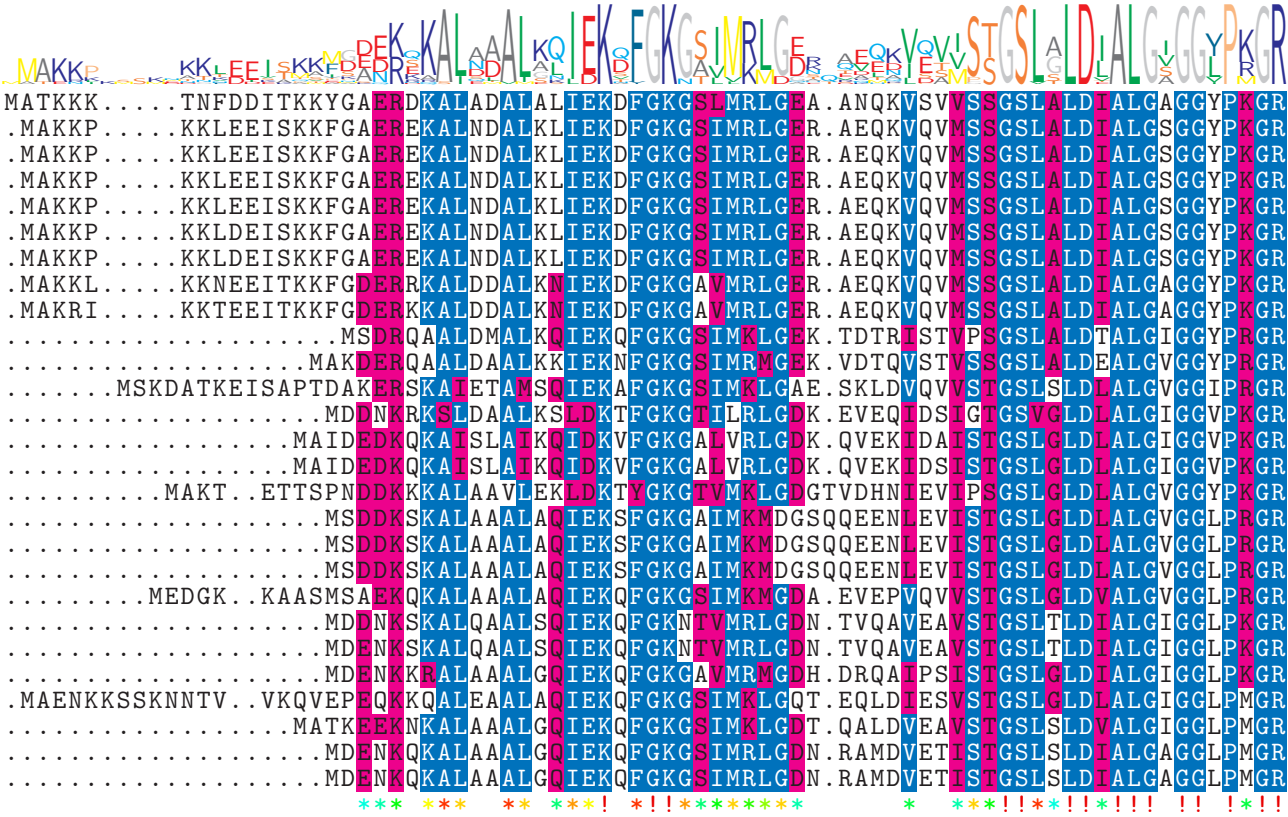


logo

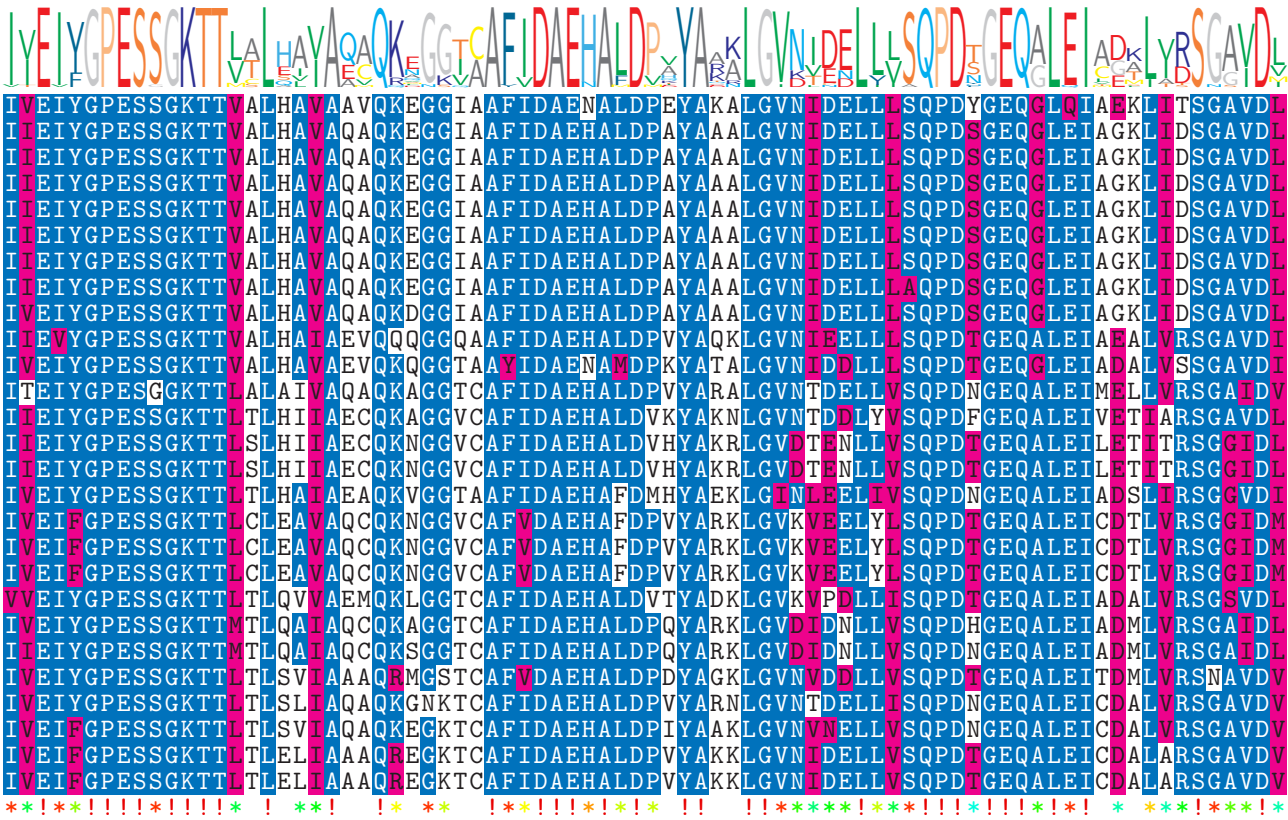
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NTDB id 222270 A6J57 RS05600 WP 005724816.1
NTDB id 1165 A1552VC RS01565 WP 000344154.1
NTDB id 1142 GTF74 RS11550 WP 000344154.1
consensus



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consensus



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NTDB id 1142 GTF74 RS11550 WP 000344154.1
consensus

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Sequence alignment

DEKIGQGGKEAKKYLYLKENPEIAEDDKYREGLDLDSEVAETEEETTASKTAKAKKEEKAVETEEIELELED... 387
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DEKIGQGSNAKKYLAEHPEIFDEIDKQVRSKFGLIDGEEVSEQDTENKKDE...PKKEEA..VNNEEVLPLDLGDELEIE 385
DEKIGQGSNAKKYLAEHPEIFDEIDKQVRSKFGLIDGEEVSEQDTENKKDE...PKKEEA..VNNEEVLPLDLGDELEIE 385
DEKIGQGSNAKKYLADNPEVFAEIDHQVRVQFGLIDGEEAAE...SKKDE...VAQVDS..VNNEEVTLDLGDELEIE 381
DEKIGQGSNAKKYLADNPEIFDEIDHQVRVQFGLIDGEEIAE...SKKDE...AAQVDS..VNNEEVTLDLGDELEIE 381
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GEKIGQGSNAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEEKSSKEK.....ETDEK..ADKNLVLELDDTIELE 382
EERLGGRENAAKQFLKENKDIMLMIQEQIREHYGLDNNGVVQQQAETQEEL.EFE.E..... 348
SERIGQGRENAKNYLADHDVEDEVRLKVRAAYGISDVPEEDLPPTTEDE.QI.NILPDDST.....EE..... 355
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DKKLGGGRENAKALLKEDKALADEITLKIKESIGSNEEI....MPLPDEPLEEME..... 347
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NTDB id 1142 GTF74 RS11550 WP 000344154.1	...	354	
consensus			

-  non conserved
-  similar
-  ≥ 50% conserved