

logo

NTDB id 1306 DR RS10060 WP 027479822.1	MSXLRIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAL	79
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAILLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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NTDB id 1168 A1552VC RS11070 WP 000957200.1	0
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	0
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consensus	0

logo

NTDB id 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIVVPSGA	157
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NTDB id 1306 DR RS10060 WP 027479822.1	SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
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NTDB id 1306 DR RS10060 WP 027479822.1	LEMLRSDALRYLAVPYGSEVVEVYVSDPRKREERYRXLNRPXSMLEPBYEELHESYYPQRNRLGELVQEGELSR	317
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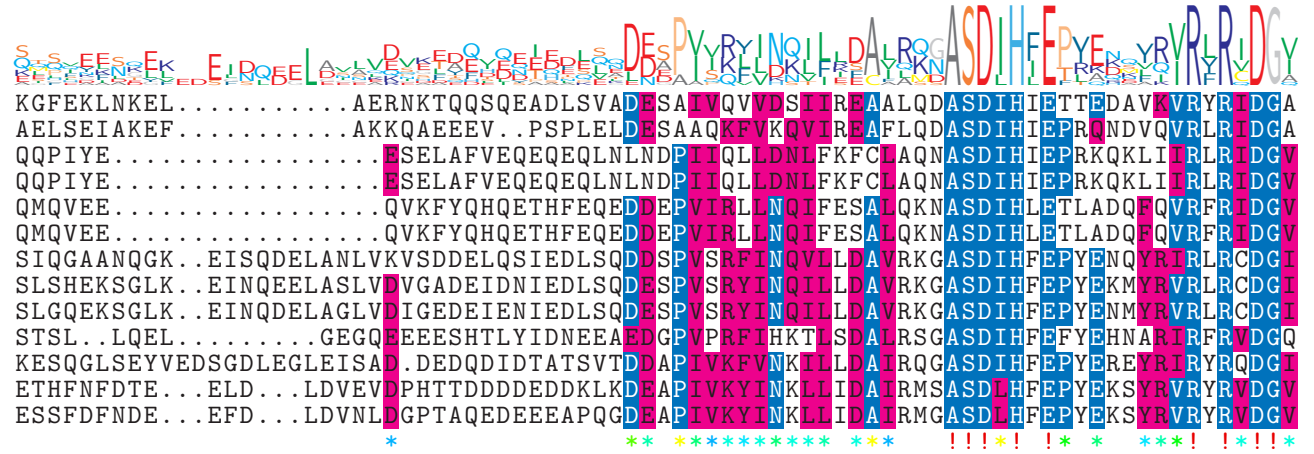
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NTDB id 1251 GC085 RS07725 WP 011946523.1MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEEKISLLQYIVKNKILSAEQIALTAQNFVGPMLDINC	74
NTDB id 1018 ACIAD RS01685 WP 004920473.1MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQALEKAKKFKQDIVPYLIDNFSISPLTIAEITISLEFCEPLLDLGV	74
NTDB id 1058 ABD1 RS01615 WP 001274986.1MSALHTSPKFTGFFRRLVEEKHVSAAATMQTALDAAKRAKQDTVAYLIEEVHLSPSLLAETISAFAEPYFDLDV	74
consensus*.....**.....*.....*.....*.....***.....*	

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NTDB id 290748 C3363 RS00935 WP 010787006.1	ISPSRWQKNCEEKQLLLRYLAVPVQETEHKLWLAVDENNLTACEIFAFMTH.....KQIEPVVIASDELKYLINALSPE	91
NTDB id 1384 A4U84 RS08525 WP 010787006.1	ISPSRWQKNCEEKQLLLRYLAVPVQETEHKLWLAVDENNLTACEIFAFMTH.....KQIEPVVIASDELKYLINALSPE	91
NTDB id 1366 NTHI RS01950 WP 011271972.1	ISPDLWERNQQQSSLLRYFALPLKEENNRLWLGVDLSNLSACETIAFITG.....KPVEPILLESSQLKELLQQLTPH	96
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	YDTSQIPKDLVDQKLLILKHRIIPLIQRGQILYVATSNPSNIEAIDAIRFNSK.....LLVEPVIVEHHKLEKVLGQHFAE	149
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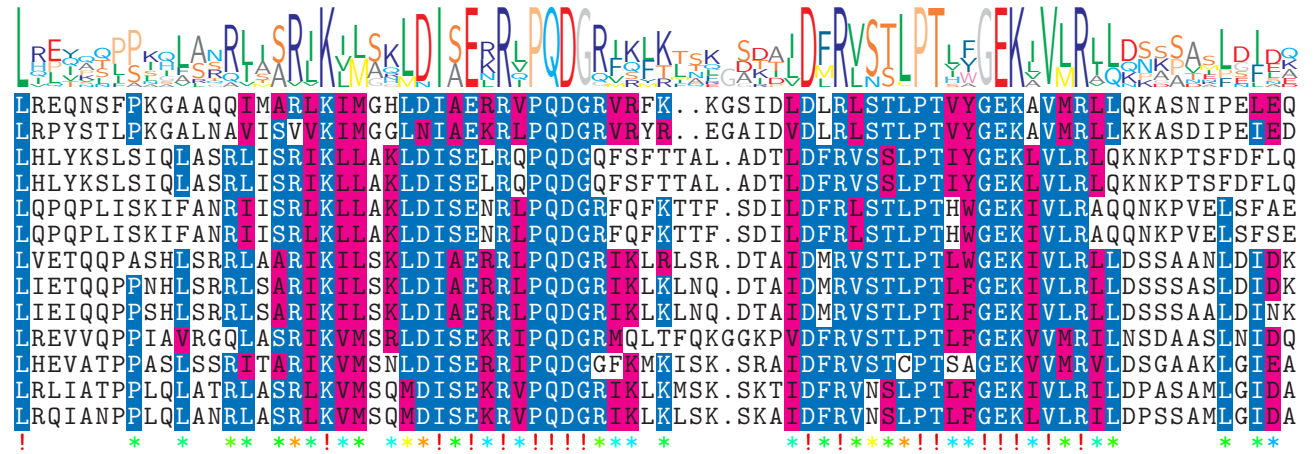
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 consensus



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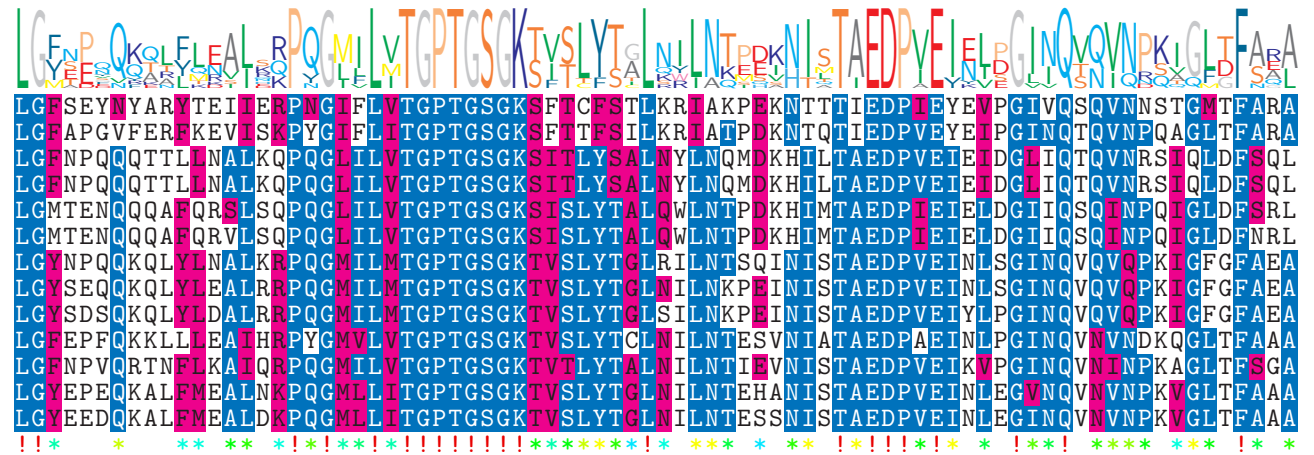
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 consensus



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 consensus



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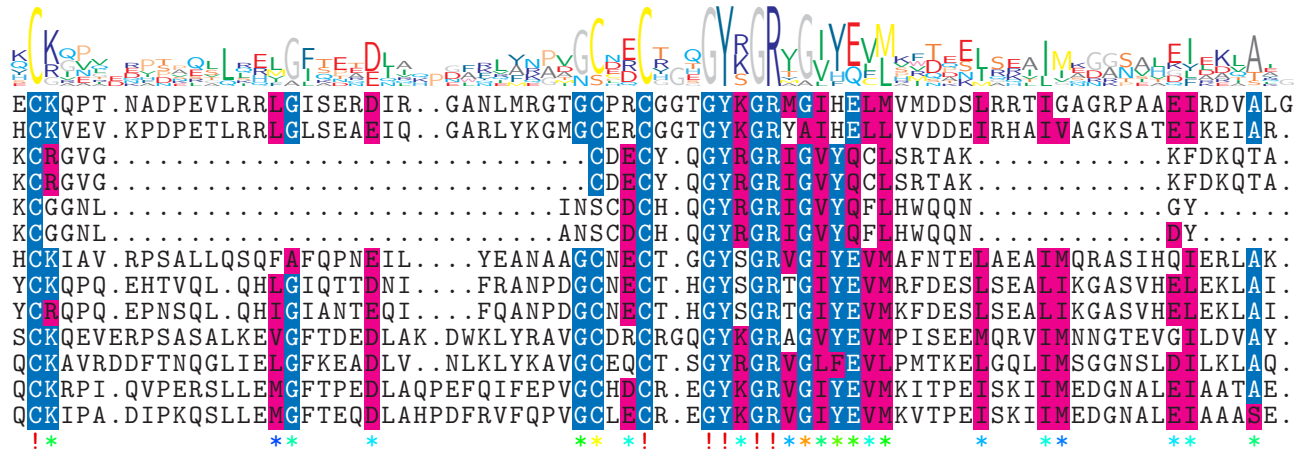
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 consensus



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 NTDB id 1058 ABD1 RS01615 WP 001274986.1
 consensus



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 consensus



887
889
461
461
464
464
562
561
561
558
574
570
570

 non conserved
 similar
 $\geq 50\%$ conserved