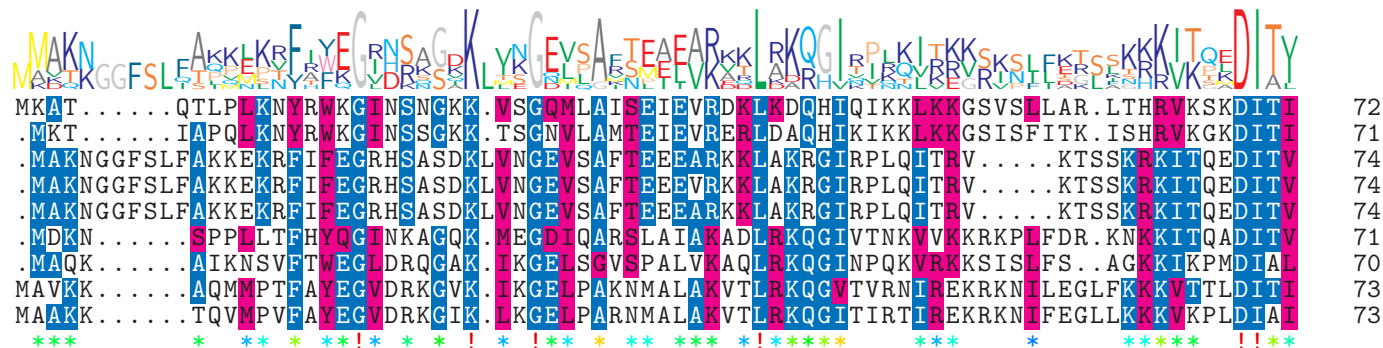


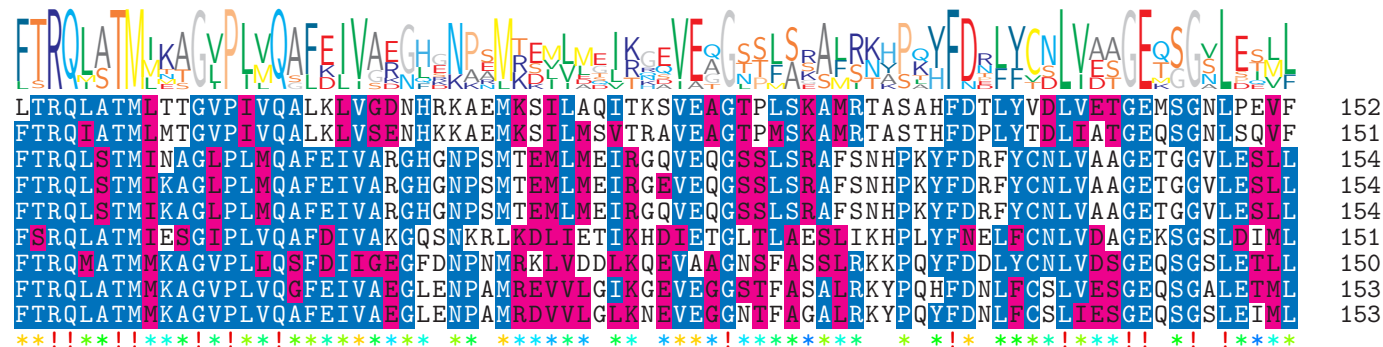
logo

NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1113 AAA85695.1 219..1451()
NTDB id 222635 A6J48 RS09470 WP 002248468.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1252 GC085 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



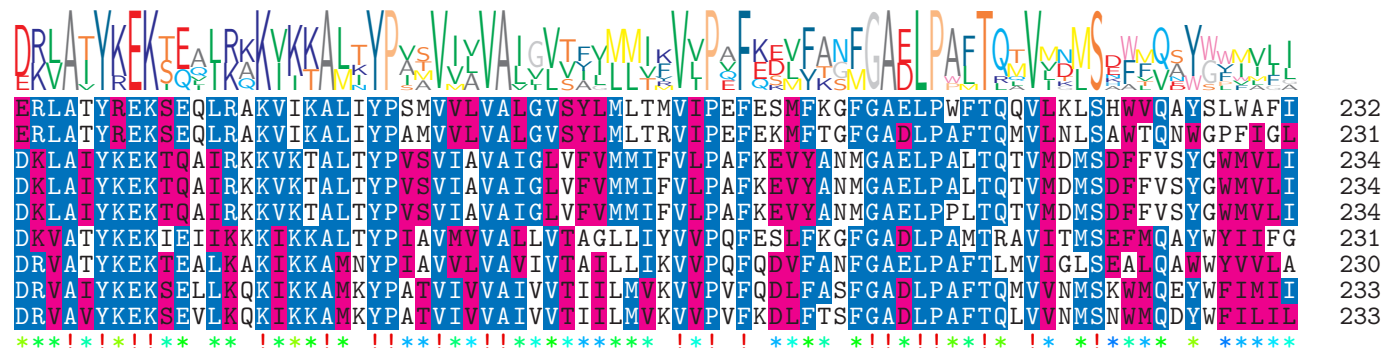
logo

NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1113 AAA85695.1 219..1451()
NTDB id 222635 A6J48 RS09470 WP 002248468.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1252 GC085 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



logo

NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1113 AAA85695.1 219..1451()
NTDB id 222635 A6J48 RS09470 WP 002248468.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1252 GC085 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



logo

NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1113 AAA85695.1 219..1451()
NTDB id 222635 A6J48 RS09470 WP 002248468.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1252 GC085 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



```

NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 222635 A6J48 RS09470 WP 002248468.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1252 GC085 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

```

392
391
394
394
394
391
390
393
393

logo

NTDB id 1169	A1552VC RS11075 WP 000648511.1	VVAMYLPIFNLMSVLG.	408
NTDB id 1402	DSB67 RS12670 WP 010643257.1	VTAMYLPIFNLMSVLG.	407
NTDB id 1113	AAA85695.1 219..1451()	LVAMYLPIFNLGNVVA*	410
NTDB id 222635	A6J48 RS09470 WP 002248468.1	LVAMYLPIFNLGNVVA.	410
NTDB id 1112	NGFG RS09215 WP 003689811.1	LVAMYLPIFNLGNVVA.	410
NTDB id 1252	GC085 RS07730 WP 011213805.1	VVAMYLPIFKLGSAV.	406
NTDB id 1198	PSJM300 03950 AFN76868.1	IIAMYLPIFQMGSSVV.	405
NTDB id 1059	ABD1 RS01610 WP 000279215.1	VIAMYLPIFQMGSSVV.	408
NTDB id 1016	ACIAD RS01680 WP 004920476.1	VIAMYLPIFQMGSSVV.	408
consensus		***!!!!!!*!*****	

Sequence logo

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