

NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDRGR.LVEATTEAEDLRTAARLLRDRCGLFAETKEPGKGLQAEVRIPALER	60
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKATQTLP LKNYRWKGINSNGKK.VSGQLLAISEIEVRDKLKDQHIIQIKKLKKGVSLSLAR....LTH	63
NTDB id 1402	DSB67 RS12670 WP 010643257.1	MKTIAPO LKNYRWKGINSCKK.TSGNVLAMTEIEVRERIDAQHIIKKLKKGKSISFITK....ISH	62
NTDB id 1112	NFGF RS09215 WP 003689811.1	MAKNGGFSLF AKKEKR FIFEGRHASADKL VNGEVS AFTEEEARKK LA KRGI RPLQITRVK.....TS....SKR	65
NTDB id 1113	AAA85695.1 219..1451()	MAKNGGFSLF AKKEKR FIFEGRHASADKL VNGEVS AFTEEEARKK LA KRGI RPLQITRVK.....TS....SKR	65
NTDB id 1198	PSJM300 03950 AFN76868.1	MAQKA I KNSVF TMEGLDRQGAK.LKGELSGVSPAL VKAQ LRKQGINPQKV RK KSIS LFS.....AGK	61
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAKKTQV MPV FAYEGVD RKG I KLKGEL PARNMAL AKVT LRKQG IT RTIR EKR KN IFEGL...LK	64
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MAVKKA QMMPT FAYEGVD RKG V KLKGEL PAKNMAL AKVT LRKQGV TVRN IREK RN ILEGL...FK	64
NTDB id 30868	XCCB100 RS05200 WP 012437601.1	MSVARSAIKKQPVD RNTSML QT FIMEGAD RKG V KLKGELT ARNAN MLRAEL RRQGIV PSVV KQKP LF GA...AGK	74
NTDB id 1252	GCO85 RS07730 WP 011213805.1	MDKNSPPLLTFHYCGINKAKGK.MEGDIQARS LA TAKAD LRKQGI VTNKV KKRP LF DR....KNK	62
consensus		* * * * *	

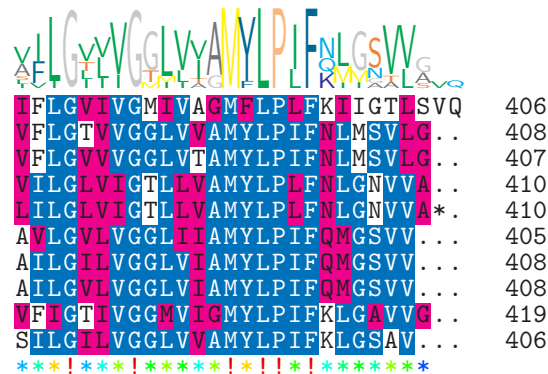
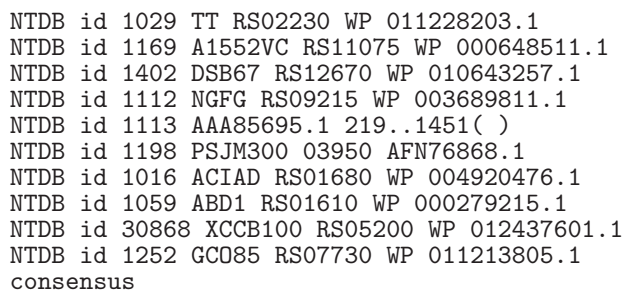
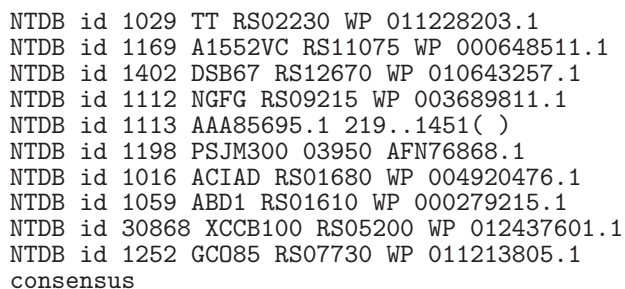
NTDB id	1029	TT RS02230 WP 011228203.1	GPGGLKDIAIFSRQLATMLGAGLTLLQALATLERQTENRKFRREILKQVRTDVEGGMAFSEALSKHK.IFSRLYVNLVRAGE	139
NTDB id	1169	A1552VC RS11075 WP 000648511.1	RVKSKDITILLRQLATMLTTGVPIVQALKLVGDNRKAEMKSILAQITKSVEAGTPLSKAMRTASAHFDTLVVDLVTETGE	143
NTDB id	1402	DSB67 RS12670 WP 010643257.1	RVKGKDITIFTTRQLATMLMTGVPIVQALKLVSENHKKAEKMSILMSVTRAVEAGTPMSKAMRTASTHFDPPLYTDLIATGE	142
NTDB id	1112	NFGF RS09215 WP 003689811.1	KITQEDITVFTRQLSTMIKAKGLPLMQAFEIVARGHGNSPMTLEIMETRGQVEQGSSLSRAFSNHPKYFYDRFYCNLVAAGE	145
NTDB id	1113	AAA85695.1 219..1451()	KITQEDITVFTRQLSTMINAGLPIMQAFEIVARGHGNSPMTLEIMEIRGQVEEQGSSLSRAFSNHPKYFYDRFYCNLVAAGE	145
NTDB id	1198	PJM300 03950 AFN76868.1	KIKPMDIALFTRQMATMMKAGVPLIQSFDIIGEGFDNPNNRKLVDLDLKQEVAAAGNSFASSLRKPQYFDDLFCYLNDVSGE	141
NTDB id	1016	ACIAD RS01680 WP 004920476.1	KVKPLDIAIFTRQLATMMKAGVPLVQAFAEIVAEGLENPAMRDVVILGKNEVEGGNTFAGALRKYPQYFDNFLFCSLIESGE	144
NTDB id	1059	ABD1 RS01610 WP 000279215.1	KVTTLDTITFTTRQLATMMKAGVPLVQCFEIVAEGLENPAMREVVLGTKEVEEGGSTFASALRKYPQHFDNFLFCSLVESGE	144
NTDB id	30868	XCCB100 RS05200 WP 012437601.1	KITPKDIAFFSRQMAMTMKSGVPITVSSLEIIGEGHKNPRMKMVGQIRTDIEGGSSLYESTSAHPVQFDELYRNLRAGE	154
NTDB id	1252	GC085 RS07730 WP 011213805.1	KITQADITVFSRQLATMLIESGIPLVQAQDIVAKGQSINKRLKDLIETIKHDJETGLTLAESLIKHPLYNFELFCNLVDAGE	142
consensus			* * * * *	

[illegible]

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NTDB id 1402 DSB67 RS12670 WP 010643257.1
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NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
NTDB id 30868 XCCB100 RS05200 WP 012437601.1
NTDB id 1252 GCO85 RS07730 WP 011213805.1
consensus

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-  non conserved
-  similar
-  > 50% conserved