

logo

```
NTDB id 25 SA RS06185 WP 000593194.1
NTDB id 599 KW2 RS05940 WP 021037268.1
NTDB id 407 SMU RS04605 WP 002262865.1
NTDB id 525 SMSK321 RS06295 WP 000705298.1
NTDB id 497 SM12261 RS05565 WP 000705318.1
NTDB id 352544 ES300 RS05785 WP 000705324.1
NTDB id 267 KZH43 RS05590 WP 000705306.1
NTDB id 226 SPD RS05990 WP 000705306.1
NTDB id 192 SPR RS05715 WP 000705306.1
NTDB id 157 SP RS06205 WP 000705306.1
consensus
```

[illegible]

logo

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NTDB id 226 SPD RS05990 WP 000705306.1
NTDB id 192 SPR RS05715 WP 000705306.1
NTDB id 157 SP RS06205 WP 000705306.1
consensus

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SFSILDDCYPWDLSEIYDAPVLLFYKGNLDLLKFKPKVAVVGSRRACSKQGAKSVEKVIQGLENELVIVSGLAKGIDTAA	158
YTIYMDSEYPPVLLKEIYQFPLLLFYKGNIKLINNMHFLAVVGARDSTSYTQQSLEFLISNDKSKYLTIVSGLAQCADAMA	159
SFSILDDLYPERLREIYNPPVLIIFYQGNIDLLKNPKLAFVGSRLAGQSQIKSVQKIVTEINLSFTIVSGLAKGIDTAS	160
SLSIDKEYPLELKNIYNPPVLLFYQGNLDLLARPKLAVVGSRRASQMGVAAVKKIITQLSKQFVITISGLARIGIDTAA	161
SFSILDDCYPWDLSEIYDAPVLLFYKGNLDLLKFKPKVAVVGSRRACSKQGAKSVEKVIQGLENELVIVSGLAKGIDTAA	162
SFSILDDCYPWDLSEIYDAPVLLFYKGNLDLLKFKPKVAVVGSRRACSKQGAKSVEKVIQGLENELVIVSGLAKGIDTAA	163
SFSILDDCYPWDLSEIYDAPVLLFYKGNLDLLKFKPKVAVVGSRRACSKQGAKSVEKVIQGLENELVIVSGLAKGIDTAA	164
SFSILDDCYPWDLSEIYDAPVLLFYKGNLDLLKFKPKVAVVGSRRACSKQGAKSVEKVIQGLENELVIVSGLAKGIDTAA	165
SFSILDDCYPWDLSEIYDAPVLLFYKGNLDLLKFKPKVAVVGSRRACSKQGAKSVEKVIQGLENELVIVSGLAKGIDTAA	166
SFSILDDCYPWDLSEIYDAPVLLFYKGNLDLLKFKPKVAVVGSRRACSKQGAKSVEKVIQGLENELVIVSGLAKGIDTAA	167
SFSILDDCYPWDLSEIYDAPVLLFYKGNLDLLKFKPKVAVVGSRRACSKQGAKSVEKVIQGLENELVIVSGLAKGIDTAA	168
SFSILDDCYPWDLSEIYDAPVLLFYKGNLDLLKFKPKVAVVGSRRACSKQGAKSVEKVIQGLENELVIVSGLAKGIDTAA	169
SFSILDDCYPWDLSEIYDAPVLLFYKGNLDLLKFKPKVAVVGSRRACSKQGAKSVEKVIQGLENELVIVSGLAKGIDTAA	170

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NTDB id 157 SP RS06205 WP 000705306.1
consensus
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	238
	229
	228
	230
	230
	230
	230
	230
	230
	230
	230

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NTDB id 226 SPD RS05990 WP 000705306.1
NTDB id 192 SPR RS05715 WP 000705306.1
NTDB id 157 SP RS06205 WP 000705306.1
consensus



☐ non conserved
☒ similar
☒ ≥50% conserved