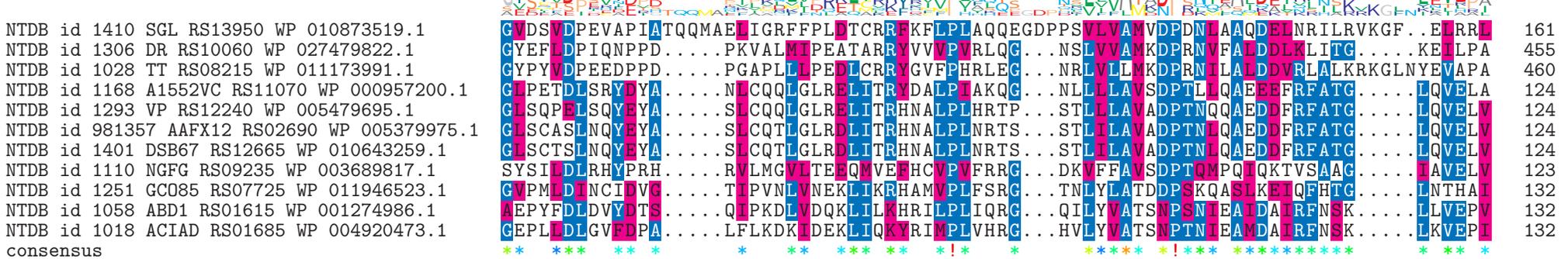
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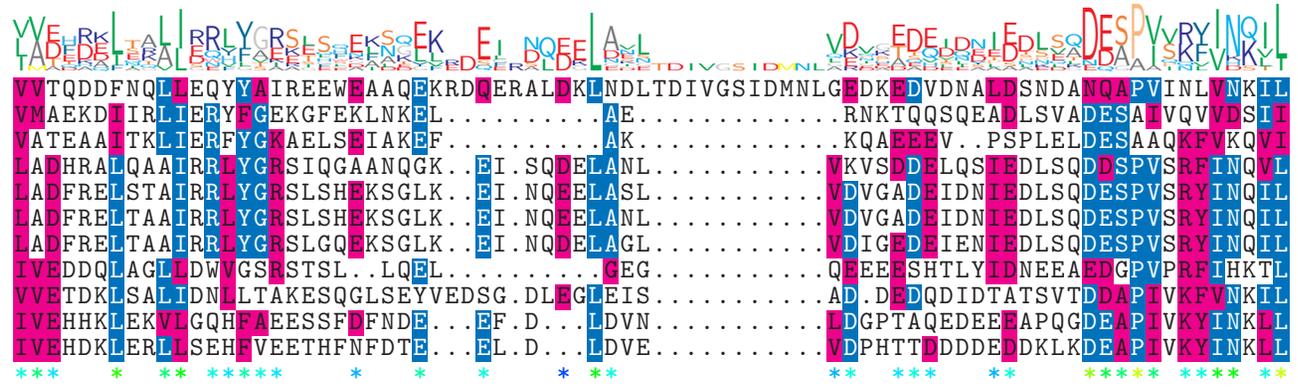


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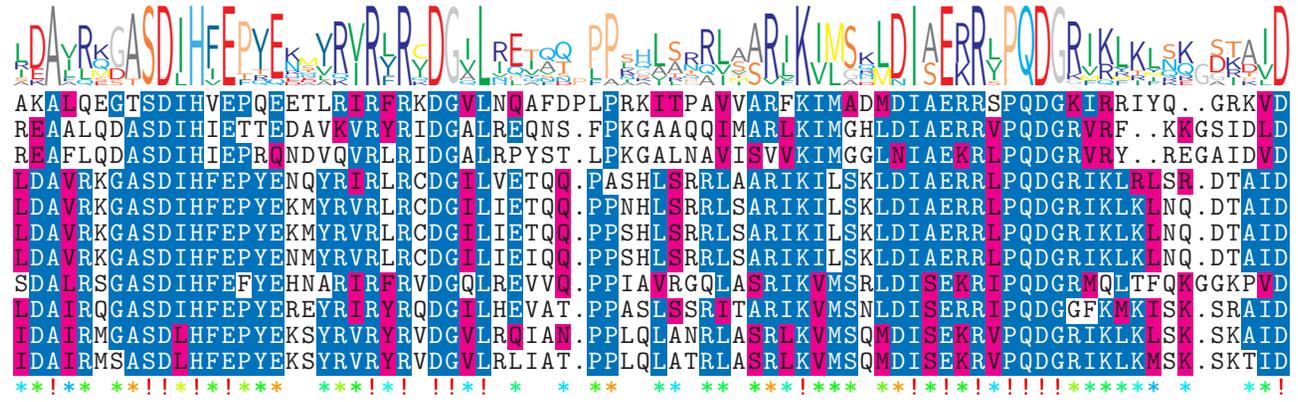
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1  
 consensus



241  
512  
515  
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NTDB id 1410 SGL RS13950 WP 010873519.1  
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 NTDB id 1251 GC085 RS07725 WP 011946523.1  
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1  
 consensus



319  
589  
592  
268  
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NTDB id 1410 SGL RS13950 WP 010873519.1  
 NTDB id 1306 DR RS10060 WP 027479822.1  
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 NTDB id 1251 GC085 RS07725 WP 011946523.1  
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 consensus



399  
669  
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348  
339  
357  
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352



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NTDB id 1410 SGL RS13950 WP 010873519.1	EMRAKRKRSALTCRTCAAELQQEWLDCPYCMTSRFS	672
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	.....	561
NTDB id 1110 NGFG RS09235 WP 003689817.1	.....	558
NTDB id 1251 GCO85 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