

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 142598 LA76x RS07150 WP 051884544.1
 NTDB id 1203 PSJM300 01605 AFN76401.1
 consensus



65
60
61
80
61

logo

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 consensus



144
138
141
159
140

logo

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 consensus



224
218
221
239
220

logo

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 consensus



304
298
301
319
300

logo

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384
368
377
392
375

logo

| | | |
|---|------------------------------------|-----|
| NTDB id 1307 DR RS10055 WP 010888596.1 | RGGDFGRGAAAAPATPAAPAAGRSEPGRAGGFGR | 418 |
| NTDB id 1177 A1552VC RS01165 WP 000422572.1 | | 368 |
| NTDB id 1019 ACIAD RS04205 WP 004922051.1 | QRRNLK..... | 383 |
| NTDB id 142598 LA76x RS07150 WP 051884544.1 | | 392 |
| NTDB id 1203 PSJM300 01605 AFN76401.1 | PGRSFR..... | 381 |
| consensus | | |

Logo showing amino acid conservation across sequences. The sequence is RGGDFGRGAAAAPATPAAPAAGRSEPGRAGGFGR. The logo indicates conservation levels for each position, with the most conserved positions (R, G, A, A, P, A, A, G, R, S, E, P, G, R, A, G, G, F, G, R) showing the highest conservation.

 non conserved
 similar
 ≥ 50% conserved