


```

NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 167664 AL475 RS17735 WP 104966495.1
NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 1252 GCO85 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
consensus

```

391
392
392
394
394
391
390
393
393

logo

NTDB id 1402 DSB67 RS12670 WP 010643257.1	VTAMYLPIFNLMSVLG.	407
NTDB id 167664 AL475 RS17735 WP 104966495.1	VVAMYLPIFNLMSVLG.	408
NTDB id 1169 A1552VC RS11075 WP 000648511.1	VVAMYLPIFNLMSVLG.	408
NTDB id 1112 NGFG RS09215 WP 003689811.1	LVAMYLPLFNLGNVVA.	410
NTDB id 1113 AAA85695.1 219..1451()	LVAMYLPLFNLGNVVA*	410
NTDB id 1252 GC085 RS07730 WP 011213805.1	VVAMYLPIFKLGSAV..	406
NTDB id 1198 PSJM300 03950 AFN76868.1	IIAMYLPIFQMGSSV..	405
NTDB id 1059 ABD1 RS01610 WP 000279215.1	VIAMYLPIFQMGSSV..	408
NTDB id 1016 ACIAD RS01680 WP 004920476.1	VIAMYLPIFQMGSSV..	408
consensus	***!!!!!!*!*****	

Sequence	Count
VTAMYLPIFNLMSVVLG	1
VTAMYLPIFNLMSVVLG	1
VVAMYLPIFNLMSVVLG	1
VVAMYLPIFNLMSVVLG	1
LVAMYLPIFLNLGNVVA	1
LVAMYLPIFLNLGNVVA	1
VVAMYLPIFLKLGSAV	1
ITAMYLPIFQMGSSV.	1
VIAMYLPIFQMGSSV.	1
VIAMYLPIFQMGSSV.	1

-  non conserved
-  similar
-  $\geq 50\%$ conserved