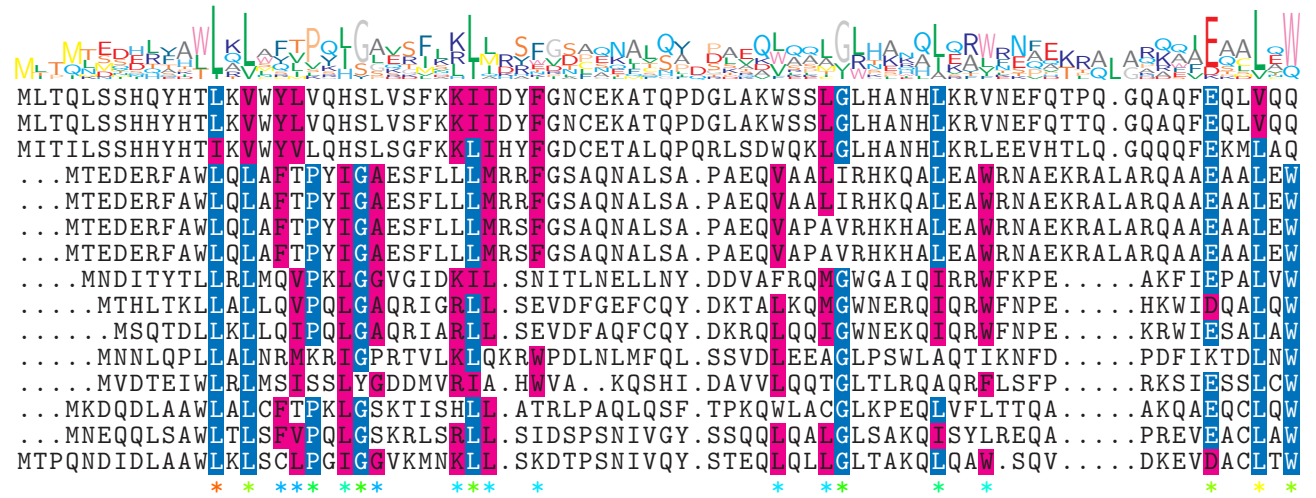


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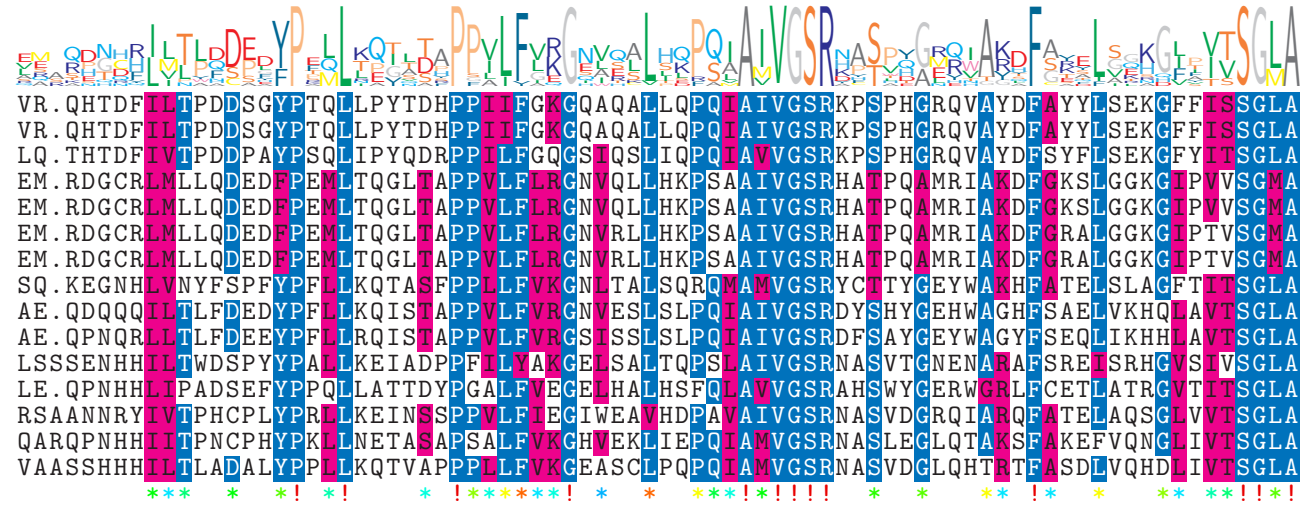
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 NTDB id 1073 ABD1 RS00865 WP 015451369.1  
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 NTDB id 1094 KZH42 RS01320 WP 002224767.1  
 NTDB id 1089 NMB RS00600 WP 002224767.1  
 NTDB id 1122 OK783 RS09595 WP 003705341.1  
 NTDB id 1120 NGFG RS10355 WP 020997408.1  
 NTDB id 1361 HI 0985 AAC22646.1  
 NTDB id 1388 A4U84 RS04945 WP 062924099.1  
 NTDB id 1466 ASU2 RS09750 WP 015674213.1  
 NTDB id 1250 GC085 RS13065 WP 027219989.1  
 NTDB id 1445 BW25113 RS17035 WP 000228551.1  
 NTDB id 1152 A1552VC RS00200 WP 000654772.1  
 NTDB id 56728 OCV56 RS15905 WP 086712415.1  
 NTDB id 1399 DSB67 RS15600 WP 010645755.1  
 consensus



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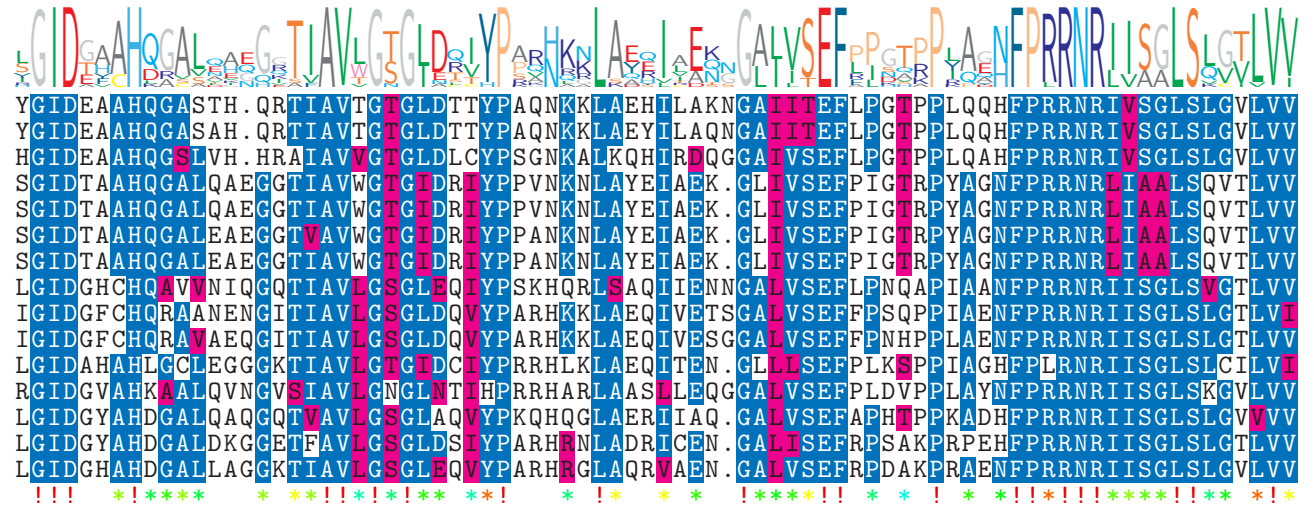
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 NTDB id 1399 DSB67 RS15600 WP 010645755.1  
 consensus



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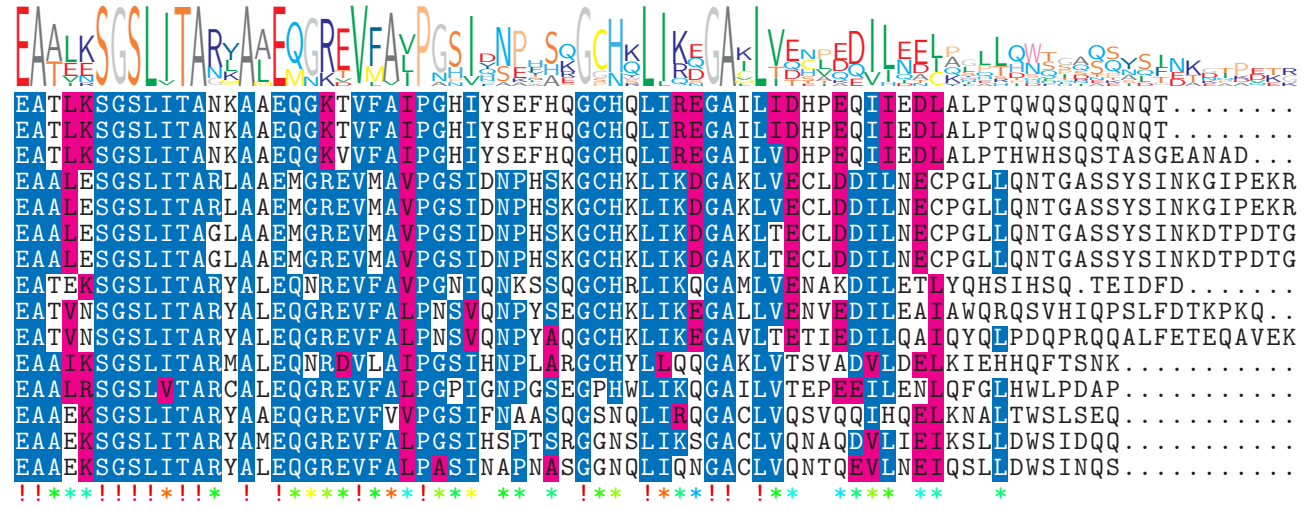
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 NTDB id 1399 DSB67 RS15600 WP 010645755.1  
 consensus



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NTDB id 1043 HON27 RS16615 WP 168727007.1  
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 consensus



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NTDB id 1043 HON27 RS16615 WP 168727007.1  
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 consensus

Sequence	Length
RS.....	376
RS.....	376
RSPY.....	383
IRT.....	397
IRT.....	397
IRT.....	395
IRT.....	395
.....	373
.....	375
.....	384
EYER.....	361
RRACHVRRNTNVFV	374
GRG.....	371
GRG.....	370
GRG.....	369

non conserved  
 similar  
 ≥ 50% conserved