

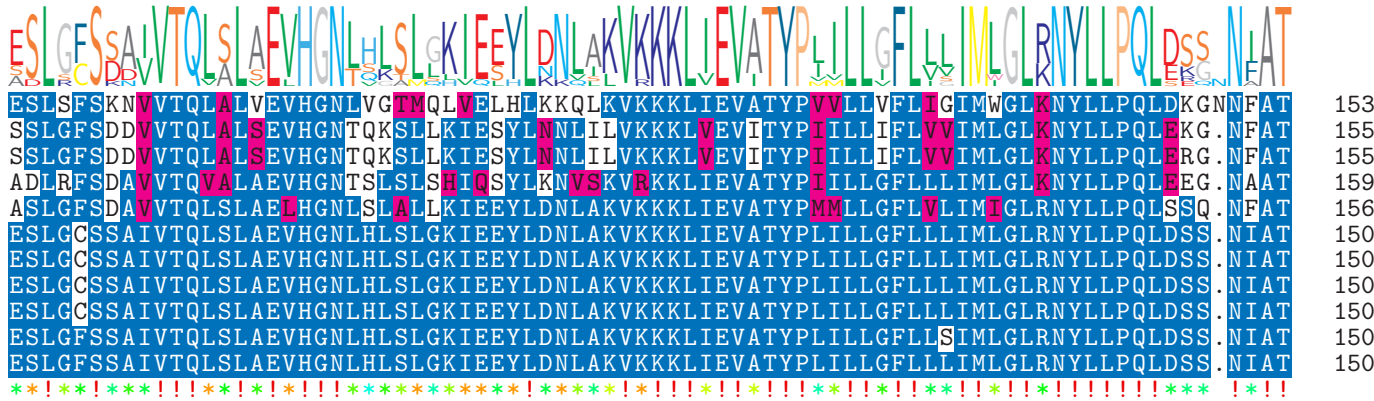
logo

NTDB id 587 KW2 RS10850 WP 042211510.1	MQT
NTDB id 395 FSA28 RS09400 WP 257638827.1	MTFLQQ
NTDB id 386 SMU RS09030 WP 255262721.1	MTFLQQ
NTDB id 1129581 GPW69 RS00750 WP 269472402.1	MRKLI AFLQQ
NTDB id 430 SGO RS09415 WP 012130941.1	...MISFLQQ
NTDB id 271 KZH43 RS09260 WP 074196785.1	
NTDB id 230 SPD RS09890 WP 074196785.1	
NTDB id 196 SPR RS09425 WP 074196785.1	
NTDB id 161 SP RS10375 WP 074196785.1	
NTDB id 529 SMSK321 RS09235 WP 080550783.1	
NTDB id 501 SM12261 RS00950 WP 004238896.1	
consensus	



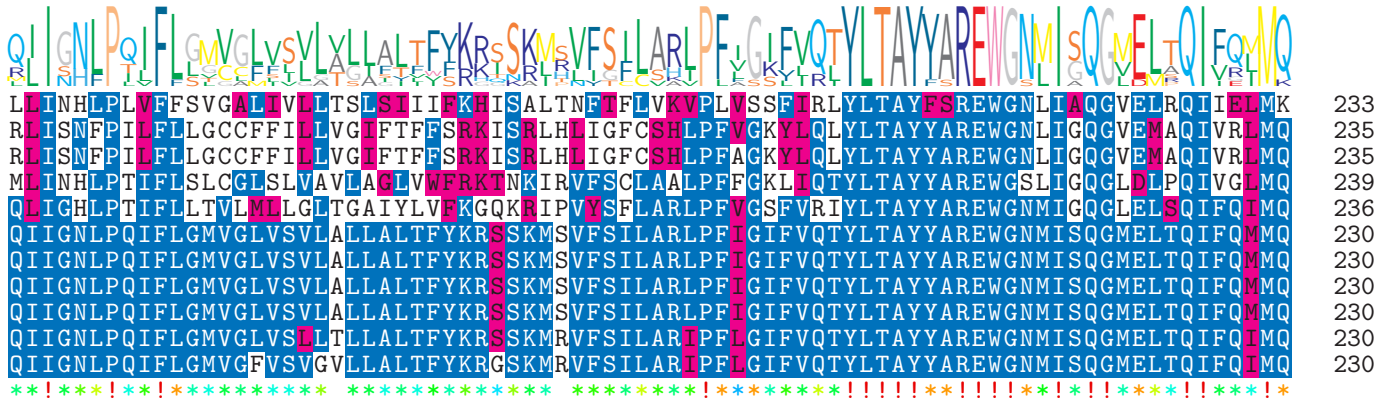
logo

NTDB id 587 KW2 RS10850 WP 042211510.1	ESLGFSSAVTQLSLAEVHGNLHLSLGLKEEYLDNLAKVKKKLEI VATYPVLLGLFLLIMGLRNYLLPQLDSSNIAT
NTDB id 395 FSA28 RS09400 WP 257638827.1	ESLGFSSAVTQLSLAEVHGNLHLSLGLKEEYLDNLAKVKKKLEI VATYPVLLGLFLLIMGLRNYLLPQLDSSNIAT
NTDB id 386 SMU RS09030 WP 255262721.1	SSLGFSDDVTQLALSEVHGNTQKSLKIESYLNLLIVKKKLEI VATYPVLLGLFLLIMGLRNYLLPQLDSSNIAT
NTDB id 1129581 GPW69 RS00750 WP 269472402.1	ADLRFSDAVVTQVALAEVHGNTSLSLHIGSYLKNVSKVRKKLEI VATYPVLLGLFLLIMGLRNYLLPQLDSSNIAT
NTDB id 430 SGO RS09415 WP 012130941.1	ASLGFSDAVVTQLSLAEVHGNLHLSLGLKEEYLDNLAKVKKKLEI VATYPVLLGLFLLIMGLRNYLLPQLDSSNIAT
NTDB id 271 KZH43 RS09260 WP 074196785.1	ESLGCSSAIVTQLSLAEVHGNLHLSLGLKEEYLDNLAKVKKKLEI VATYPVLLGLFLLIMGLRNYLLPQLDSSNIAT
NTDB id 230 SPD RS09890 WP 074196785.1	ESLGCSSAIVTQLSLAEVHGNLHLSLGLKEEYLDNLAKVKKKLEI VATYPVLLGLFLLIMGLRNYLLPQLDSSNIAT
NTDB id 196 SPR RS09425 WP 074196785.1	ESLGCSSAIVTQLSLAEVHGNLHLSLGLKEEYLDNLAKVKKKLEI VATYPVLLGLFLLIMGLRNYLLPQLDSSNIAT
NTDB id 161 SP RS10375 WP 074196785.1	ESLGCSSAIVTQLSLAEVHGNLHLSLGLKEEYLDNLAKVKKKLEI VATYPVLLGLFLLIMGLRNYLLPQLDSSNIAT
NTDB id 529 SMSK321 RS09235 WP 080550783.1	ESLGFSSAIVTQLSLAEVHGNLHLSLGLKEEYLDNLAKVKKKLEI VATYPVLLGLFLLIMGLRNYLLPQLDSSNIAT
NTDB id 501 SM12261 RS00950 WP 004238896.1	ESLGFSSAIVTQLSLAEVHGNLHLSLGLKEEYLDNLAKVKKKLEI VATYPVLLGLFLLIMGLRNYLLPQLDSSNIAT
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NTDB id 587 KW2 RS10850 WP 042211510.1	LLINHLPLVFFSVGALIVLLTSLIIFKHISALTNFTFLVKVPLVSSFIRLYLTAYPSREWGNIQAQGVELRQIIEIMK
NTDB id 395 FSA28 RS09400 WP 257638827.1	RLISNFPILFLLGCCFFILLVGI FTFFSRKISRLHLIGFCSHLPFVCKYLQLYLTAYYAREWGNIIGQGVEMAQIVRLMQ
NTDB id 386 SMU RS09030 WP 255262721.1	RLISNFPILFLLGCCFFILLVGI FTFFSRKISRLHLIGFCSHLPFVCKYLQLYLTAYYAREWGNIIGQGVEMAQIVRLMQ
NTDB id 1129581 GPW69 RS00750 WP 269472402.1	MLINHLPTIFLSLGLSLVAVLACLVMFRKTNKIRVFSCLAAALPFFGKLIQTYLTAYYAREWGNSLIGQGLDLPQIVGLMQ
NTDB id 430 SGO RS09415 WP 012130941.1	QLIGHLPTIFLLTVLMLLGLTGAIYLVFKGQKHIPVVSFLARLPFVGSFVRIYLTAYYAREWGNIIGQGLDLPQIVGLMQ
NTDB id 271 KZH43 RS09260 WP 074196785.1	QIIGNLPQIFLGMVGLVSVLALLALTIFYKRS SKMSVFSILARLPFV GIFVQTYLTAYYAREWGNIISQGMELTQIFQMMQ
NTDB id 230 SPD RS09890 WP 074196785.1	QIIGNLPQIFLGMVGLVSVLALLALTIFYKRS SKMSVFSILARLPFV GIFVQTYLTAYYAREWGNIISQGMELTQIFQMMQ
NTDB id 196 SPR RS09425 WP 074196785.1	QIIGNLPQIFLGMVGLVSVLALLALTIFYKRS SKMSVFSILARLPFV GIFVQTYLTAYYAREWGNIISQGMELTQIFQMMQ
NTDB id 161 SP RS10375 WP 074196785.1	QIIGNLPQIFLGMVGLVSVLALLALTIFYKRS SKMSVFSILARLPFV GIFVQTYLTAYYAREWGNIISQGMELTQIFQMMQ
NTDB id 529 SMSK321 RS09235 WP 080550783.1	QIIGNLPQIFLGMVGLVSVLALLALTIFYKRS SKMSVFSILARLPFV GIFVQTYLTAYYAREWGNIISQGMELTQIFQMMQ
NTDB id 501 SM12261 RS00950 WP 004238896.1	QIIGNLPQIFLGMVGLVSVLALLALTIFYKRS SKMSVFSILARLPFV GIFVQTYLTAYYAREWGNIISQGMELTQIFQMMQ
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consensus



 non conserved
 similar
 ≥ 50% conserved