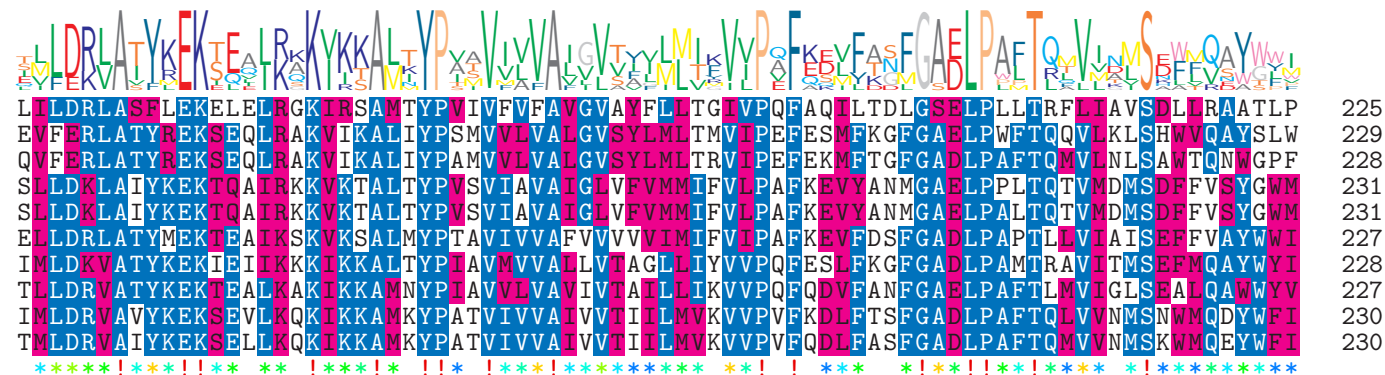
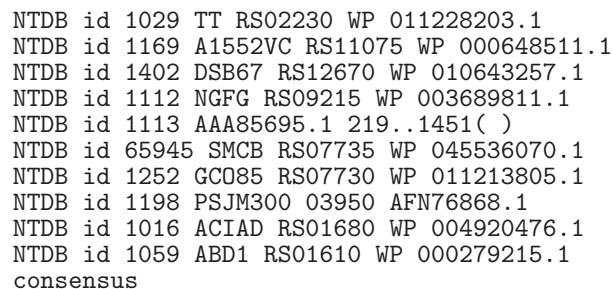
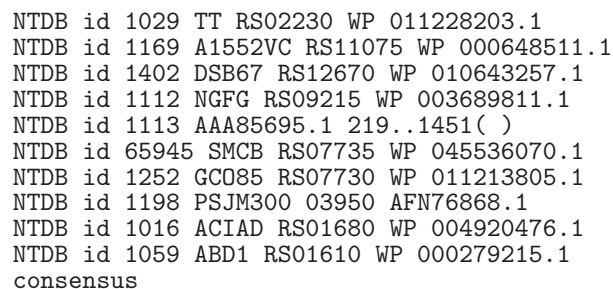


```

NTDB id 1029 TT RS02230 WP 011228203.1
NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 65945 SMCB RS07735 WP 045536070.1
NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus

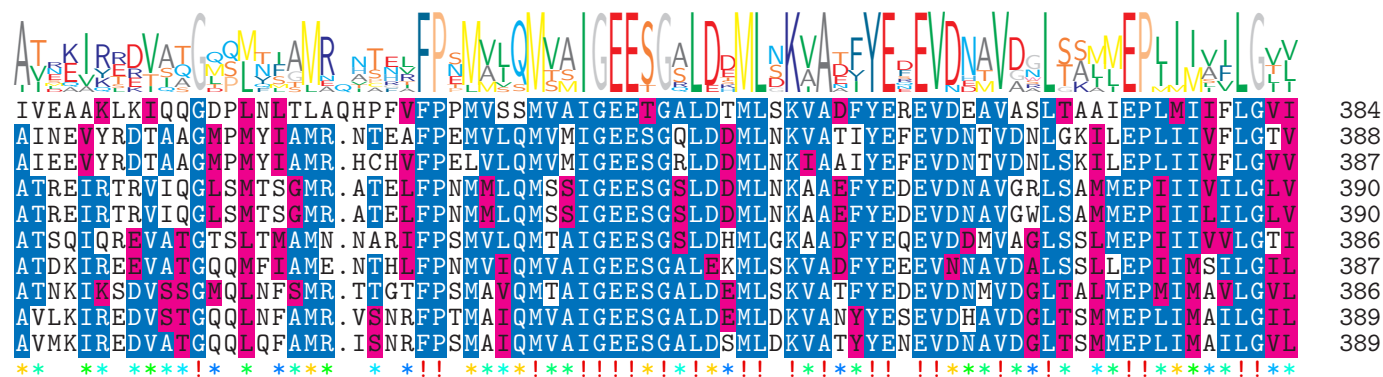
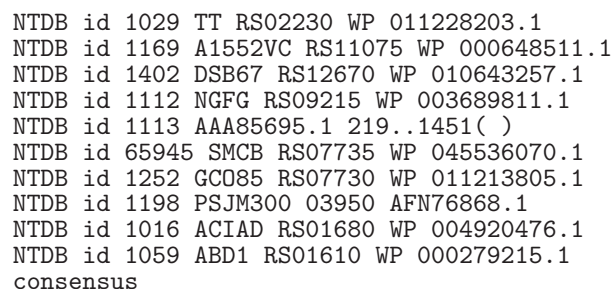
```



```

NTDB id 1029 TT RS02230 WP 011228203.1
NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 65945 SMCB RS07735 WP 045536070.1
NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus

```



NTDB id 1029	TT RS02230 WP 011228203.1	VGMIIVAGMFLPLFKITIGTL	LSVQ	406
NTDB id 1169	A1552VC RS11075 WP 000648511.1	VGGLVVMYLPINFNLMSV	LG..	408
NTDB id 1402	DSB67 RS12670 WP 010643257.1	VGGLVTAMYLPINFNLMSV	LG..	407
NTDB id 1112	NGFG RS09215 WP 003689811.1	IGTLLVAMYLPLFLNGLN	VVA..	410
NTDB id 1113	AAA85695.1 219..1451()	IGTLVAMYLPLFLNGLN	VVA*	410
NTDB id 65945	SMCB RS07735 WP 045536070.1	IGGIIVVSMYLPFIQLGQ	VVV..	405
NTDB id 1252	GC085 RS07730 WP 011213805.1	VGGLVVMYLPFIKLGSAV	...	406
NTDB id 1198	PSJM300 03950 AFN76868.1	VGGLIIAMYLPFIQMGSVV	...	405
NTDB id 1016	ACIAD RS01680 WP 004920476.1	VGGLVIAMYLPIFIQMGSVV	...	408
NTDB id 1059	ABD1 RS01610 WP 000279215.1	VGGLVIAMYLPIFIQMGSVV	...	408
consensus		* ! * * * * * ! * ! ! * ! * * * * *		

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