

logo

		MSXLSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAL	
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NTDB id 1028 TT RS08215 WP 011173991.1		MSVLTIGDKRLGAILLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
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consensus			

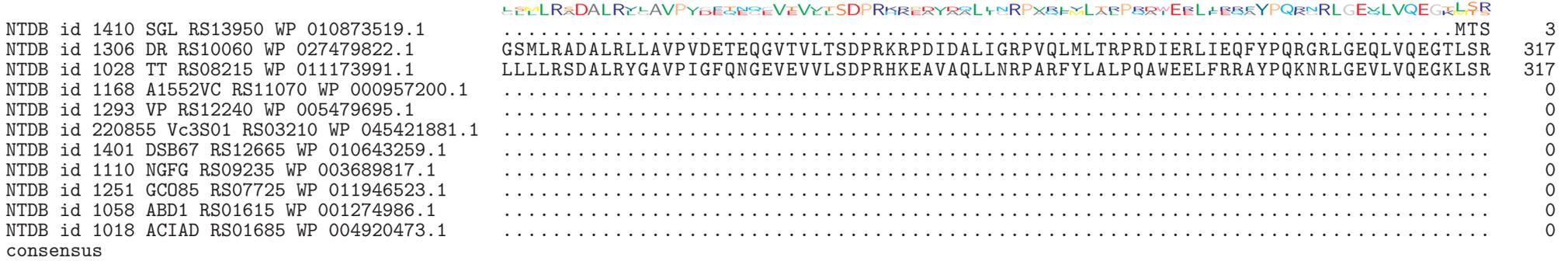
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		RSIRPQTALNLQAFPFALE.GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLIIVVPSGA	
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NTDB id 1028 TT RS08215 WP 011173991.1		ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDPLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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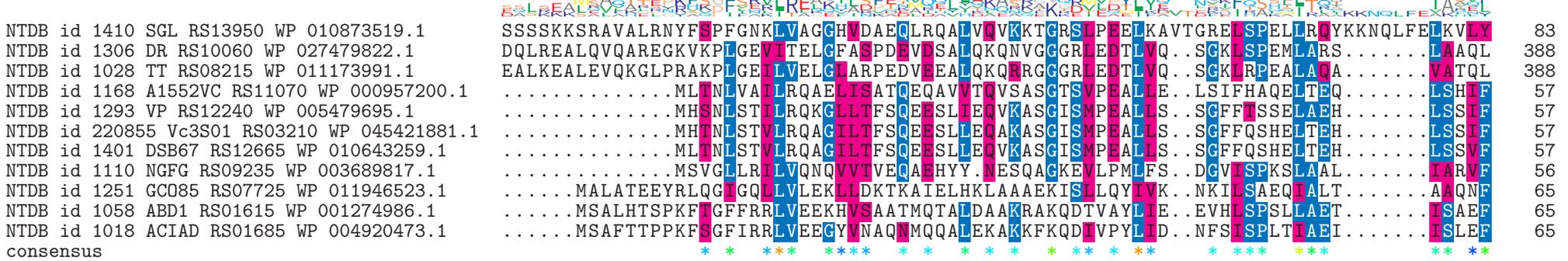
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NTDB id 1028 TT RS08215 WP 011173991.1		SGEGQKDLKLGELLQKQGWISREALEEALVEQEKTDLLGRILV.RKGLPEEALYRALAEQKGLFLESTEGIVDPDPSAA	237
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consensus			

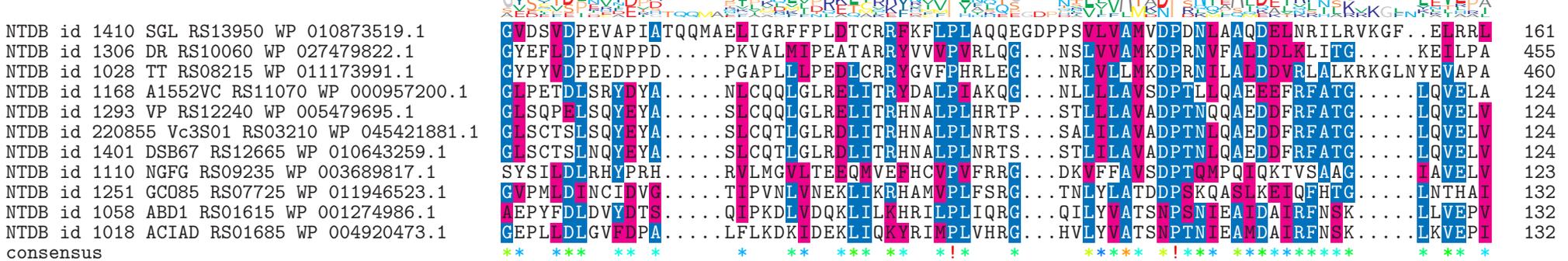
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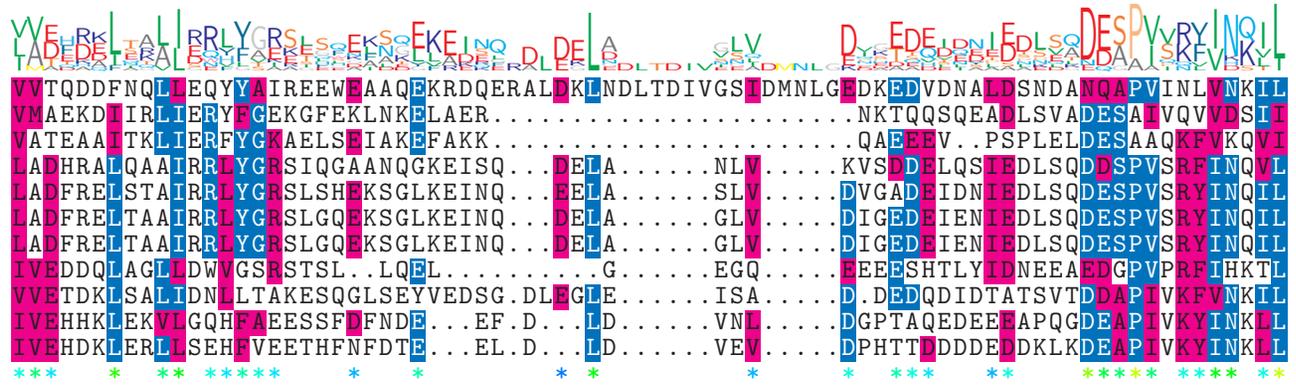


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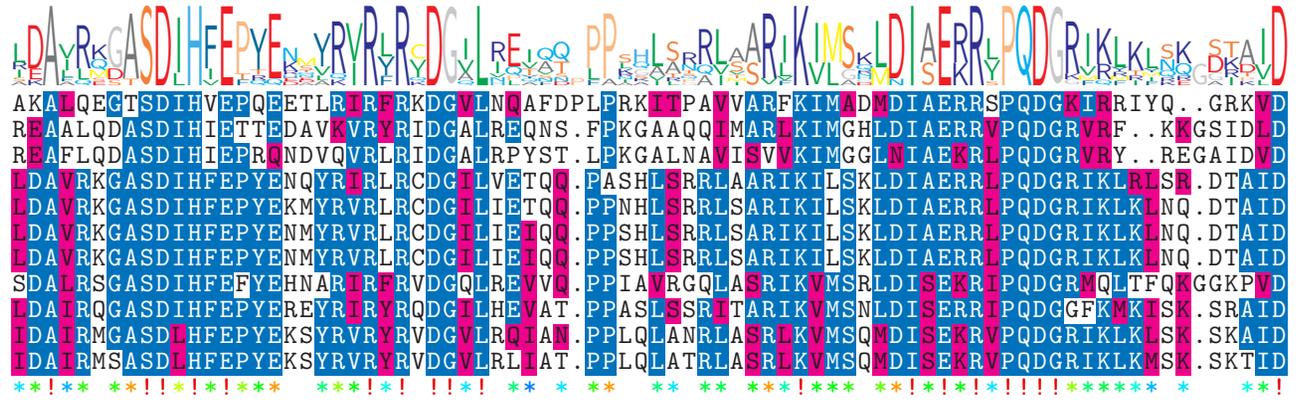
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1  
 consensus



241  
512  
515  
190  
190  
190  
190  
180  
199  
194  
194

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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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 consensus



319  
589  
592  
268  
268  
268  
268  
259  
277  
272  
272

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NTDB id 1410 SGL RS13950 WP 010873519.1  
 NTDB id 1306 DR RS10060 WP 027479822.1  
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 consensus



399  
669  
672  
348  
348  
348  
348  
339  
357  
352  
352



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NTDB id 1410 SGL RS13950 WP 010873519.1	EMRAKRKRSALTCRTCAAELQQEWLDCPYCMTSRFS	672
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NTDB id 220855 Vc3S01 RS03210 WP 045421881.1	.....	561
NTDB id 1401 DSB67 RS12665 WP 010643259.1	.....	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
- $\geq 50\%$  conserved