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		MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVEVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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NTDB id 1366	NTHI RS01950 WP 011271972.1	0
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NTDB id 1384	A4U84 RS08525 WP 010787006.1	0
NTDB id 1468	ASU2 RS05090 WP 014991734.1	0
NTDB id 22018	SCH RS00730 WP 000650610.1	0
NTDB id 1436	BW25113 RS00530 WP 001025145.1	0
NTDB id 1168	A1552VC RS11070 WP 000957200.1	0
NTDB id 1293	VP RS12240 WP 005479695.1	0
NTDB id 1401	DSB67 RS12665 WP 010643259.1	0
NTDB id 1251	GCO85 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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		ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSEEEVDTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
NTDB id 1028	TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSEEEVDTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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NTDB id 1251	GCO85 RS07725 WP 011946523.1	0
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consensus			

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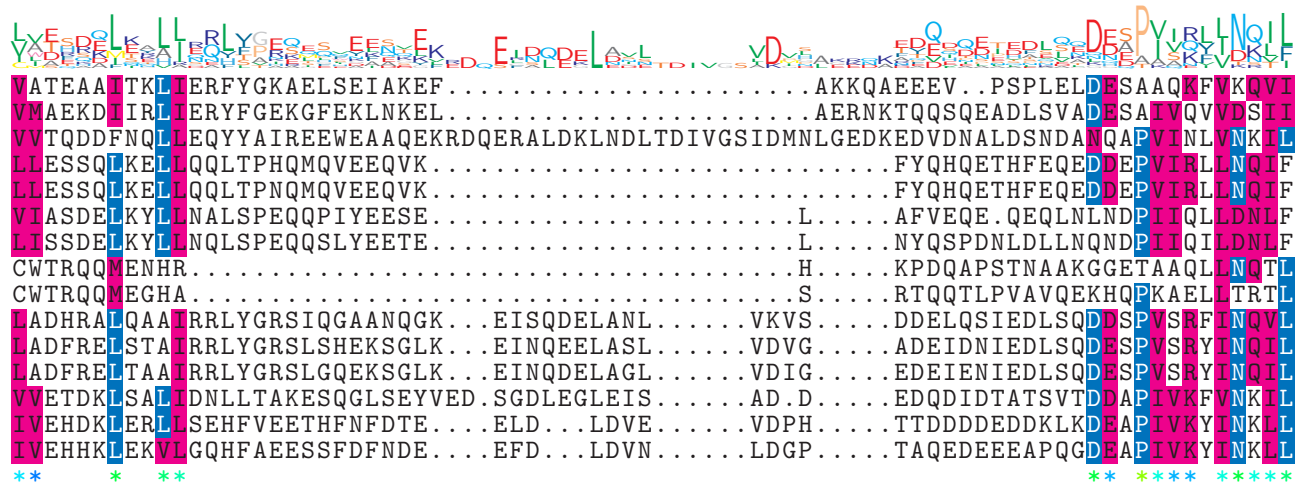
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consensus		

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consensus		

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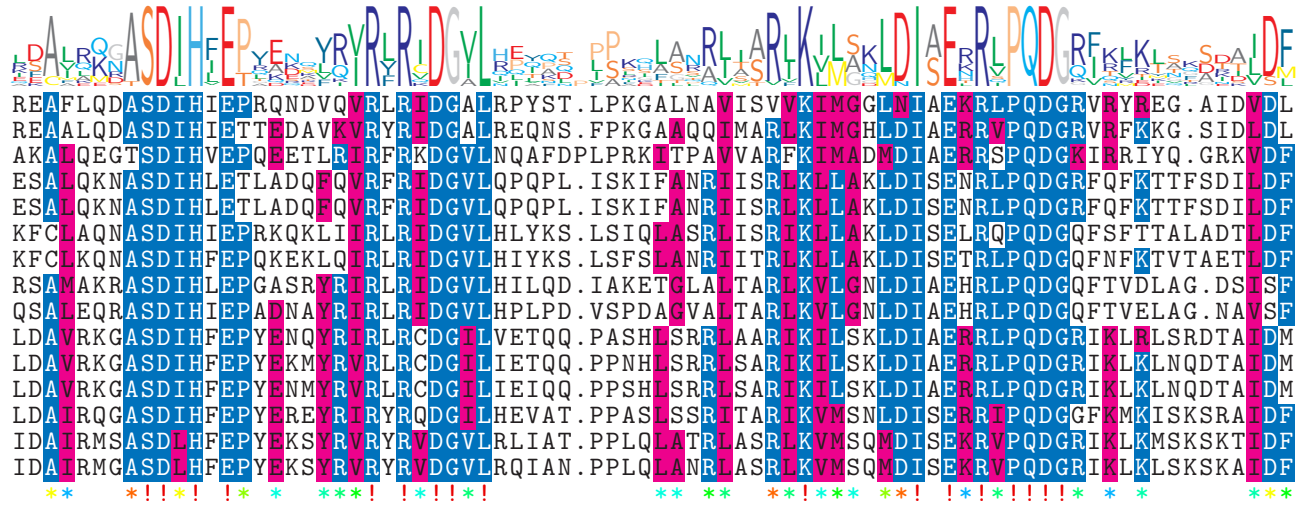
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 consensus



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logo

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 consensus



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consensus



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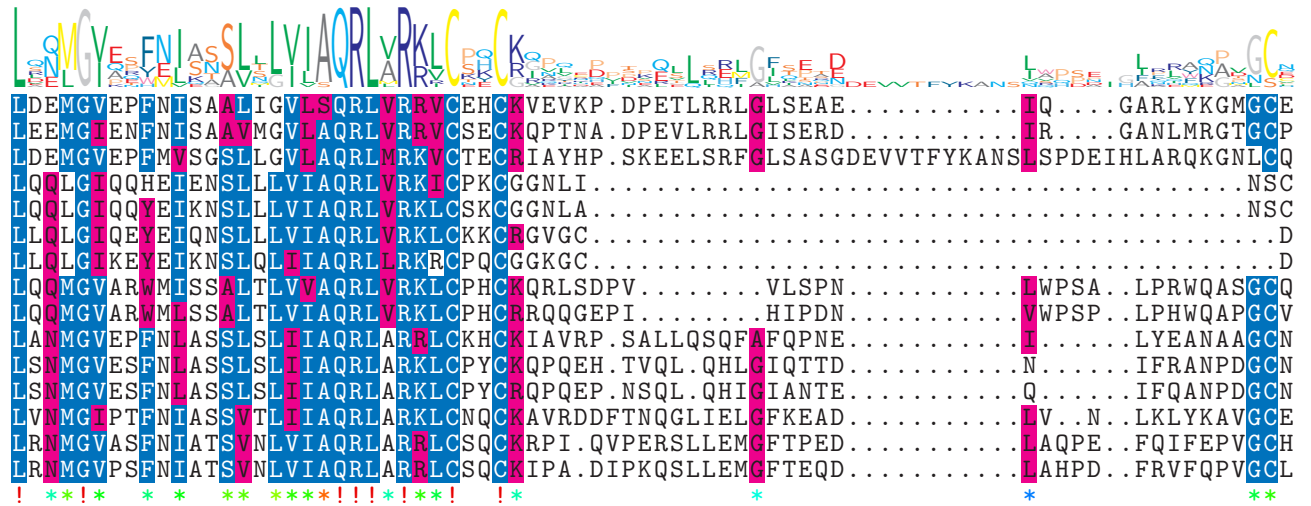
consensus



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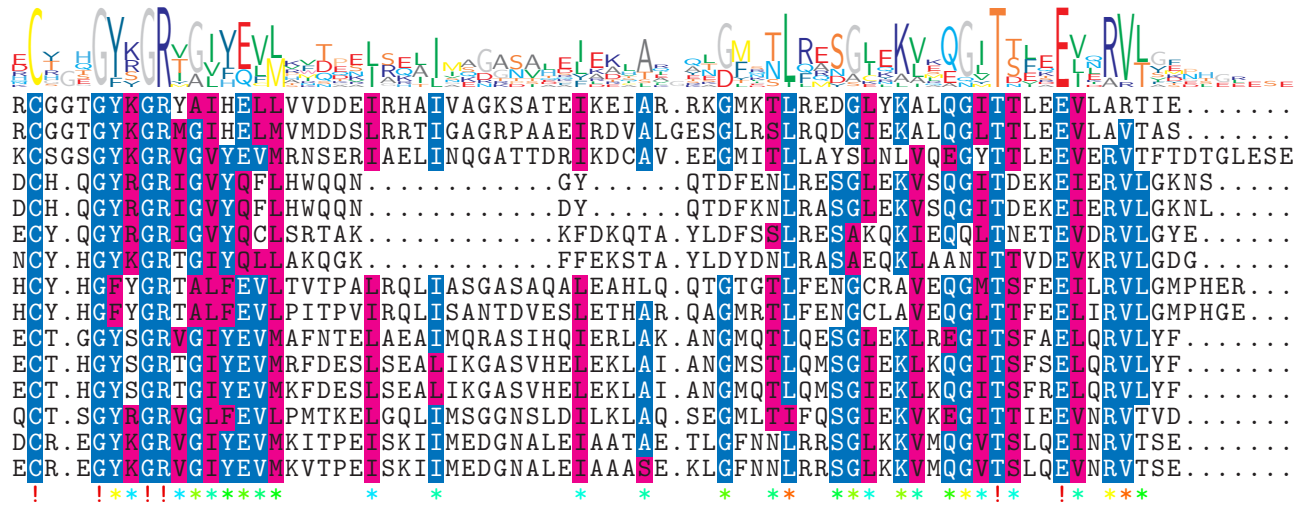
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 consensus



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NTDB id 1028 TT RS08215 WP 011173991.1
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 consensus



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logo

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consensus		

 non conserved
 similar
 ≥ 50% conserved