

logo

NTDB id 27580 PANA RS14530 WP 013026737.1

NTDB id 1457 ASU2 RS00455 WP 014990913.1

NTDB id 1392 A4U84 RS02730 WP 010786053.1

consensus

MSEQIQATDASCNSTKTEKINLLNLRQQMRELFVSLGKPKFRADQLMKWIYHFGEDNFDNMTNINKVLRREKLRIAE

81

85

78

logo

NTDB id 27580 PANA RS14530 WP 013026737.1

NTDB id 1457 ASU2 RS00455 WP 014990913.1

NTDB id 1392 A4U84 RS02730 WP 010786053.1

consensus

TRAPEVAEEMRSTDGTIKWAIIRVGDQLVETVYIPEGDRATLCVSSQVGCALACKFCSTAQQGFNRNLRVSEIIGQVWRASKIIGN

166

170

163

logo

NTDB id 27580 PANA RS14530 WP 013026737.1

NTDB id 1457 ASU2 RS00455 WP 014990913.1

NTDB id 1392 A4U84 RS02730 WP 010786053.1

consensus

AKVTCQRPITNVVMMGMGEPLNLRNVIPAMEIMLDDFAYGLSKRRVTLSTAGVVPALDKMREKIDVALAISLHAPNDELRLDEIM

251

255

248

logo

NTDB id 27580 PANA RS14530 WP 013026737.1

NTDB id 1457 ASU2 RS00455 WP 014990913.1

NTDB id 1392 A4U84 RS02730 WP 010786053.1

consensus

PINKKYNIETFLAAVKRYIGKSNANQGRVTIEYVLLDHVNDSTDAHELAAALKEKTPCKINLIPWNPFPPEAPYKSSNSRIDRFQ

336

340

333

logo

NTDB id 27580 PANA RS14530 WP 013026737.1

NTDB id 1457 ASU2 RS00455 WP 014990913.1

NTDB id 1392 A4U84 RS02730 WP 010786053.1

consensus

KVLMDYGFTHIVRKTRGDDIDAACGQLAGDVLDRTKRTMEKRKFGEGLAVQNH

389

393

386

- non conserved
- similar
- ≥ 50% conserved