

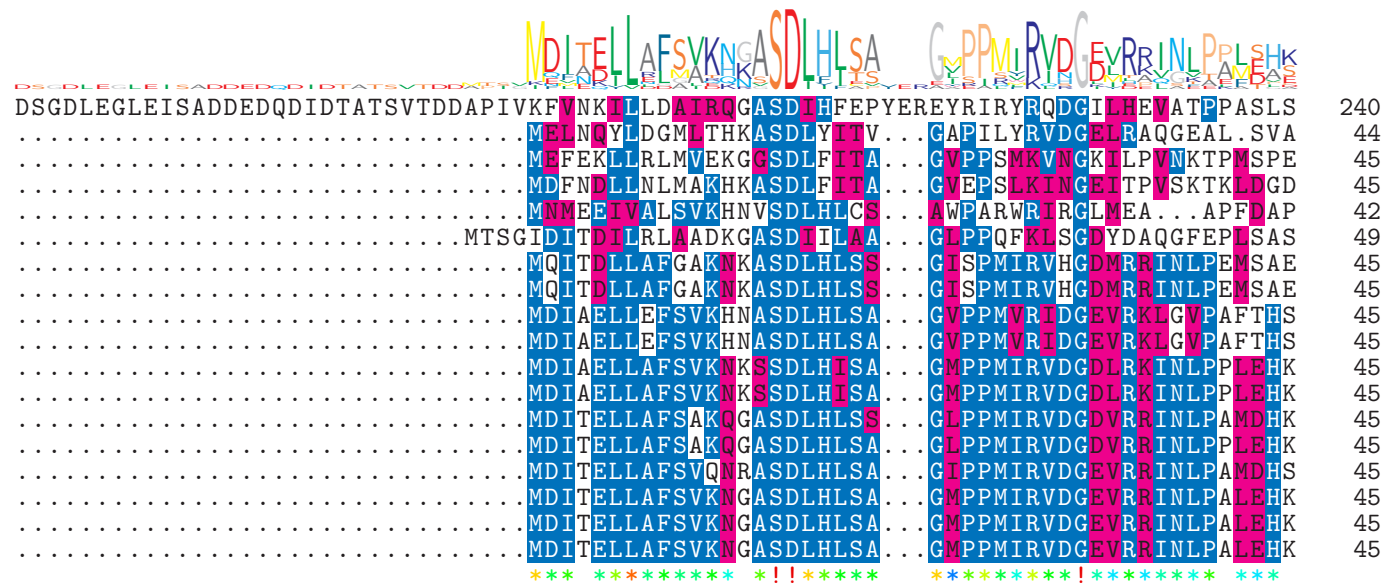
logo

NTDB id 1251 GC085 RS07725 WP 011946523.1	MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEEKISLLQYIVKNKILSAEQIALTAAGNFQVPMLDINCLDVGTI	80
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NTDB id 1203 PSJM300 01605 AFN76401.1		0
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NTDB id 277299 C6W73 RS15575 WP 001055620.1		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
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NTDB id 1040 HON27 RS13360 WP 000355489.1		0
NTDB id 1081 FDQ49 RS05335 WP 006581387.1		0
consensus		

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NTDB id 1202 PSJM300 01600 AFN76400.1		0
NTDB id 1194 PAKAF RS01995 WP 003084552.1		0
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consensus		

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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	277299	C6W73	RS15575	WP	001055620.1
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NTDB id	1086	NMV	RS00245	WP	002218487.1
NTDB id	1116	NGFG	RS10615	WP	003690200.1
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NTDB id	1283	ACE17W	RS02630	WP	010947729.1
NTDB id	1259	GC085	RS10305	WP	010947729.1
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consensus					

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.....DKNKGRHG VF EVL LNSPRIADLIRRGELHELKATMARSQEVGM	312
.....TPDGKRRRAVIEVLINTPLAADLIRKGEVHELKGLMKRSTDLCGM	314
.....THDQSSRRAAVEVLINTPLLSDYIRQGEITYKIKELMARSQELGM	315
.....D.KQECRVALFELLINTPAVGNLIREGKTHQLPHVIQTGQQVGM	310
.....RADGQGRVLAYEILLANPAVRALIREGKTFQITSVMQTGAREGM	318
.....THDGNGRVASHEILIANPAVRNLIRENKITTQINSVLQTGQASGM	314
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.....R.VGGGRVACHEIMLATPAIRNLIREDKVAQMYSIIQTGA AHGM	313
.....K.NKGG RVA ALEIMMCTGAIRNLIREDKIAQMYSIIQTGQAKGM	313
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.....K.VGGGRVAAHEIMIGTPAIRNLIREDKVAQMYSIIQTGGS LGM	313
.....K.IGGGRVAAHEIMIGTPAIRNLIREDKVAQMYSIIQTGGS LGM	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
consensus					

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QTFDQALYNLYVQGEITYEDALLHADSSNDLRLMIKLGS.ETDGEHLTSVS.QGLTL.....EMSDDD PGRSFR...	381
QTFDQALFDLYDHQHISYADALKHADSPNNLRLQIKLSE.EANKPLDNRERMSEMER.....KMTFDGQRRNLK...	383
LTFFQSYQQRVGEGRL.....	326
VTMDAYLAGLFRHHLITYDKGLERAIDPKEFARLANDASAGLGG....ASMTPAAPPVPAYNSASSSSTGRGGDFGRGA	393
QTMDQSLQSLVRQGLIAPEVARRRAQNSESMF.....	347
QTMDQSLQSLVRQGLIAPEAARRRAQNSESMF.....	347
QTMEQNAKQLIARGVVDAQEVQSKI ELDLKA F.....	345
QTMEQNAKQLIARGVVDAQEVQSKI ELDLKA F.....	345
QTLDQHLTELNVNKNIIISKHTAHEAALNK.SLF.....	344
QTLDQHLTELNVNKNIIISKHTAHEAALNK.SLF.....	344
QTLDQHLTELNVNKNIIISKHTAHEAALNK.SLF.....	344
QTLDQSLKGLIAGKGLISRDSAKEKAKQP.ENF.....	344
QTLDQSLKGLIAGKGLISRDSAKEKAKQP.ENF.....	344
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NTDB id 1251 GC085 RS07725 WP 011946523.1		574
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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