

NTDB id 1029	TT RS02230 WP 011228203.1	MPVYQYKARDRQGR.LVEATIEAEDLRTAARLLRDRGLFVAEIKPEPKGLQAEVRIPALERGPGLK	65
NTDB id 1402	DSB67 RS12670 WP 010643257.1	.MKT.....IAPQLKNYRWKGINSSGKK.TSGNVLAMTEIEVRERLDAQHIKIKKLKKGSIKFITK.....ISHRVKGGK	67
NTDB id 1169	A1552VC RS11075 WP 000648511.1	MKAT.....QTLPLKNYRWKGINSGKK.VSGQMLAISEIEVRDCLKDQHIQIKKLKKGSVSLAR.....LTHRVKSK	68
NTDB id 1112	NGFG RS09215 WP 003689811.1	.MAKNGGFSLFAKKEKRFIFEGRHSASDKLNVGEVSAFTEEEARKKLAKRGIRPLQITRV.....KTSKKRKITQE	70
NTDB id 1113	AAA85695.1 219..1451()	.MAKNGGFSLFAPKKEKRFIFEGRHSASDKLNVGEVSAFTEEEARKKLAKRGIRPLQITRV.....KTSKKRKITQE	70
NTDB id 1252	GC085 RS07730 WP 011213805.1	.MDKN.....SPPLLTFHYQGINKAGQK.MEGDIQARSLATAKADLRKQGVITNKKVKKRPLFDR.....KNKKITQA	67
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NTDB id 1016	ACIAD RS01680 WP 004920476.1	MAAKK.....TQVMPVFAYEGVDRKGIK.LKGELPARNMALAKVTLRKQGITIRTIREKRKNIFEG....LLKKVKPL	69
NTDB id 391899	GBN68 RS01425 WP 234955119.1	MAVKK.....AQMMPTFNVEGIDRKGAK.IKGELPARNLALAKVTLRKQGITIKSIREKKKNILEG....ILKKKVTTL	69
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consensus		*** * * * * ! * * * * * * * * * *	

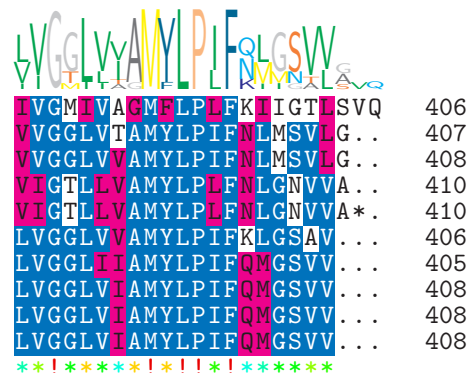
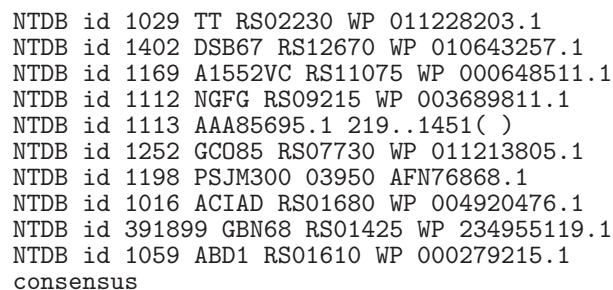
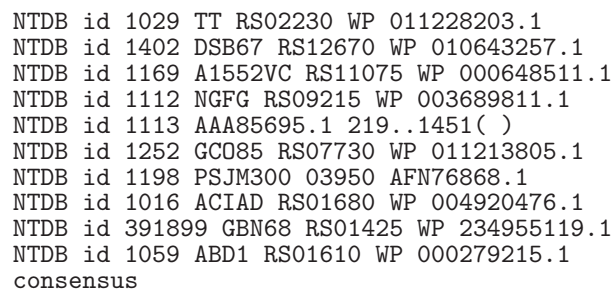
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NTDB id	1402	DSB67 RS12670 WP	010643257.1	S Q V F E R L A T Y R E K S E Q L R A K V I K A L I Y P A M V V L V A L G V S Y L M L T R V I P E F E K M F T G F G A D L P A F T Q M V L N L S A W T Q N W G P	227
NTDB id	1169	A1552VC RS11075 WP	000648511.1	P E V F E R L A T Y R E K S E Q L R A K V I K A L I Y P S M V V L V A L G V S Y L M L T M V I P E F E S M F K G F G A E L P W F T Q Q V L K L S H W V Q A Y S L	228
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consensus				* * * * *	

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NTDB id 391899 GBN68 RS01425 WP 234955119.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus

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- X non conserved
- X similar
- X > 50% conserved