

logo

NTDB id 32733 SGGB RS00670 WP 009853237.1

NTDB id 402 FSA28 RS09365 WP 002263443.1

NTDB id 393 SMU RS08995 WP 002263443.1

consensus

MAKTISINAGSSSLKWQLYNMPEEVLAKGLIERIGLVNGVSTVKFNGQTKSETKDTIPDHTAAVKILLDDLIHLDIISKSYDEITG

85

85

85

logo

NTDB id 32733 SGGB RS00670 WP 009853237.1

NTDB id 402 FSA28 RS09365 WP 002263443.1

NTDB id 393 SMU RS08995 WP 002263443.1

consensus

VGHRVWAGGT YFNESALVDSEEVIKKVEELALLAPLHNKANAAGIRAFKKILPDITSVVFDTAFHTTMPEVAYRYPLPNKYFTE

169

170

170

logo

NTDB id 32733 SGGB RS00670 WP 009853237.1

NTDB id 402 FSA28 RS09365 WP 002263443.1

NTDB id 393 SMU RS08995 WP 002263443.1

consensus

NQVRKYGAHGTSHYYVAHEAAKILEKPIEKLKLITVHMGNGVSLTAVDGGKSVDTSMGFTPLGGVMMGTRTGDLDPAVIPYLMND

254

255

255

logo

NTDB id 32733 SGGB RS00670 WP 009853237.1

NTDB id 402 FSA28 RS09365 WP 002263443.1

NTDB id 393 SMU RS08995 WP 002263443.1

consensus

TEDFKTPEDIRRFNNSGLLGISSELSNDMREIEAATAAGNKNTLAYNMFIDRIIKHIGAYAAMNGVDAIVFTAGIGENDAH

339

340

340

logo

NTDB id 32733 SGGB RS00670 WP 009853237.1

NTDB id 402 FSA28 RS09365 WP 002263443.1

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consensus

RSEIMKHFDWL GADIVTEKNEKRPVYGVISNAAKVKVLVIPTDEELVIARDVERLKTK

399

399

399

- non conserved
- similar
- ≥ 50% conserved