

logo

NTDB id 1306 DR RS10060 WP 027479822.1	MSXLRIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAL	79
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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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
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NTDB id 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIVVPSGA	157
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NTDB id 1306 DR RS10060 WP 027479822.1	SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEEQEGTTFLPNPSGFGHPGEEVL	237
NTDB id 1028 TT RS08215 WP 011173991.1	SGEGQKDKLKGELLQKQWISREALEEALVEQEKTGDLGRILV.RKGLPEEALYRALAEQKGLFLESTEGIVPDPSAA	237
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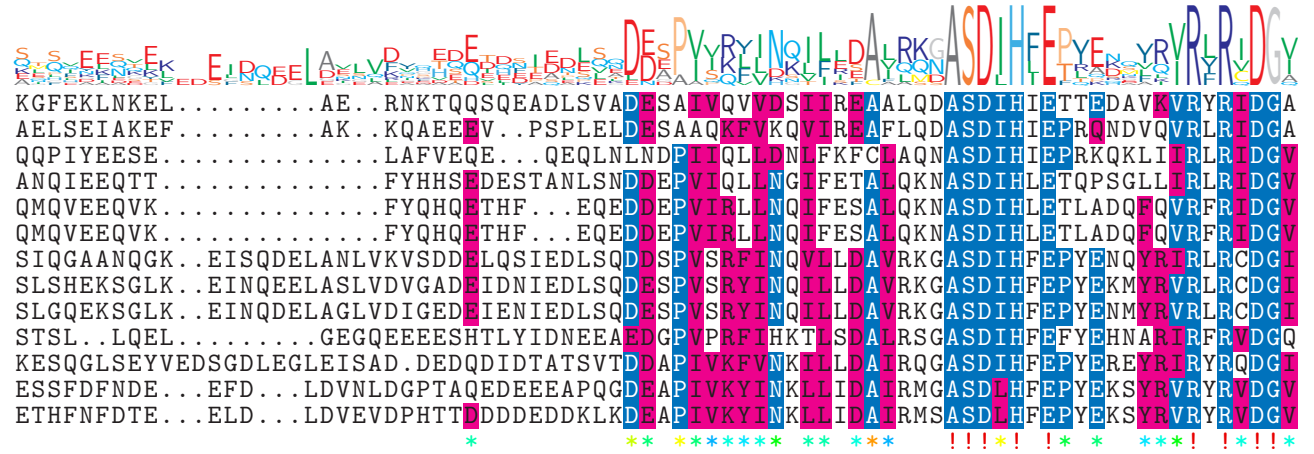
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NTDB id 1018 ACIAD RS01685 WP 004920473.1MSAFTTPPKFSGFIRRLVVEEGYVNAQNMQQALEKAKKFKQDIVPYLIDNFSISPLTIAEII SLEFGEPLLDLGV	74
consensus*.....**.....*.....*.....*.....**	

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NTDB id 1384 A4U84 RS08525 WP 010787006.1	ISPSRWQKNCEEKQLLLRYLAVPVQETEHKLWLAVD DENNLTA CEIFAFMTH.....KQIEPVVITASDELKYL NALSPE	91
NTDB id 42225 PARA RS00540 WP 014064061.1	ISPQYWQQNQQQQALLLRYFALPLKEDEHHLWLAVD SLN NLAACETFAFLSG.....KLVEPILFETAELKQLLQSLAPK	89
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NTDB id 1355 HI 0298 AAC21962.1	ISPDLWERNQQQSSLLLRYFALPLKEENNRLLWLGVD SL SNLSACETIAFITG.....KPVEPILLESSQLKELLQQLTPN	96
NTDB id 1168 A1552VC RS11070 WP 000957200.1	YDYANLCQQLGLRELITRYDALPIAKGNL LLLAVSDPTLLQAE EEFRAF TG.....LQVELALADHRAIQAAIRRLYGR	141
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	FDPALFLKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTNIEAMD A I R F N S K.....LLVEPITIVEHDKLERLLSEHFVE	149
consensus*.....**.....**.....*!.....*****.....**.....*.....*.....*.....*.....*.....*	

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 consensus



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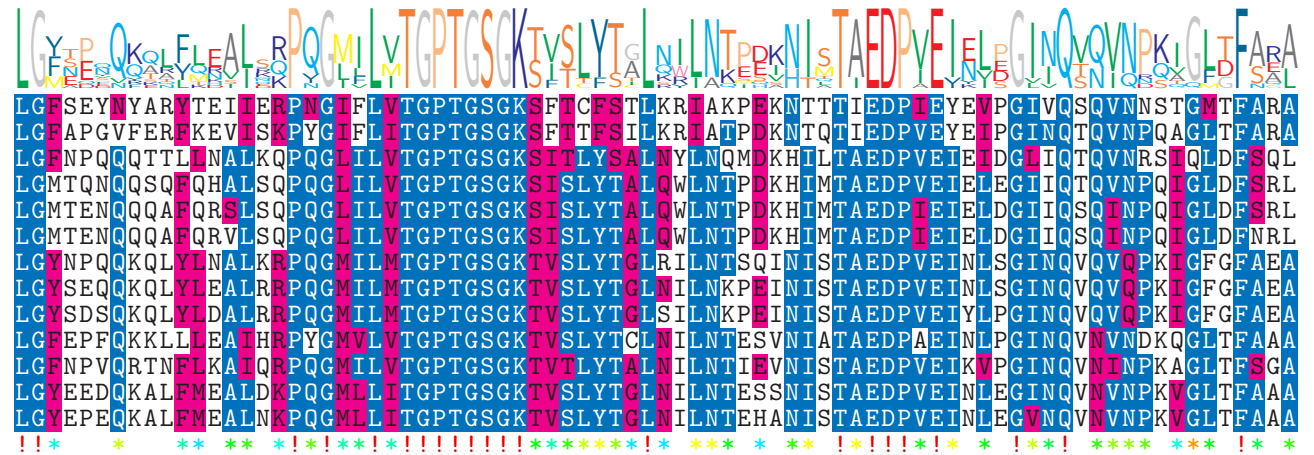
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 consensus



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 consensus



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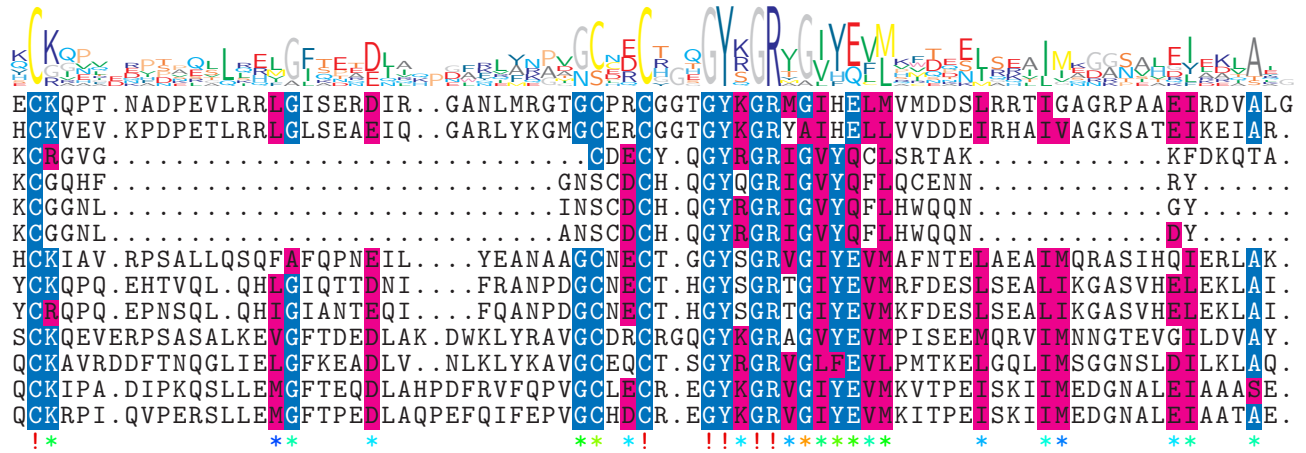
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 consensus



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 consensus



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 consensus



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570

 non conserved
 similar
 $\geq 50\%$ conserved