

logo

NTDB id 1307 DR RS10055 WP 010888596.1
 NTDB id 1177 A1552VC RS01165 WP 000422572.1
 NTDB id 1019 ACIAD RS04205 WP 004922051.1
 NTDB id 125171 HV99 RS09605 WP 003100959.1
 NTDB id 1203 PSJM300 01605 AFN76401.1
 consensus



80
75
76
76
76

logo

NTDB id 1307 DR RS10055 WP 010888596.1
 NTDB id 1177 A1552VC RS01165 WP 000422572.1
 NTDB id 1019 ACIAD RS04205 WP 004922051.1
 NTDB id 125171 HV99 RS09605 WP 003100959.1
 NTDB id 1203 PSJM300 01605 AFN76401.1
 consensus



159
153
156
155
155

logo

NTDB id 1307 DR RS10055 WP 010888596.1
 NTDB id 1177 A1552VC RS01165 WP 000422572.1
 NTDB id 1019 ACIAD RS04205 WP 004922051.1
 NTDB id 125171 HV99 RS09605 WP 003100959.1
 NTDB id 1203 PSJM300 01605 AFN76401.1
 consensus



239
233
236
235
235

logo

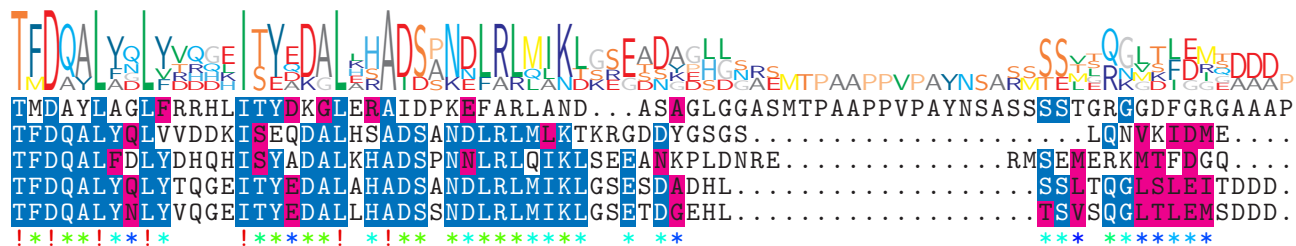
NTDB id 1307 DR RS10055 WP 010888596.1
 NTDB id 1177 A1552VC RS01165 WP 000422572.1
 NTDB id 1019 ACIAD RS04205 WP 004922051.1
 NTDB id 125171 HV99 RS09605 WP 003100959.1
 NTDB id 1203 PSJM300 01605 AFN76401.1
 consensus



319
313
316
315
315

logo

NTDB id 1307 DR RS10055 WP 010888596.1
 NTDB id 1177 A1552VC RS01165 WP 000422572.1
 NTDB id 1019 ACIAD RS04205 WP 004922051.1
 NTDB id 125171 HV99 RS09605 WP 003100959.1
 NTDB id 1203 PSJM300 01605 AFN76401.1
 consensus



396
368
378
375
375

logo

NTDB id 1307 DR RS10055 WP 010888596.1	ATPAAI	PA	GRS	NEPGRAGGL	FR	418
NTDB id 1177 A1552VC RS01165 WP 000422572.1	ATPAAI	PA	GRS	SEPGRAGG	FGR	368
NTDB id 1019 ACIAD RS04205 WP 004922051.1	RRN	L	383
NTDB id 125171 HV99 RS09605 WP 003100959.1	PA	GR	FR	382
NTDB id 1203 PSJM300 01605 AFN76401.1	PG	RS	FR	381
consensus		**	*		**	

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