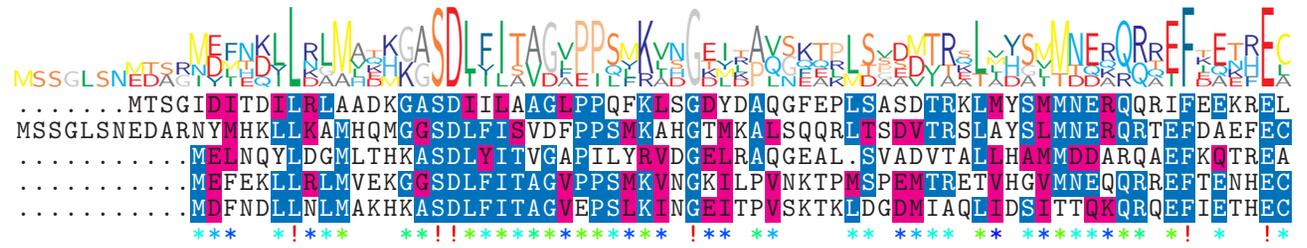


logo

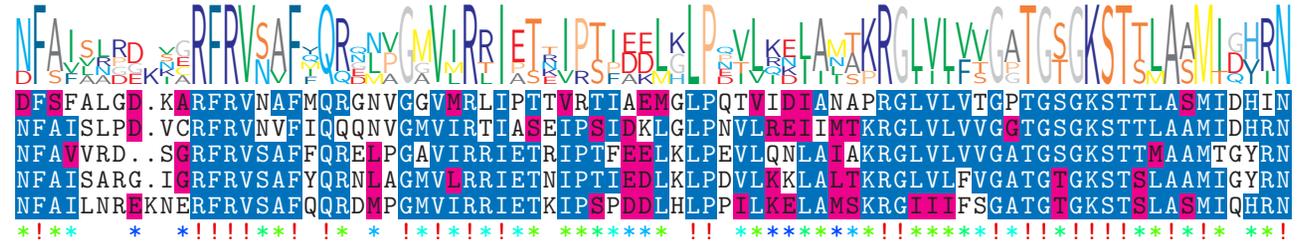
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 NTDB id 56709 AZKH RS03635 WP 015434384.1
 NTDB id 1177 A1552VC RS01165 WP 000422572.1
 NTDB id 1203 PSJM300 01605 AFN76401.1
 NTDB id 1019 ACIAD RS04205 WP 004922051.1
 consensus



73
80
68
69
69

logo

NTDB id 1307 DR RS10055 WP 010888596.1
 NTDB id 56709 AZKH RS03635 WP 015434384.1
 NTDB id 1177 A1552VC RS01165 WP 000422572.1
 NTDB id 1203 PSJM300 01605 AFN76401.1
 NTDB id 1019 ACIAD RS04205 WP 004922051.1
 consensus



152
159
146
148
149

logo

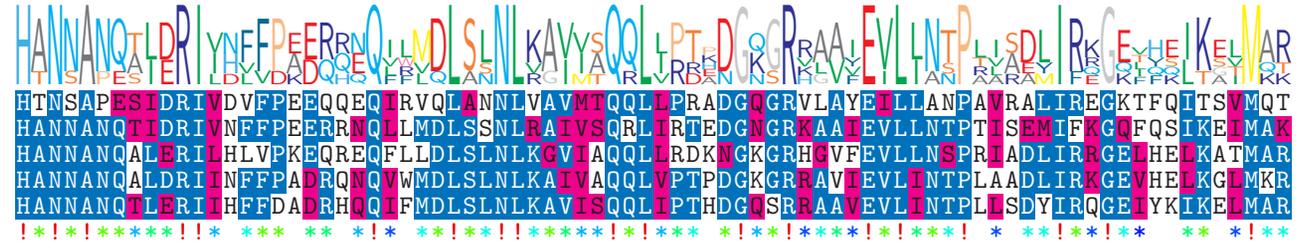
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 NTDB id 56709 AZKH RS03635 WP 015434384.1
 NTDB id 1177 A1552VC RS01165 WP 000422572.1
 NTDB id 1203 PSJM300 01605 AFN76401.1
 NTDB id 1019 ACIAD RS04205 WP 004922051.1
 consensus



232
239
226
228
229

logo

NTDB id 1307 DR RS10055 WP 010888596.1
 NTDB id 56709 AZKH RS03635 WP 015434384.1
 NTDB id 1177 A1552VC RS01165 WP 000422572.1
 NTDB id 1203 PSJM300 01605 AFN76401.1
 NTDB id 1019 ACIAD RS04205 WP 004922051.1
 consensus



312
319
306
308
309

logo

NTDB id 1307 DR RS10055 WP 010888596.1
 NTDB id 56709 AZKH RS03635 WP 015434384.1
 NTDB id 1177 A1552VC RS01165 WP 000422572.1
 NTDB id 1203 PSJM300 01605 AFN76401.1
 NTDB id 1019 ACIAD RS04205 WP 004922051.1
 consensus



385
386
368
376
378

logo

NTDB id 1307 DR RS10055 WP 010888596.1	GGDFGRGAAAPATPAAPAAGRSEPGRAGGFGR	418
NTDB id 56709 AZKH RS03635 WP 015434384.1	DEDEE.GA..PDKAAV.....	399
NTDB id 1177 A1552VC RS01165 WP 000422572.1	368
NTDB id 1203 PSJM300 01605 AFN76401.1	GRSFR.....	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	RRNLK.....	383
consensus		

GRDFR
BENEERGA A A A P P T P A X
AKA A A P A A G R S E P G R A G G F G R R

 non conserved
 similar
 ≥ 50% conserved