

logo

NTDB id 162252 WG70 RS22380 WP 010104670.1
NTDB id 125 BSU 00870 NP 387968.1
NTDB id 279 KZH43 RS00140 WP 074017595.1
NTDB id 238 SPD RS00140 WP 074017595.1
NTDB id 204 SPR RS00140 WP 074017595.1
NTDB id 169 SP RS00155 WP 074017595.1
NTDB id 537 SMSK321 RS07120 WP 080550752.1
NTDB id 509 SM12261 RS00130 WP 078228442.1
consensus



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consensus

ASLIMAVLEKRAGLLLQNQDAYLKSAGGVKLDPAIDLAVAVAIASSYKDKPTNPQECFVGELGLTGEIRRVNRIEQRINEAAKL

LAMILAVLHRRHAGTIACFDQDVFNLAVGGVKITEPAADLAVLLAIHSSMRNKPLPKGLIVFGFVGLAGEIRPSPRGQERIREAAKL

VSLIMAVLEKRVGLLLLQNQDAYLKVAGGVKLDPAIDLAVIVISIASFFRDTPPNPADCFIIGEVLGTGEVRRVSRIEQRVKEAAKL

ASLIMAVLEKRAGLLLQNQDAYLKSAGGVKLDPAIDLAVAVAIASSYKDKPTNPQECFVGELGLTGEIRRVNRIEQRINEAAKL

ASLIMAVLEKRAGLLLQNQDAYLKSAGGVKLDPAIDLAVAVAIASSYKDKPTNPQECFVGELGLTGEIRRVNRIEQRINEAAKL

ASLIMAVLEKRAGLLLQNQDAYLKSAGGVKLDPAIDLAVAVAIASSYKDKPTNPQECFVGELGLTGEIRRVNRIEQRINEAAKL

ASLIMAVLEKRAGLLLQNQDAYLKSAGGVKLDPAIDLAVAVAIASSYKDKPTNPQECFVGELGLTGEIRRVNRIEQRINEAAKL

ASLIMAVLEKRAGLLLQNQDAYLKSAGGVKLDPAIDLAVAVAIASSYKDKPTNPQECFVGELGLTGEIRRVNRIEQRINEAAKL

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consensus

GFTKIYVPKNSLTGITLPKEIQVIGVTTIQEVLKKVFA

KRXITIIAANADKQPKIGLEHAERVECAISSELE

GFTTALIPKANAPKQPI.DGLRVHVERIEQAIDQIRVLE

GFKRMIIPAANLDGWTKPKGIEVIGVANVAEALRTSLGG.

GFTKIYVPKNSLTGITLPKEIQVIGVTTIQEVLKKVFA..

GFTKIYVPKNSLTGITLPKEIQVIGVTTIQEVLKKVFA..

GFTKIYVPKNSLTGITLPKEIQVIGVTTIQEVLKKVFA..

GFTKIYVPKNSLTGITLPKEIQVIGVTTIQEVLKKVFA..

GFTKIYVPKNSLTGITPPMEIQVIGVTTIQEVLKKVFA..

GFTKIYVPKNSLTGITPPKEIQVIGVTTIQEVLKKVFA..

!!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!

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453

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453

453

- X

 non conserved
- X

 similar
- X

 ≥ 50% conserved