

logo

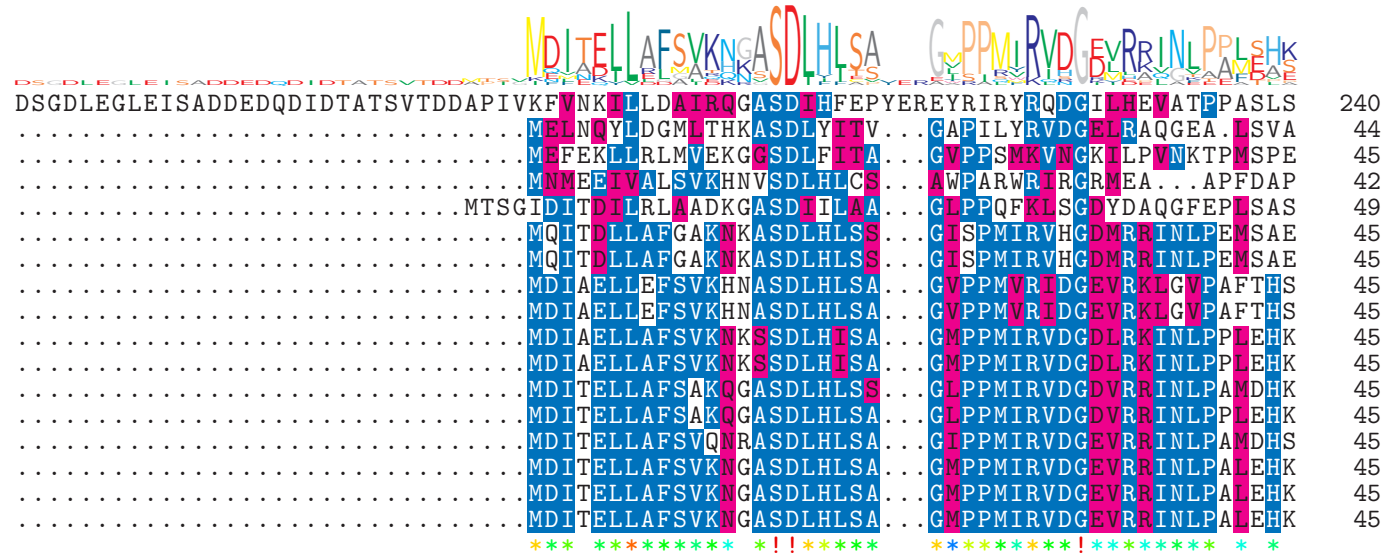
NTDB id 1251 GC085 RS07725 WP 011946523.1	MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEEKISLLQYIVKNKILSAEQIALTAAGNFGVPMLDINCLDVGTI	80
NTDB id 1177 A1552VC RS01165 WP 000422572.1	0
NTDB id 1203 PSJM300 01605 AFN76401.1	0
NTDB id 223913 A9X72 RS04235 WP 001324408.1	0
NTDB id 1307 DR RS10055 WP 010888596.1	0
NTDB id 1086 NMV RS00245 WP 002218487.1	0
NTDB id 1116 NGFG RS10615 WP 003690200.1	0
NTDB id 1185 GTF74 RS11940 WP 000350195.1	0
NTDB id 1176 A1552VC RS01160 WP 000350195.1	0
NTDB id 1283 ACE17W RS02630 WP 010947729.1	0
NTDB id 1259 GC085 RS10305 WP 010947729.1	0
NTDB id 1202 PSJM300 01600 AFN76400.1	0
NTDB id 1194 PAKAF RS01995 WP 003084552.1	0
NTDB id 1020 ACIAD RS04210 WP 011182164.1	0
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	0
NTDB id 1060 ABD1 RS04380 WP 000355489.1	0
NTDB id 1040 HON27 RS13360 WP 000355489.1	0
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NTDB id 1251 GC085 RS07725 WP 011946523.1	PVNLVNEKLIKRIHAMVPLFSRGTNLYLATDDPSKQASLKEIQFHTGLNTHAIVVETDKLSALIDNLLTAKESQGLSEYVE	160
NTDB id 1177 A1552VC RS01165 WP 000422572.1	0
NTDB id 1203 PSJM300 01605 AFN76401.1	0
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NTDB id 1307 DR RS10055 WP 010888596.1	0
NTDB id 1086 NMV RS00245 WP 002218487.1	0
NTDB id 1116 NGFG RS10615 WP 003690200.1	0
NTDB id 1185 GTF74 RS11940 WP 000350195.1	0
NTDB id 1176 A1552VC RS01160 WP 000350195.1	0
NTDB id 1283 ACE17W RS02630 WP 010947729.1	0
NTDB id 1259 GC085 RS10305 WP 010947729.1	0
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AVRDDFTNQGLIELGFKEADLVNKLKLYKAVGCEQCTSGYRGRVGLFEVLPMTKELGQLIMSGGNS.LDILKLAQSEGML 548
KNKGRHGVEVFLNSPRIADLIRRGELHELKATMARSQEVGMQ 313
PDGKRRRAVIEVLINTPLAADLIRKGEVHELKGLMKRSTDLGMQ 315
KQEGCVALFELLINTPAVGNLIREGKTHQLPHVIQTGGQVGM 311
ADGQGRVLAYEILLANPAVRALIREGKTFQITSVMTGAREGMV 319
HDGNGRVASHEILIANPAVRNLIRENKITQINSVLQTGGASGMQ 315
HDGNGRVASHEILIANPAVRNLIRENKITQINSVLQTGGASGMQ 315
VGGGRVACHEIMLATPAIRNLIREDKVAQMYSIIQTGAAHGMQ 314
VGGGRVACHEIMLATPAIRNLIREDKVAQMYSIIQTGAAHGMQ 314
NKGGRVAALEIMMCTGAIRNLIREDKIAQMYSIIQTGQAKGMQ 314
NKGGRVAALEIMMCTGAIRNLIREDKIAQMYSIIQTGQAKGMQ 314
VGGGRVAAHEIMIGTPAIRNLIREDKVAQMYSIIQTGGSLGMQ 314
IGGGRVAHEIMIGTPAIRNLIREDKVAQMYSIIQTGGSLGMQ 314
EGGGRVAAHEIMIGIPAIRNLIREGKVAQMYSIIQTGATHGMT 314
NCGGRVAAHEIMIGIPAIRNLIRENKVAQMYSIIQTGANHGMT 314
NCGGRVAAHEIMIGIPAIRNLIRENKVAQMYSIIQTGANHGMT 314
NCGGRVAAHEIMIGIPAIRNLIRENKVAQMYSIIQTGANHGMT 314

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 NTDB id 1040 HON27 RS13360 WP 000355489.1

TLDQSLKGLVARGVLSQETAREAKSPESFL 574
 TIFQSGIEKVKEGITIEEVNRVTVD..... 368
 TFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKRGDD.YGSG..SLQNVKI...DME..... 381
 TFDQALYNLYVQGEITYEDALLHADSSNDLRLMIKLGSETD.GEHLTSVVSQGLTL...EMSDDDPGRSFR..... 326
 TFQQSFQQRVGEGRLL..... 399
 TMDAYLAGLFRHLLITYDKGLERAIDPKEFARLANDASAGLGGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAAPATP 399
 TMDQSLQSLVLRQGLIAPEVARRRAQNSESMSF..... 347
 TMDQSLQSLVLRQGLIAPEAARRRAQNSESMSF..... 347
 TMEQNAKQLIARGVVDAQEVQSKIELDLKAF..... 345
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 TLDQHLTELVNKNIISKHTAHEAALNK.SLF..... 344
 TLDQHLTELVNKNIISKHTAHEAALNK.SLF..... 344
 TLDQSLKGLIAKGLISRDSAKEKAKQP.ENF..... 344
 TLDQSLKGLIAKGLISRDSAKEKAKQP.ENF..... 344
 TLDQSLKGLVARGVISPQTARTAAKQP.ESFL..... 345
 TLDQSLKGLVARGVISPQTARTAAKQP.ESFL..... 345
 TLDQSLKGLVARGVISPQTARTAAKQP.ESFL..... 345

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NTDB id 1251 GCO85 RS07725 WP 011946523.1	574
NTDB id 1177 A1552VC RS01165 WP 000422572.1	368
NTDB id 1203 PSJM300 01605 AFN76401.1	381
NTDB id 223913 A9X72 RS04235 WP 001324408.1	326
NTDB id 1307 DR RS10055 WP 010888596.1	AAPAAGRSEPGRAGGFGRR	418
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NTDB id 1116 NGFG RS10615 WP 003690200.1	347
NTDB id 1185 GTF74 RS11940 WP 000350195.1	345
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NTDB id 1283 ACE17W RS02630 WP 010947729.1	344
NTDB id 1259 GCO85 RS10305 WP 010947729.1	344
NTDB id 1202 PSJM300 01600 AFN76400.1	344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	345
NTDB id 1040 HON27 RS13360 WP 000355489.1	345
consensus		

^A^P^A^A^G^R^S^E^P^G^R^A^G^G^F^G^R^R

- X non conserved
- X similar
- X $\geq 50\%$ conserved