

logo

NTDB id 1306 DR RS10060 WP 027479822.1	MSXLRIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	79
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1384 A4U84 RS08525 WP 010787006.1	0
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	0
NTDB id 1293 VP RS12240 WP 005479695.1	0
NTDB id 1110 NGFG RS09235 WP 003689817.1	0
NTDB id 1251 GC085 RS07725 WP 011946523.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
consensus	0

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NTDB id 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIVVPSGA	157
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consensus	0

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NTDB id 1306 DR RS10060 WP 027479822.1	SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFFHPGEEVL	237
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consensus	0

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NTDB id 1306 DR RS10060 WP 027479822.1	LSMLRSDALRYLAVPYGSEVVEVYVSDPRKREYRXLNRPXSHMLXEPBYEELHESYYPQRNRLGEXLVQEGELSR	317
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NTDB id 1058 ABD1 RS01615 WP 001274986.1MSALHTSPKFTGFFRRLVEEKHVSAAATMQTALDAAKRAKQDTVAYLIEEVHLSPSLLAETISAFAEPYFDLDV	74
consensus*.....*	

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NTDB id 70936 K6J57 RS03935 WP 011271972.1	ISPDLWERNQQQSSLLRLRYFALPLKEENNRLWLGVDLSNLSACETIAFITG.....KPVEPILLESSQLKELLQQLTPH	96
NTDB id 1366 NTHI RS01950 WP 011271972.1	ISPDLWERNQQQSSLLRLRYFALPLKEENNRLWLGVDLSNLSACETIAFITG.....KPVEPILLESSQLKELLQQLTPH	96
NTDB id 1355 HI 0298 AAC21962.1	ISPDLWERNQQQSSLLRLRYFALPLKEENNRLWLGVDLSNLSACETIAFITG.....KPVEPILLESSQLKELLQQLTPN	96
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NTDB id 1251 GC085 RS07725 WP 011946523.1	IDVGTIPVNLVNEKLIKRRHAMVPLFRSGTNLYLATDPPSKQASLKEITQFHTG.....LNTHAIVVETDKLSALIDNLLTA	149
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	YDTSQIPKDLVDQKLLILKHRIIPLIQRGQILYVATSNPSNIEAIDAIRFNSK.....LLVEPVIVEHHKLEKVLGQHFAE	149
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 consensus



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logo

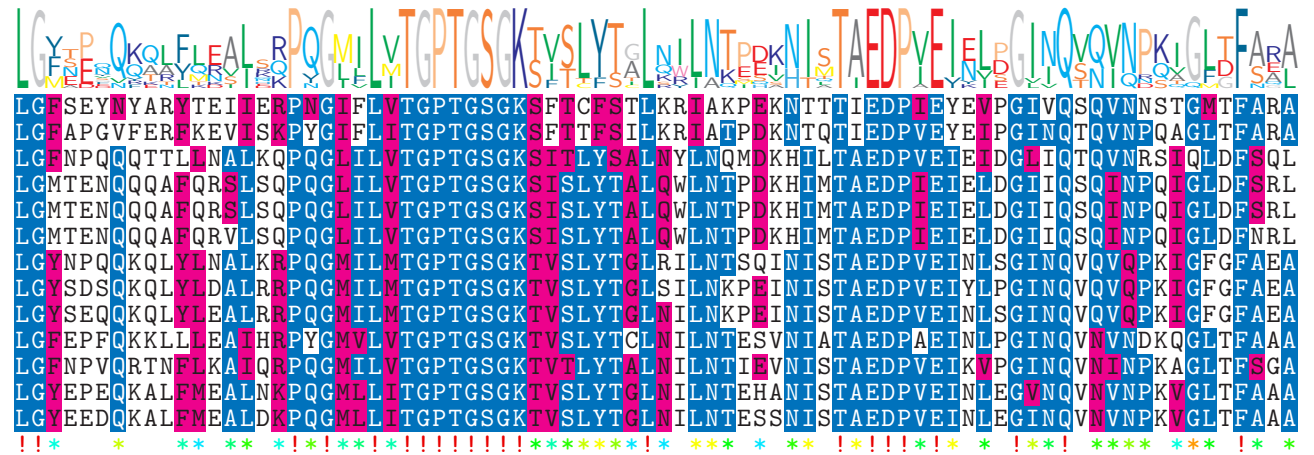
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 consensus



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 consensus



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logo

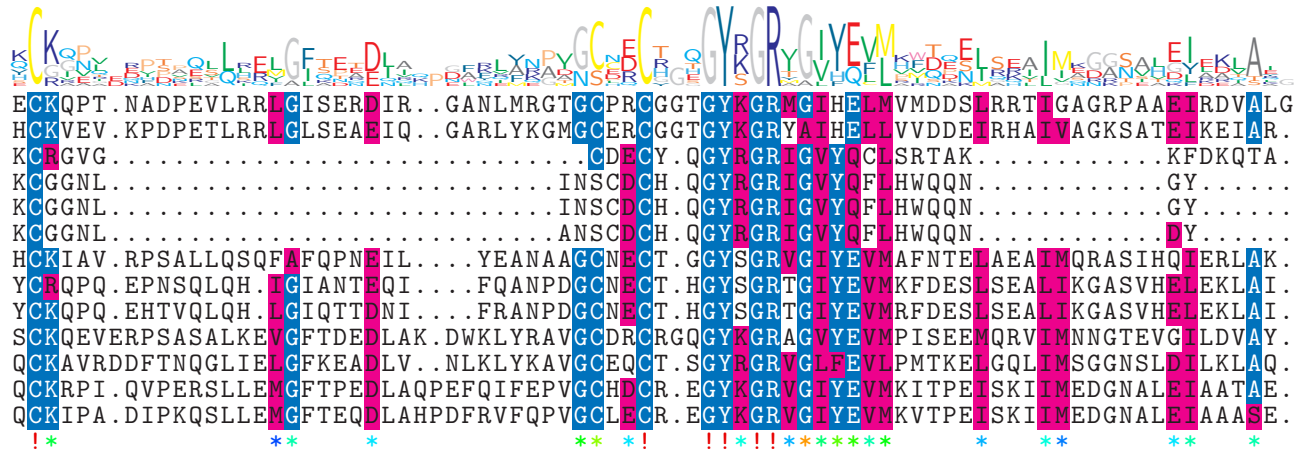
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 consensus



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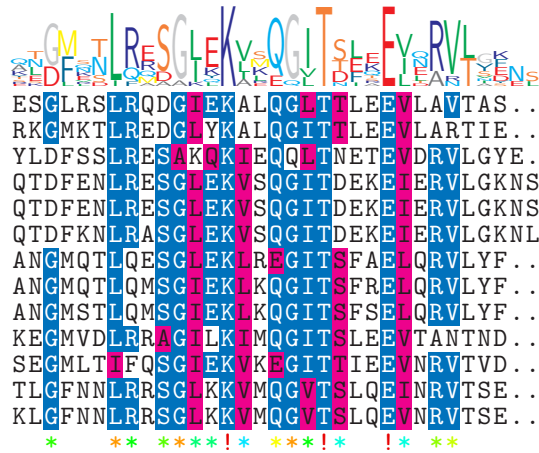
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 consensus



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NTDB id 1306 DR RS10060 WP 027479822.1
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 consensus



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561
561
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574
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570

 non conserved
 similar
 $\geq 50\%$ conserved