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NTDB id 1306 DR RS10060 WP 027479822.1	LSMLRSDALRYLAVPYBESNREVEVYVSDPRKREERYRRLYRXPXSHMLTSPBYEELHESYYPQERRLGEXLVQEGELSR	317
NTDB id 1028 TT RS08215 WP 011173991.1	GSMLRADALRLLAVPVDETEQQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLLGEQLVQEGTLSR	317
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 1017622 ABX025 RS07575 WP 371751717.1	0
NTDB id 1110 NGFG RS09235 WP 003689817.1	0
NTDB id 1251 GC085 RS07725 WP 011946523.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
consensus	0

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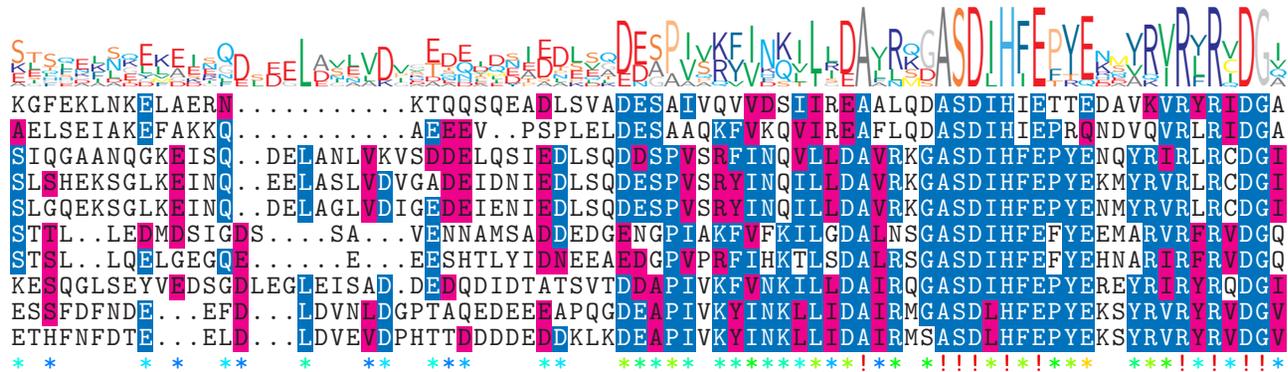
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NTDB id 1028 TT RS08215 WP 011173991.1	EALKEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRGGGRLEDTLVQSGKLRPEALACAVATQLGYPYVDPEE	397
NTDB id 1168 A1552VC RS11070 WP 000957200.1	66
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NTDB id 1251 GC085 RS07725 WP 011946523.1	74
NTDB id 1058 ABD1 RS01615 WP 001274986.1	74
NTDB id 1018 ACIAD RS01685 WP 004920473.1	74
consensus	74

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NTDB id 1306 DR RS10060 WP 027479822.1	NPPDPKVALMIPAEATARRVVPVRLQGNLSLVAMKDPNRFALDDLKLITG.....KEILPAVMAEKDIIRLIERYFGE	472
NTDB id 1028 TT RS08215 WP 011173991.1	DPPDPGAPLLIPEDLCRRYGVFPHRLEGNRLVLLMKDPRNILALDDVRLALKRKGKLNVEVAPAVATEAAITKLIERYFGK	477
NTDB id 1168 A1552VC RS11070 WP 000957200.1	YDYANLCQQQLGLRELITRYDALPIAKQGNLLLLAVSDPTLLQAEFEFRFATG.....LQVELALADHRALQAAIRRLYGR	141
NTDB id 1293 VP RS12240 WP 005479695.1	YDYANLCQQQLGLRELITRHNA LPHRTPSTLLAVADPTNQQAEDDFRFATG.....LQVELVLA DFRELSTAIRRLYGR	141
NTDB id 1401 DSB67 RS12665 WP 010643259.1	YDYANLCQTLGLRDLITRHNA LPLNRTSSTLILAVADPTNIQAEDDFRFATG.....LQVELVLA DFRELTAAIRRLYGR	141
NTDB id 1017622 ABX025 RS07575 WP 371751717.1	YNQNLVPLDLINLKIIMQYICVPLFKRGNRLFVAVSDPTSIQHYQKIATDAG.....CVLNLVIVRDSQLRQLLENLGQE	140
NTDB id 1110 NGFG RS09235 WP 003689817.1	YPRHRVLMGVLTEEQMVEFHCVVFRGDKVFFAVSDPTQMPQIQKTVSAAG.....IAVELVIVEDDQLAGLLDWVGSR	140
NTDB id 1251 GC085 RS07725 WP 011946523.1	IDVGTIPVNLVNEKLIKRRHAMVPLFSRGTNLYLATDDPSKQASLKEIQFHTG.....LNTHAIVVETDKLSALIDNLLTA	149
NTDB id 1058 ABD1 RS01615 WP 001274986.1	YDTSQIPKDLVDQKLIKHRILPLIQRGQILYVATSNPSNIEAIDAIRFNSK.....LLVEPVIVEHHKLEKVLGQHFAE	149
NTDB id 1018 ACIAD RS01685 WP 004920473.1	FDPALFLKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTNIEAMD AIRFNSK.....LKVEPILIVEHDKLERLISEHFVE	149
consensus	149

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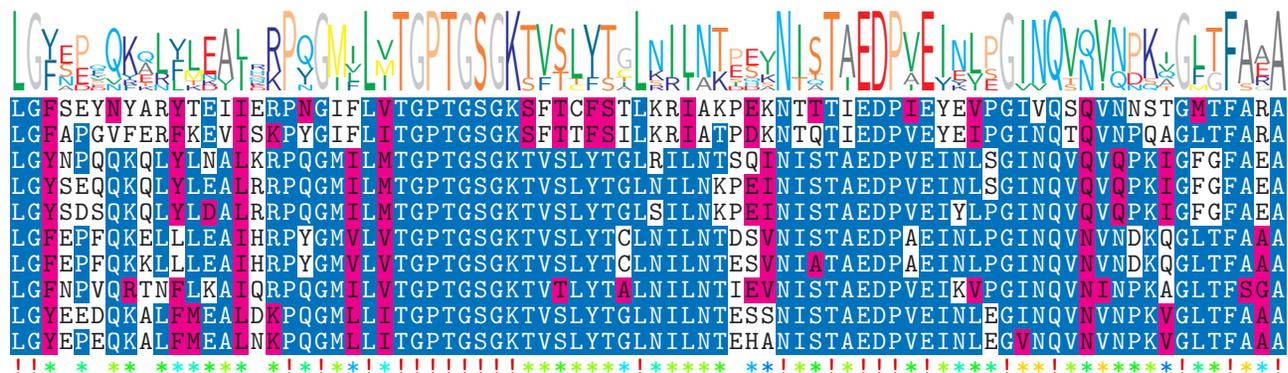
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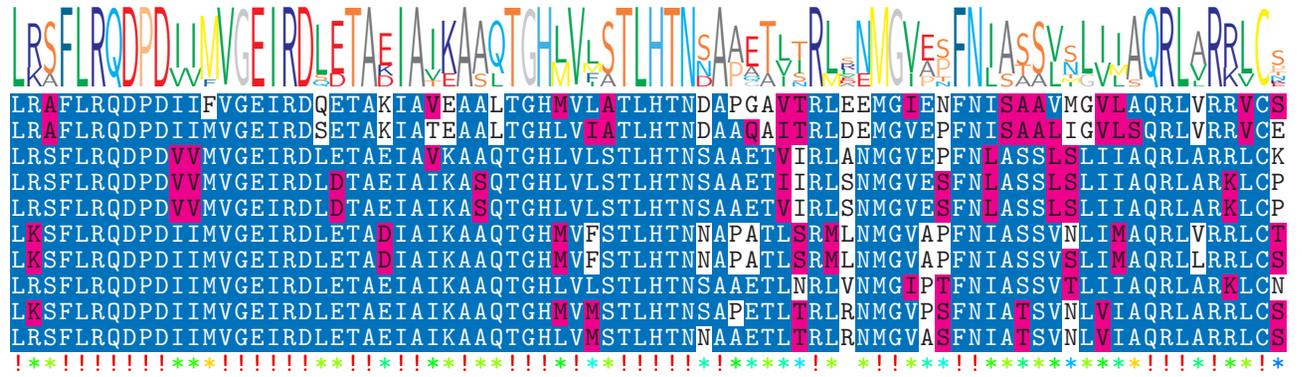
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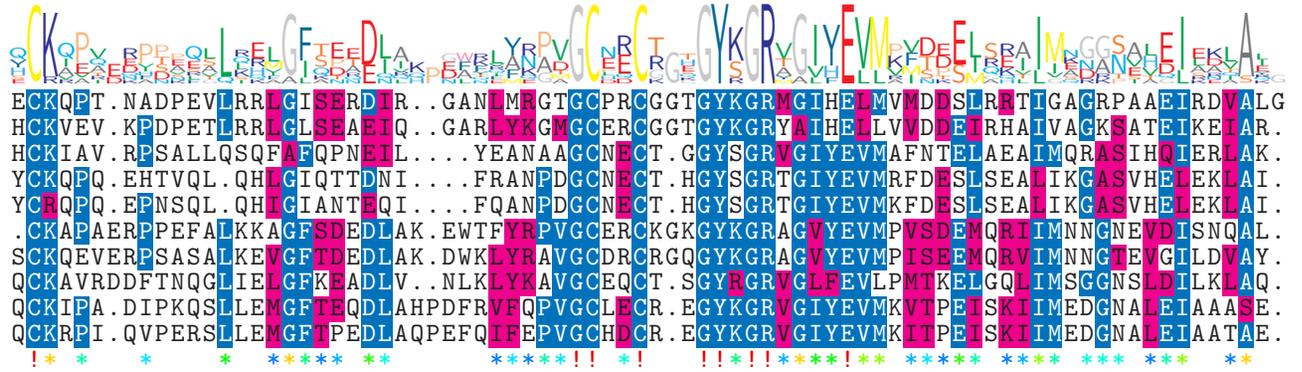
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 consensus



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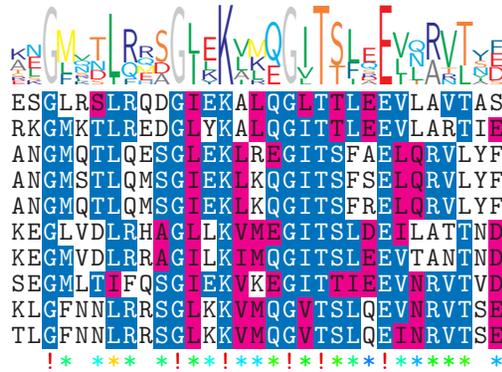
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 consensus



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 consensus



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- non conserved
- similar
- ≥ 50% conserved