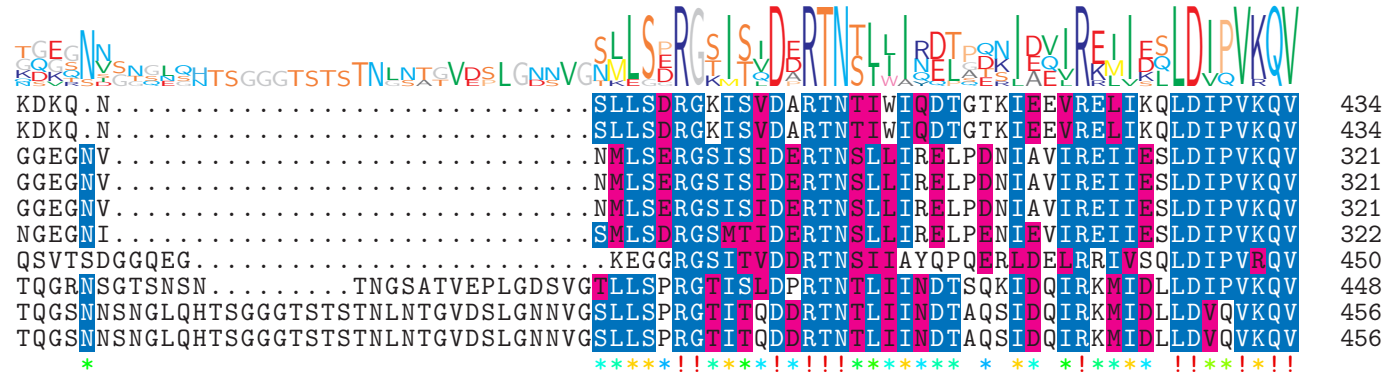
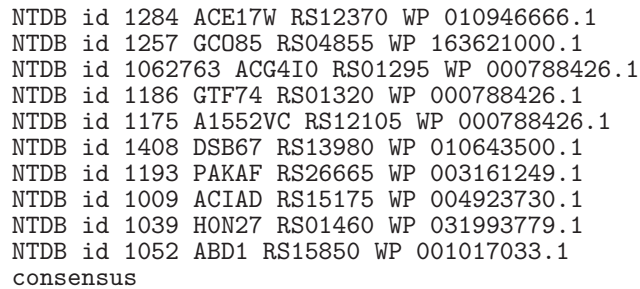
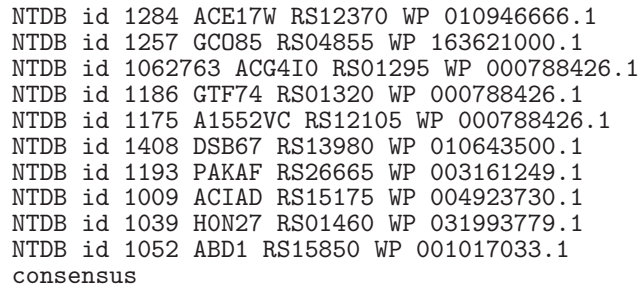


NTDB id 1284	ACE17W	RS12370	WP	010946666.1	MRKTIAILIL...ISASISMAYSKDNSLSISKVLPPEPENVRRIDFQFSKPLKQLPASFITQKPPRLVLDLISTDMQIPSDQ	78
NTDB id 1257	GC085	RS04855	WP	163621000.1	MRKTIAILIL...ISASISLAYSKDNSLSISKVLPPEPENVRRIDFQFSKPLKQLPASFITQKPPRLVLDLISTDMQIPSDQ	78
NTDB id 1062763	ACG4IO	RS01295	WP	000788426.1		0
NTDB id 1186	GTF74	RS01320	WP	000788426.1		0
NTDB id 1175	A1552VC	RS12105	WP	000788426.1		0
NTDB id 1408	DSB67	RS13980	WP	010643500.1		0
NTDB id 1193	PAKAF	RS26665	WP	003161249.1	MNSGLSRLGIALAAWFAPALLAADTEKDVAAPGDRVEIKLQFDEPVAAPRGYTIEQPARIALDLPGVQNKIG..T	76
NTDB id 1009	ACIAD	RS15175	WP	004923730.1	MKQIRQFSIGTVAYAIMQVASAQVAITNVPMNIPGIGTEIRVMFNGLPPQPQAYQLEQPARLILDFDKAGQKLA..K	77
NTDB id 1039	HON27	RS01460	WP	031993779.1	MNHVFRQFSMGAVAIAIMQAASAQVSMTNIVPMQIAGGTEIRVMFNGLPPQPQAYQLENPSRLILDFDKAQQGLK..Q	77
NTDB id 1052	ABD1	RS15850	WP	001017033.1	MNHVFRQFSMGAVAIAIMQAASAQVSMTNIVPMQIAGGTEIRVMFNGLPPQPQAYQLENPSRLILDFDKAQQGLK..Q	77
consensus					* * * * *	

NTDB id	1284 ACE17W RS12370 WP 010946666.1	KTKT IQLGSLHSYNIVAVGDVRRAILELDRSVSYSGSTAGSIIYSLVINGKSNDFENSKE.....VFITNQV	145
NTDB id	1257 GC085 RS04855 WP 163621000.1	KTKTIQLGSLHSYNIVAVGDVRRAILELDRSVSYSGSTAGSVIYSLVINGKSNDFENSKE.....VFITNQV	145
NTDB id	1062763 ACG4IO RS01295 WP 000788426.1	MKNGLKT..YVAQTLWLTLWG....LAICASS.....MVFSA.E	32
NTDB id	1186 GTF74 RS01320 WP 000788426.1	MKNGLKT..YVAQTLWLTLWG....LAICASS.....MVFSA.E	32
NTDB id	1175 A1552VC RS12105 WP 000788426.1	MKNGLKT..YVAQTLWLTLWG....LAICASS.....MVFSA.E	32
NTDB id	1408 DSB67 RS13980 WP 010643500.1	MKMRHGLTLPKRKL TGLFS LWFS....LL LAP.....FVAA.E	33
NTDB id	1193 PAKAF RS26665 WP 003161249.1	KNREISVGNTRS VTVVEAKDRTLRILINLTTLSSYTTRVEGNLFVVGVNSPAGASVASTAPVKASPAPASYAQPIKPYPY	156
NTDB id	1009 ACIAD RS15175 WP 004923730.1	SLIPVTTN EASSIDVSSDDKRARVVVN LKD AGAFTTRVEGN VFILK INNI QSTAT.....PVAT...SA.....	138
NTDB id	1039 HON27 RS01460 WP 031993779.1	SKIS VATNE ASSVDVTSDDQRSRL TVNL KDTGA FT TRVEGN TFILKINS AQTSNKPLPV.VS.....	138
NTDB id	1052 ABD1 RS15850 WP 001017033.1	SKIS VATNE ASSVDVTSDDQRSRL TVNL KDG AF GT TRVEGN TFILKINS AQTSNKPLPV.VS.....	138
consensus		* * * * *	

[illegible]

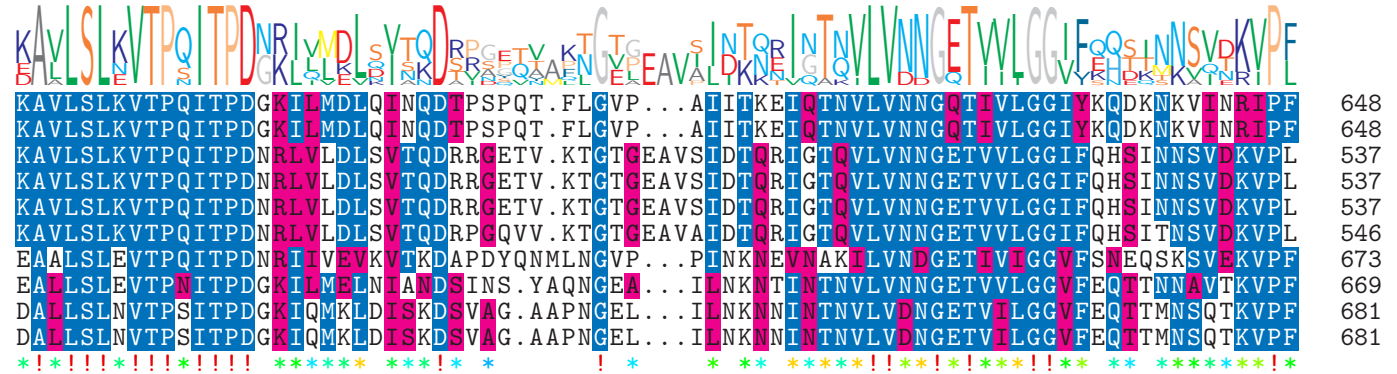
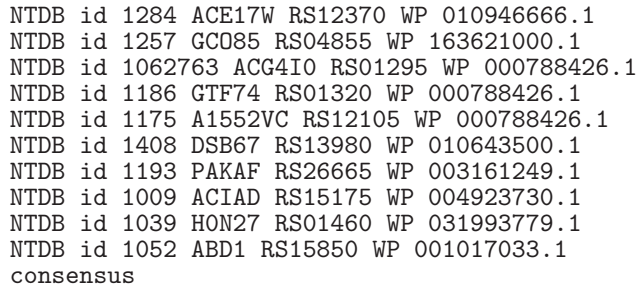
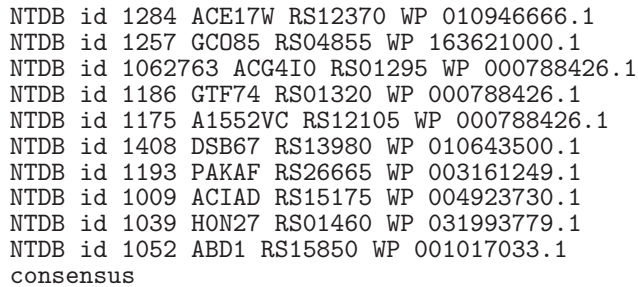
```
NTDB id 1284 ACE17W RS12370 WP 010946666.1
NTDB id 1257 GC085 RS04855 WP 163621000.1
NTDB id 1062763 ACG4IO RS01295 WP 000788426.1
NTDB id 1186 GTF74 RS1320 WP 000788426.1
NTDB id 1175 A1552VC RS12105 WP 000788426.1
NTDB id 1408 DSB67 RS13980 WP 010643500.1
NTDB id 1193 PAKAF RS26665 WP 003161249.1
NTDB id 1009 ACIAD RS15175 WP 004923730.1
NTDB id 1039 HON27 RS01460 WP 031993779.1
NTDB id 1052 ABD1 RS15850 WP 001017033.1
consensus
```



```

NTDB id 1284 ACE17W RS12370 WP 010946666.1
NTDB id 1257 GC085 RS04855 WP 163621000.1
NTDB id 1062763 ACG4I0 RS01295 WP 000788426.1
NTDB id 1186 GTF74 RS01320 WP 000788426.1
NTDB id 1175 A1552VC RS12105 WP 000788426.1
NTDB id 1408 DSB67 RS13980 WP 010643500.1
NTDB id 1193 PAKAF RS26665 WP 003161249.1
NTDB id 1009 ACIAD RS15175 WP 004923730.1
NTDB id 1039 HON27 RS01460 WP 031993779.1
NTDB id 1052 ABD1 RS15850 WP 001017033.1
consensus

```



logo



X non conserved
X similar
X ≥50% conserved