

logo

NTDB id 1307 DR RS10055 WP 010888596.1  
 NTDB id 1177 A1552VC RS01165 WP 00422572.1  
 NTDB id 1019 ACIAD RS04205 WP 004922051.1  
 NTDB id 96458 GLA29479 RS03185 WP 057970755.1  
 NTDB id 1203 PSJM300 01605 AFN76401.1  
 consensus



65  
60  
61  
80  
61

logo

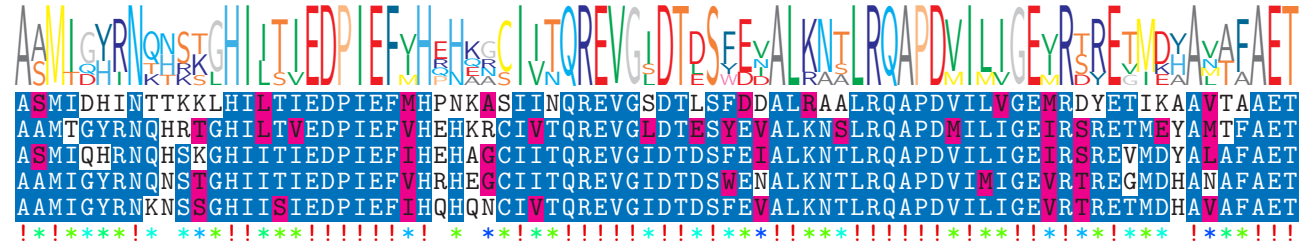
NTDB id 1307 DR RS10055 WP 010888596.1  
 NTDB id 1177 A1552VC RS01165 WP 00422572.1  
 NTDB id 1019 ACIAD RS04205 WP 004922051.1  
 NTDB id 96458 GLA29479 RS03185 WP 057970755.1  
 NTDB id 1203 PSJM300 01605 AFN76401.1  
 consensus



144  
138  
141  
159  
140

logo

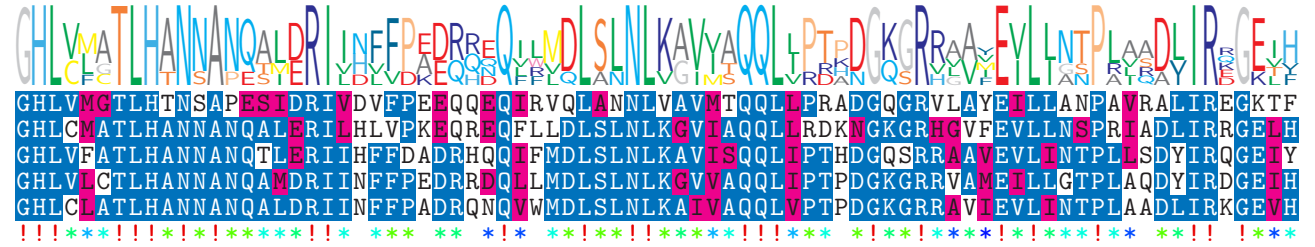
NTDB id 1307 DR RS10055 WP 010888596.1  
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 NTDB id 96458 GLA29479 RS03185 WP 057970755.1  
 NTDB id 1203 PSJM300 01605 AFN76401.1  
 consensus



224  
218  
221  
239  
220

logo

NTDB id 1307 DR RS10055 WP 010888596.1  
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 NTDB id 1019 ACIAD RS04205 WP 004922051.1  
 NTDB id 96458 GLA29479 RS03185 WP 057970755.1  
 NTDB id 1203 PSJM300 01605 AFN76401.1  
 consensus



304  
298  
301  
319  
300

logo

NTDB id 1307 DR RS10055 WP 010888596.1  
 NTDB id 1177 A1552VC RS01165 WP 00422572.1  
 NTDB id 1019 ACIAD RS04205 WP 004922051.1  
 NTDB id 96458 GLA29479 RS03185 WP 057970755.1  
 NTDB id 1203 PSJM300 01605 AFN76401.1  
 consensus



384  
368  
377  
392  
375

logo

NTDB id 1307 DR RS10055 WP 010888596.1	RGGDFGRGAAAPATPAAPAAGRSEPGRAGGFGR	418
NTDB id 1177 A1552VC RS01165 WP 000422572.1	.....	368
NTDB id 1019 ACIAD RS04205 WP 004922051.1	QRRNLK.....	383
NTDB id 96458 GLA29479 RS03185 WP 057970755.1	.....	392
NTDB id 1203 PSJM300 01605 AFN76401.1	PGRSFR.....	381
consensus		

Logo showing amino acid conservation for the sequence RGGDFGRGAAAPATPAAPAAGRSEPGRAGGFGR. The logo uses color coding: blue for non-conserved, red for similar, and green for ≥50% conserved.

- non conserved
- similar
- ≥ 50% conserved