

logo

NTDB id 1306 DR RS10060 WP 027479822.1	MSXLSLIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAL	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 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIVVPSGA	157
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLTSLLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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NTDB id 1028 TT RS08215 WP 011173991.1	SGEGQKDKLKGELLQKQWISREALEEALVEQEKTDGDLGRILV.RKGLPEEALYRALAEQKGLFLESTEGIVPDPSAA	237
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NTDB id 1028 TT RS08215 WP 011173991.1	LLLLRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSR	317
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NTDB id 1293 VP RS12240 WP 005479695.1	.....MHSNLSTILRQKGLLTFQSQESLIEQVKASGISMPEALLSSGFFTSSELAEHLSSIFGLSQPELSQ	66
NTDB id 1110 NGFG RS09235 WP 003689817.1	.....MSVGLLRILVQNQVVTVEQAEHYY.NESQAGKEVLPMLFSDGVTISPKSLAALIAARVFSYSILDLRH	65
NTDB id 1251 GC085 RS07725 WP 011946523.1	.....MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEEKISLLQYIVKNKILSAEQIALTAAQNFVPMPLDINC	74
NTDB id 1018 ACIAD RS01685 WP 004920473.1	.....MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQALEKAKKFKQDIVPYLIDNFSISPLTIAEITISLEFCEPLLDLGV	74
NTDB id 1058 ABD1 RS01615 WP 001274986.1	.....MSALHTSPKFTGFFRRLVEEKHVSAAATMQTALDAAKRAKQDTVAYLIEEVHLSPSLLAETISAFAEPYFDLDV	74
consensus	.....*.....**.....*	

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NTDB id 1028 TT RS08215 WP 011173991.1	DPPDPGAPLILPEDLCRRYGVFPHRLEGNRLVLLMKDPRNILALDDVRLALKRKLNYEVAPAVATEAAITKLIERYFYGK	477
NTDB id 1384 A4U84 RS08525 WP 010787006.1	ISPSRWQKNCEEKQLLLRYLAVPVQETEHKLWLAVDDENNLTACEIFAFMTH.....KQIEPVVITASDELKYLINALSPE	91
NTDB id 71395 K6J71 RS03985 WP 005666685.1	ISPDLWERNQQQSSLLRLRYFALPLKEENNRLWLGVDLSNLASACETIAFITG.....KPVEPILLESSQLKELLQKLTTPC	96
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NTDB id 1293 VP RS12240 WP 005479695.1	YEYASLCQQIGLRELITRHNAIPLHRTPTSTLLAVADPTNQAEDDFRFATG.....LQVELVLADFRELSTAIRRLYGR	141
NTDB id 1110 NGFG RS09235 WP 003689817.1	YPRHRVLMGVLTEEQMVVEFHCVPVFRRGDKVFAVSDPTQMPQIQKTVSAAG.....IAVELVIVEDDQLAGLLDWVGRS	140
NTDB id 1251 GC085 RS07725 WP 011946523.1	IDVGTIPVNLVNEKLIKRRHAMVPLFRSGTNLYLATDPPSKQASLKEITQFHTG.....LNTHAIVVETDKLSALIDNLLTA	149
NTDB id 1018 ACIAD RS01685 WP 004920473.1	FDPALFLDKIDEKLIKRYRIMPLVHRGHVLYVATSNPTNIEAMDAIRFNSK.....LKVEPIVIVEHDKLERLLSEHFVE	149
NTDB id 1058 ABD1 RS01615 WP 001274986.1	YDTSQIPKDLVDQKLLILKHRIPLIQRGQILYVATSNPSNIEAIDAIRFNSK.....LLVEPVIVEHHKLEKVLGQHFAE	149
consensus	.....*.....**.....*!*.....*****.....*.....*	

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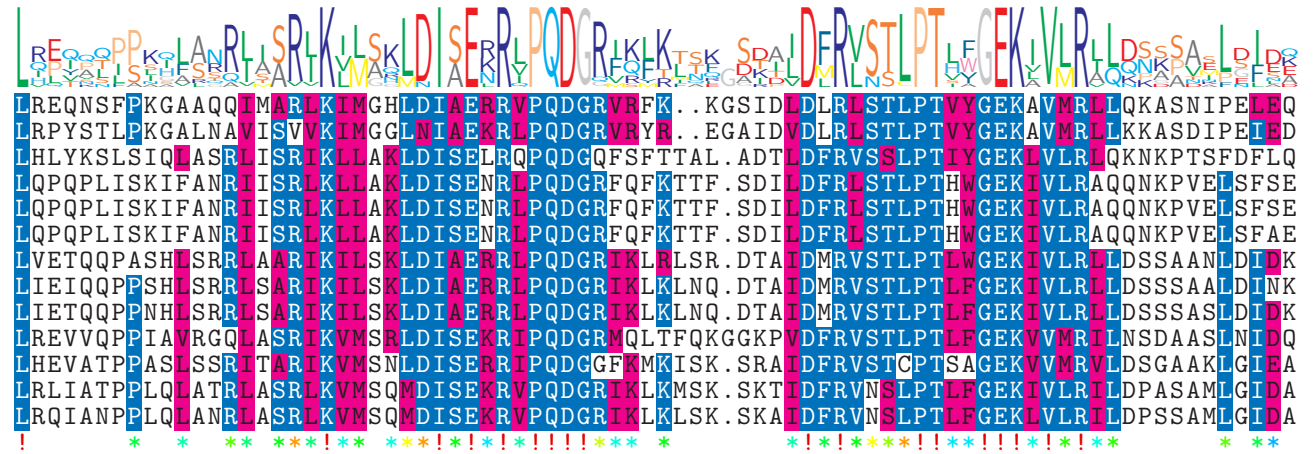
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 consensus



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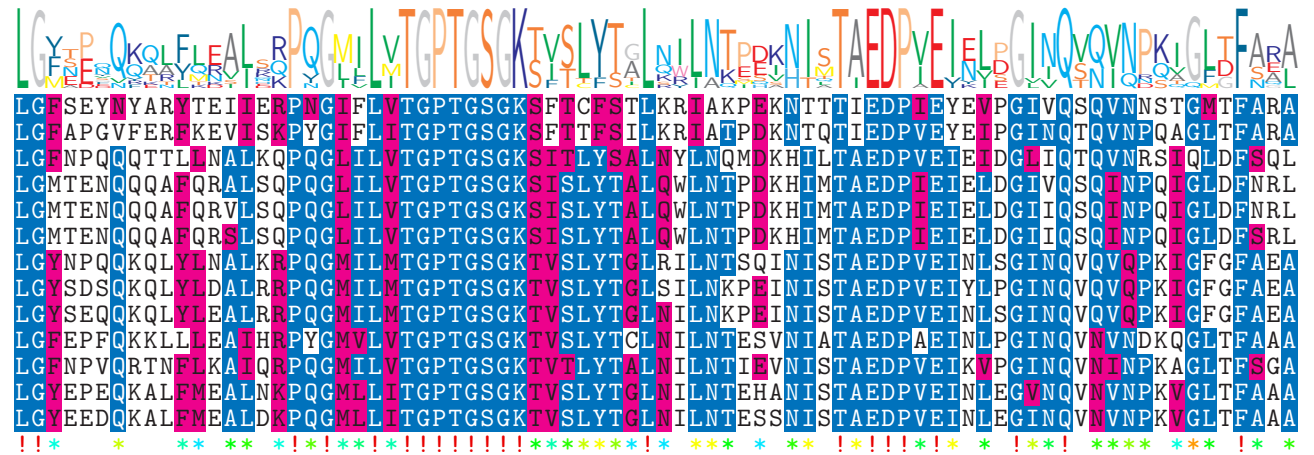
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 consensus



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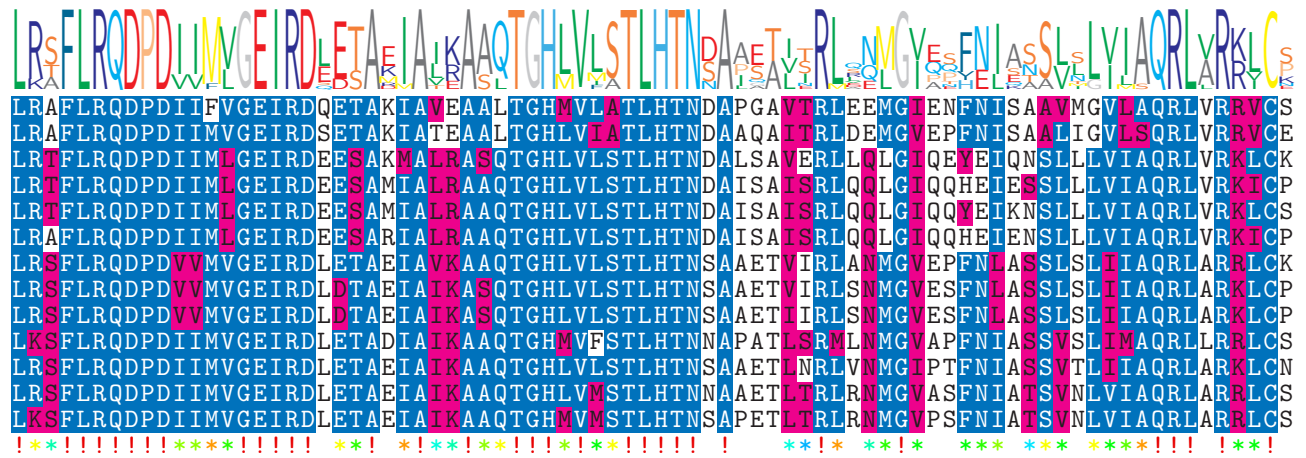
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 consensus



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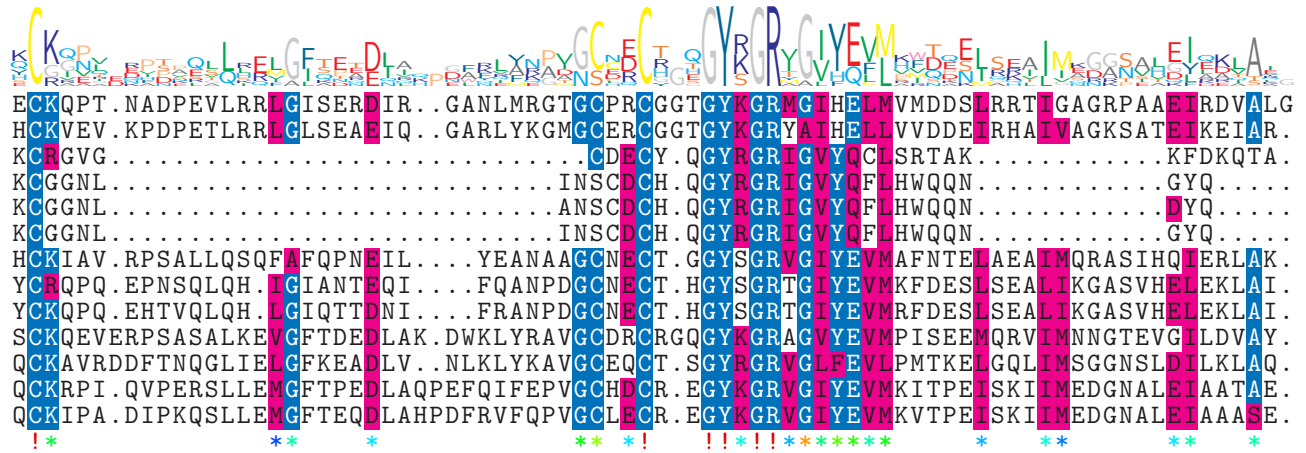
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NTDB id 1306 DR RS10060 WP 027479822.1  
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 NTDB id 1384 A4U84 RS08525 WP 010787006.1  
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 consensus



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 NTDB id 1384 A4U84 RS08525 WP 010787006.1  
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 consensus



887  
889  
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 non conserved  
 similar  
  $\geq 50\%$  conserved