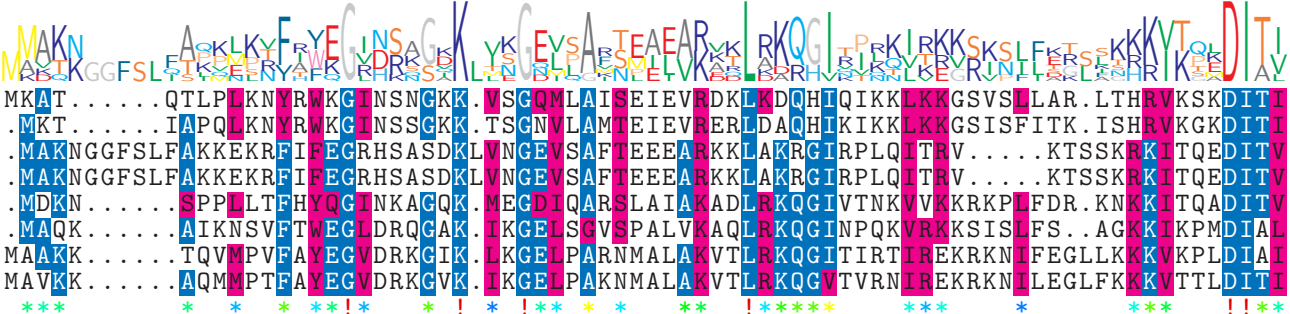


logo

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	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					



72
71
74
74
71
70
73
73

logo

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	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					



152
151
154
154
151
150
153
153

logo

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	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					



232
231
234
234
231
230
233
233

logo

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	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					



312
311
314
314
311
310
313
313

```

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 1252 GCO85 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

```

[illegible]

logo

```

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 1252 GCO85 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

```

Sequence	Score
VVAMYLP I FNLM S VLG.	408
VTAMYLP I FNLM S VLG.	407
LVAMYLP I FNLMGNVVA.	410
LVAMYLP I FNLMGNVVA.	410
VVAMYLP I FKLGS A V..	406
IIAMYLP I FQMGSSV..	405
VIAMLYLP I FQMGSSV..	408
VIAMLYLP I FQMGSSV..	408

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