

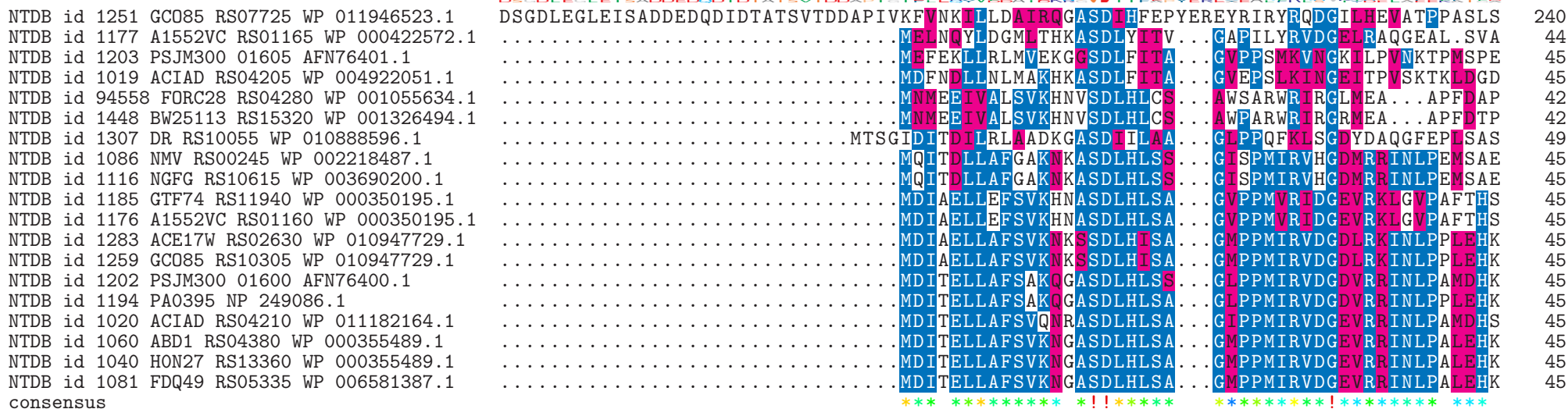
logo

NTDB id 1251 GC085 RS07725 WP 011946523.1	MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEEKISLLQYIVKNKILSAEQIALTAQNFQVPMLDINCLDVGTTI	80
NTDB id 1177 A1552VC RS01165 WP 000422572.1	0
NTDB id 1203 PSJM300 01605 AFN76401.1	0
NTDB id 1019 ACIAD RS04205 WP 004922051.1	0
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NTDB id 1448 BW25113 RS15320 WP 001326494.1	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
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NTDB id 1194 PA0395 NP 249086.1	0
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NTDB id 1060 ABD1 RS04380 WP 000355489.1	0
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 1251 GC085 RS07725 WP 011946523.1	PVNLVNEKLIKRRHAMVPLFSRGTNLYLATDDPSKQASLKEIQFHTGLNTHAIVVETDKLSALIDNLLTAKESQGLSEYVE	160
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NTDB id 1307 DR RS10055 WP 010888596.1	0
NTDB id 1086 NMV RS00245 WP 002218487.1	0
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NTDB id 1185 GTF74 RS11940 WP 000350195.1	0
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consensus		

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NTDB id 1251 GC085 RS07725 WP 011946523.1	KAVRDDFTNQGLIELGFKEADLVNLKLYKAVGCEQCTSGYRGRVGLFEVLPMTKELGQLIMSGGNS.LDILKLAQSEGM	547
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NTDB id 1081 FDQ49 RS05335 WP 006581387.1K.NGGGRVAACHEIMIGIPAIRNLIRENKVAQMYSIIQTGANHGM	313
consensus	* **!*!!!!!!*****!!!!* *****!!!!	

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NTDB id 1251 GC085 RS07725 WP 011946523.1	LTIFQSGIEKVKECITITIEEVNRVTVD	574
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NTDB id 1307 DR RS10055 WP 010888596.1	VTMDAYLAGLFRHLITIDYKGLERAIDPKFARLANDASAGLGG....ASMTPAAPPVPAYNSASSSSTGRGGDFGRGA	393
NTDB id 1086 NMV RS00245 WP 002218487.1	QTMDDQLQSLVRQGLIAPEVARRRAQNSESMSF.....	347
NTDB id 1116 NGFG RS10615 WP 003690200.1	QTMDDQLQSLVRQGLIAPEAARRRAQNSESMSF.....	347
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consensus	*!*** * ** ***** * *	

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NTDB id 1251 GC085 RS07725 WP 011946523.1	574
NTDB id 1177 A1552VC RS01165 WP 000422572.1	368
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consensus		

 non conserved
 similar
 ≥ 50% conserved