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NTDB id 1306 DR RS10060 WP 027479822.1	MSXLSIGDRRLGAALLDQGYLGDNDLQRALERHSEVGGRLADVLIIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	79
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
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NTDB id 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPPFALE..GDLRLVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	157
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	SDAGRTGGKLGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFGHPGEEVL	237
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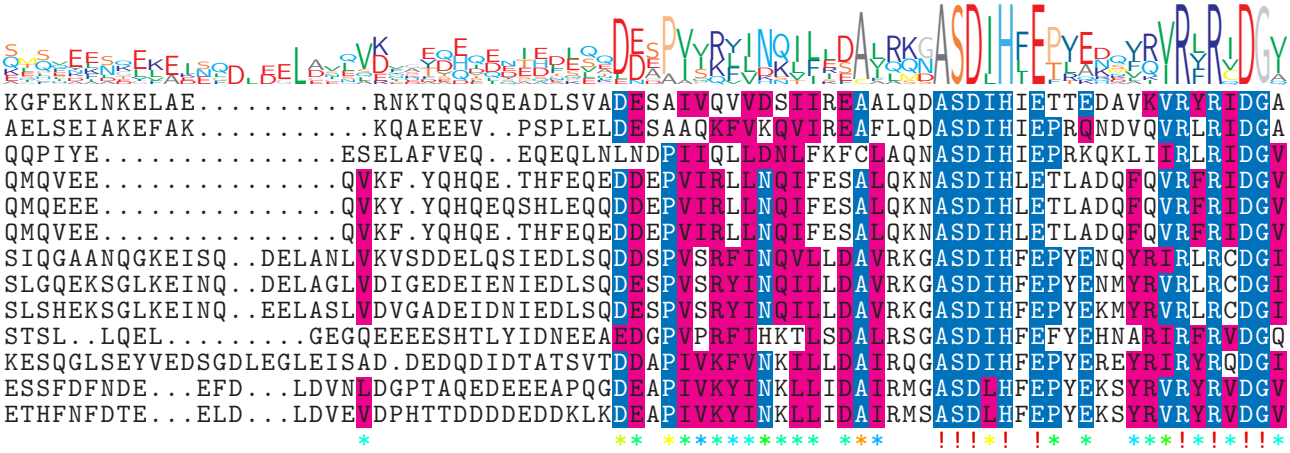
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NTDB	id	1306	DR	RS10060	WP	O27479822.1	NPPDPKVALMIPEATARRYVVVPVRLQGNSLVAMKDPRNVFALDDKLITG....KEILPAVMAEKDIIRLIERFYGE	472
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NTDB	id	1401	DSB67	RS12665	WP	O10643259.1	YEYASLCQTLGLRDLITRHNALPLNRSTLILAVADPTNLQAEDDFRFATG....LQVELVLADFRELTAATRLLYGR	141
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NTDB	id	1058	ABD1	RS01615	WP	O01274986.1	YDTSQIPKDLVDQKLILKHRIPLIQRGQILYVATSNPSTIEAIDAIRFNASK....LLVEPVIVEHHKLEKVLGGHFAE	149
NTDB	id	1018	ACIAD	RS01685	WP	O04920473.1	FDPALFLKDKIDEKLIKQYRIMPLVLRGHVLVYATSNPTNIEAMDRAIRFNASK....LKVEPUIIVEHDKLERLSEHFVE	149
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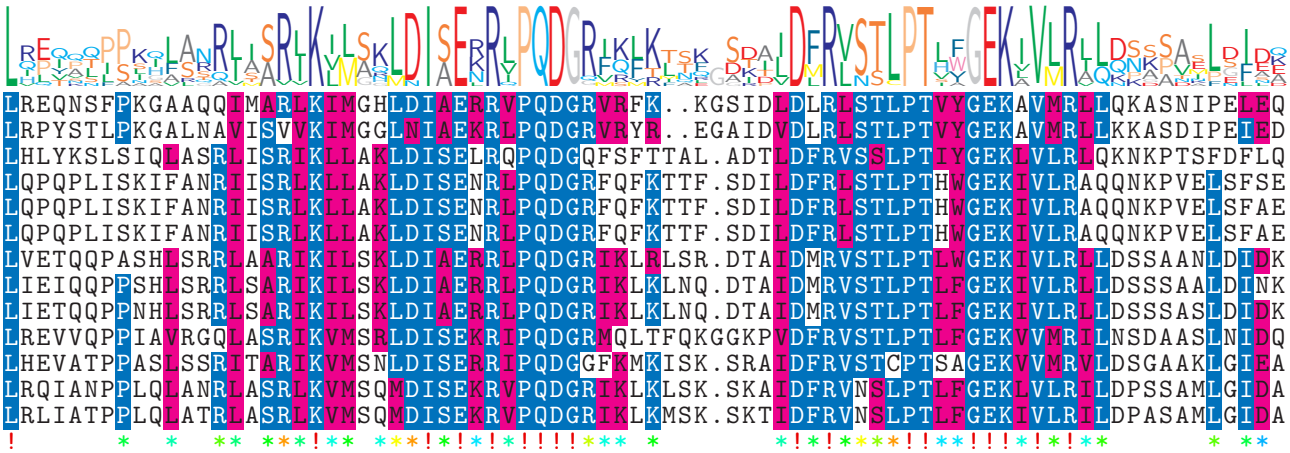
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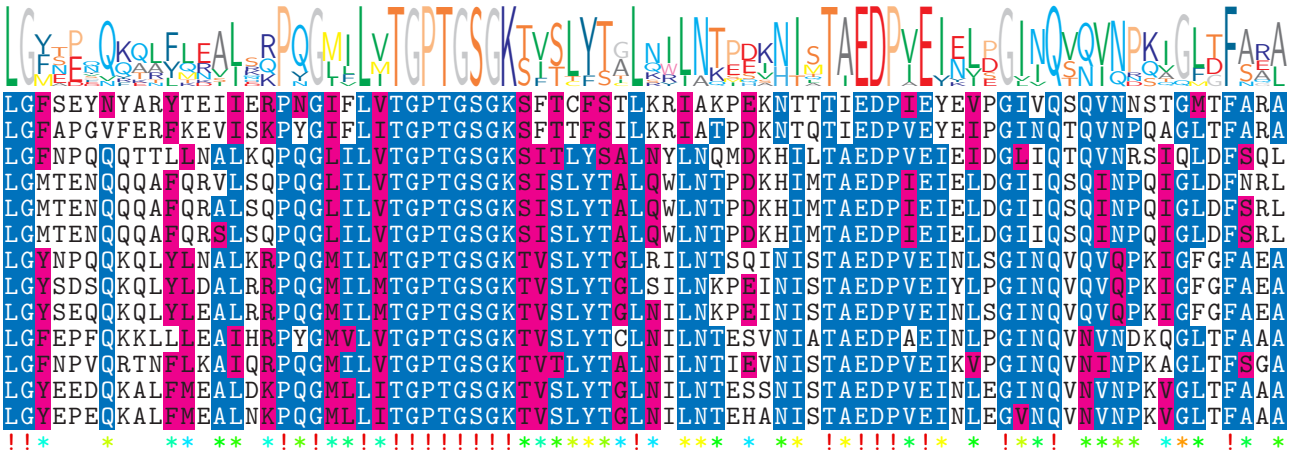
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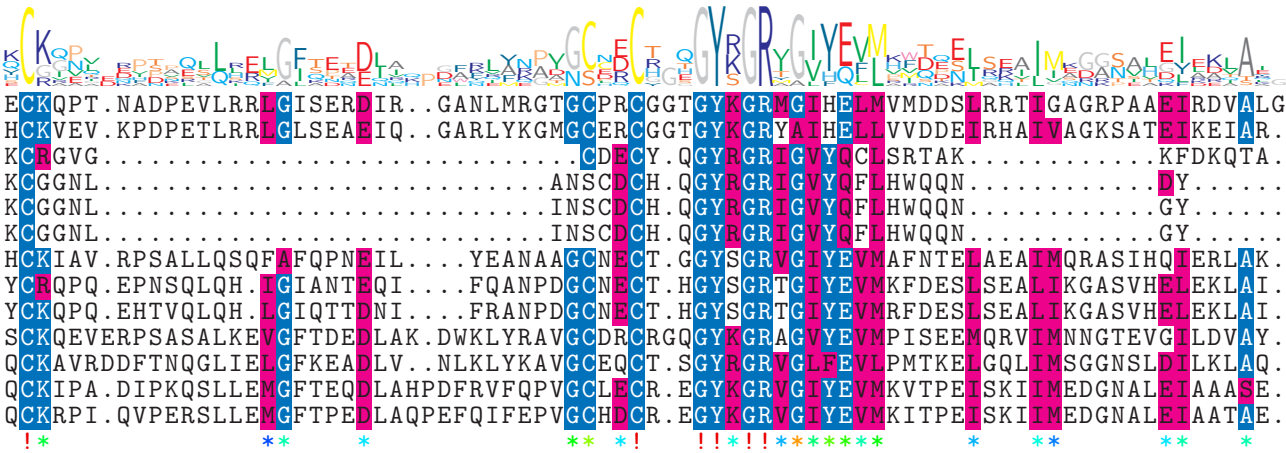
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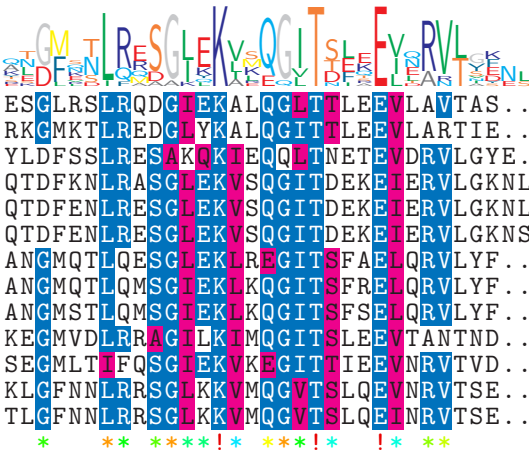
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consensus



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consensus



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