

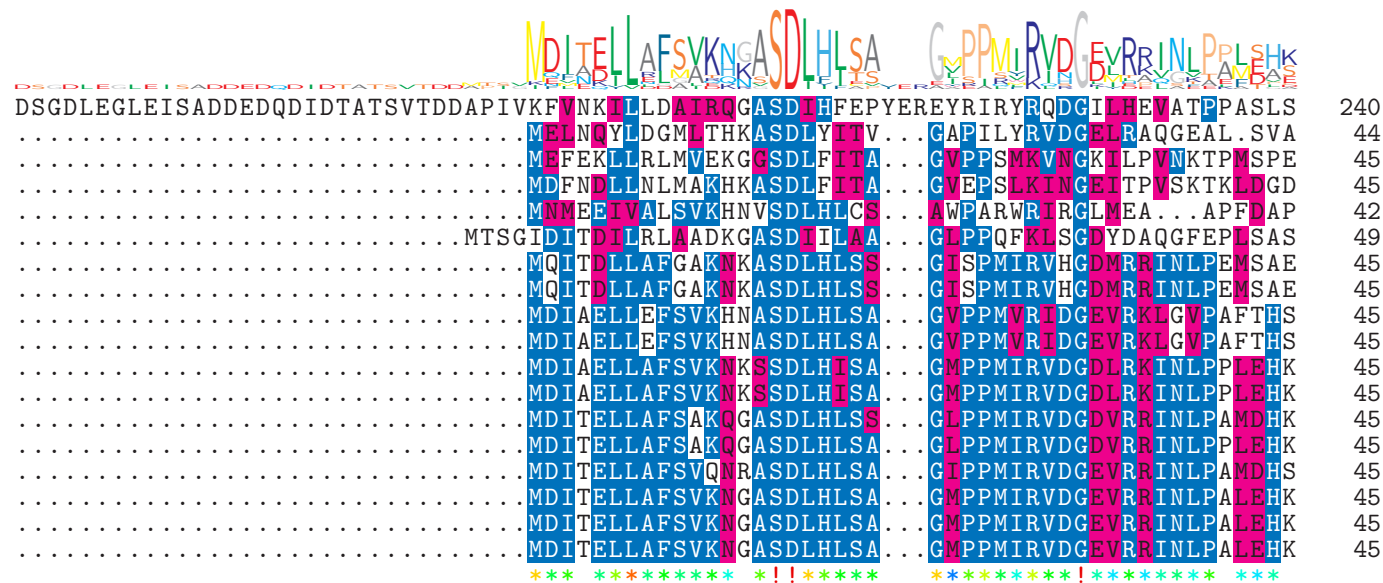
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consensus		

logo

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consensus		

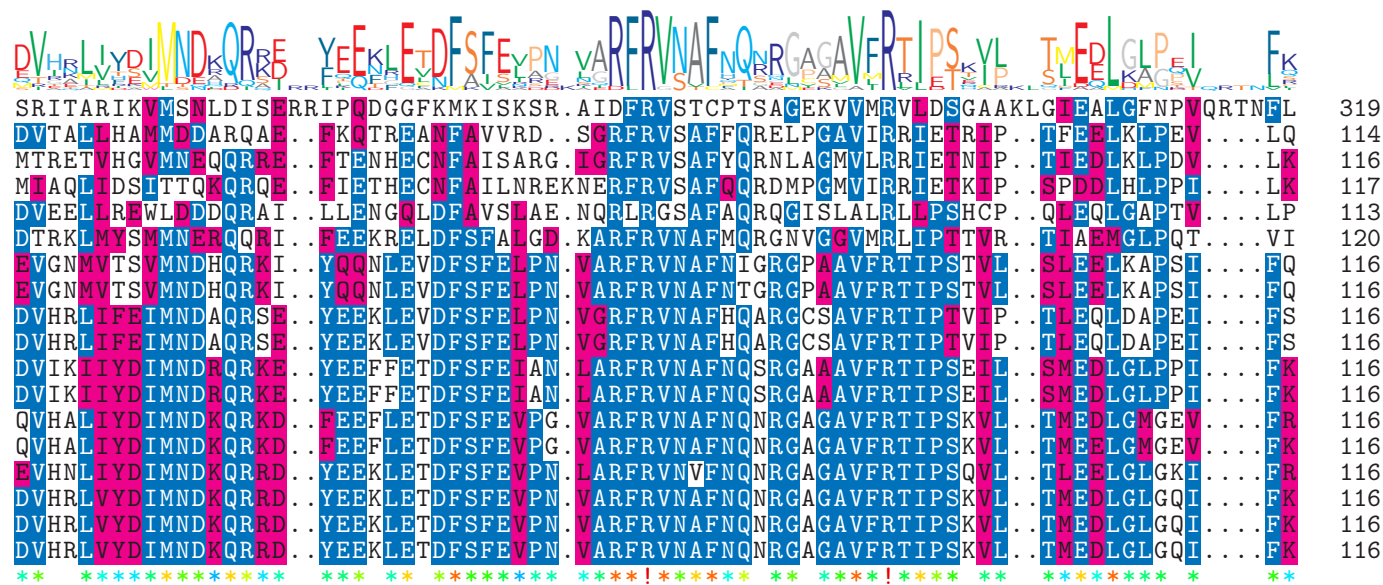
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consensus



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NTDB id 1060 ABD1 RS04380 WP 000355489.1
NTDB id 1040 HON27 RS13360 WP 000355489.1
NTDB id 1081 FDQ49 RS05335 WP 006581387.1
consensus

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NTDB id 1081 FDQ49 RS05335 WP 006581387.1
consensus

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NTDB id	1251	GC085	RS07725	WP	011946523.1
NTDB id	1177	A1552VC	RS01165	WP	000422572.1
NTDB id	1203	PSJM300	01605	AFN	76401.1
NTDB id	1019	ACIAD	RS04205	WP	004922051.1
NTDB id	264194	CXX14	RS10775	WP	001055620.1
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consensus					

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.....DKNKGRHGVEVLNSPRIADLIRRGELHELKATMARSQEVGM	312
.....TPDGKRRRAVIEVLINTPLAADLIRKGEVHELKGLMKRSTDLCGM	314
.....THDQSSRRAAVEVLINTPLLSDYIRQGEITYKIKELMARSQELGM	315
.....D.KQECGRVALFELLINTPAVGNLIREGKTHQLPHVIQTGQQVGM	310
.....RADGQGRVLAYEILLANPAVRALIREGKTFQITSVMQTGAREGM	318
.....THDGNGRVASHEILIANPAVRNLIRENKITTQINSVLQTGQASGM	314
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.....R.VGGGRVACHEIMLATPAIRNLIREDKVAQMYSIIQTGAAHGM	313
.....R.VGGGRVACHEIMLATPAIRNLIREDKVAQMYSIIQTGAAHGM	313
.....K.NKGGRVAALEIMMCTGAIRNLIREDKIAQMYSIIQTGQAKGM	313
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NTDB id	1251	GC085	RS07725	WP	011946523.1
NTDB id	1177	A1552VC	RS01165	WP	000422572.1
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NTDB id	1081	FDQ49	RS05335	WP	006581387.1
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QTFDQALYNLYVQGEITYEDALLHADSSNDLRLMIKLGS.ETDGEHLTSVS.QGLTL.....EMSDDDPGRSFR...	381
QTFDQALFDLYDHQHISYADALKHADSPNNLRLQIKLSE.EANKPLDNRERMSEMER.....KMTFDGQRRNLK...	383
LTFFQSYQQRVGEGRL.....	326
VTMDAYLAGLFRHHLITYDKGLERAIDPKEFARLANDASAGLGG.....ASMTPAAPPVPAYNSASSSSTGRGGDFGRGA	393
QTMDDQLQSLVRQGLIAPEVARRRAQNSESMSF.....	347
QTMDDQLQSLVRQGLIAPEAARRRAQNSESMSF.....	347
QTMENAKQLIARGVVDQEVQSKIELDLKAF.....	345
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QTLDDHLTELNVKNIIISKHTAHEAALNK.SLF.....	344
QTLDSCLKGLIAKGLISRDSAKEKAKQP.ENF.....	344
QTLDMCLKGLIAKGLISRAREKAKIP.ENF.....	344
TTLDDQLKQLIAKGIINLPTARAVAKSP.ESF.....	344
TTLDDQLKGLVARGVISPQTARTAAKQP.ESFL.....	345
TTLDDQLKGLVARGVISPQTARTAAKQP.ESFL.....	345
TTLDDQLKGLVSRGVISPQTARTAAKQP.ESFL.....	345
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NTDB id 1040 HON27 RS13360 WP 000355489.1		345
NTDB id 1081 FDQ49 RS05335 WP 006581387.1		345
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved