

logo

NTDB id 133058 RY29 RS08595 WP 004193568.1
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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logo

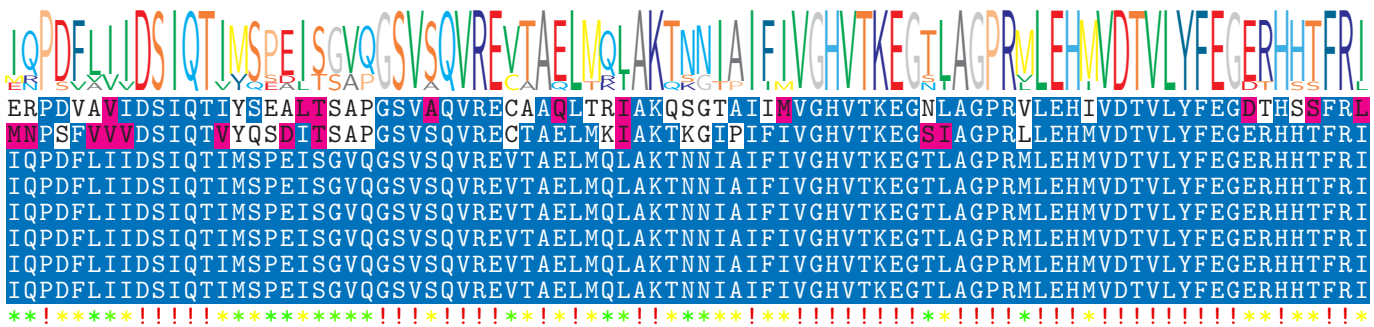
NTDB id 133058 RY29 RS08595 WP 004193568.1
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consensus



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consensus



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logo

NTDB id 133058 RY29 RS08595 WP 004193568.1
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consensus



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logo

NTDB id 133058 RY29 RS08595 WP 004193568.1	SLIMAVLEKRAGLLLQNQDAYLKSAGGVKLDEPAIDLAVAVAIASSYKDKPTNPQECFVGELGLTGEIRRVNRIEQRINEAAKLG	420
NTDB id 125 BSU 00870 NP 387968.1	AMILLAVLHRRHAGTIACFDQDDVFLNAVGGVKITEPAADLAVLLAIHSSMRNKPLPKGLIVFGFVGLAGEIRPSPRGQERIREAAKLG	420
NTDB id 279 KZH43 RS00140 WP 074017595.1	SLIMAVLEKRVGLLLLQNQDAYLKVAGGVKLDEPAIDLAIVISIIASSFRDTPPNPADCFIIGEVLGTGEVRRVSRIEQRVKEAAKLG	416
NTDB id 238 SPD RS00140 WP 074017595.1	SLIMAVLEKRAGLLLQNQDAYLKSAGGVKLDEPAIDLAIVISIIASSYKDKPTNPQECFVGELGLTGEIRRVNRIEQRINEAAKLG	416
NTDB id 204 SPR RS00140 WP 074017595.1	SLIMAVLEKRAGLLLQNQDAYLKSAGGVKLDEPAIDLAIVISIIASSYKDKPTNPQECFVGELGLTGEIRRVNRIEQRINEAAKLG	416
NTDB id 169 SP RS00155 WP 074017595.1	SLIMAVLEKRAGLLLQNQDAYLKSAGGVKLDEPAIDLAIVISIIASSYKDKPTNPQECFVGELGLTGEIRRVNRIEQRINEAAKLG	416
NTDB id 537 SMSK321 RS07120 WP 080550752.1	SLIMAVLEKRAGLLLQNQDAYLKSAGGVKLDEPAIDLAIVISIIASSYKDKPTNPQECFVGELGLTGEIRRVNRIEQRINEAAKLG	416
NTDB id 509 SM12261 RS00130 WP 078228442.1	SLIMAVLEKRAGLLLQNQDAYLKSAGGVKLDEPAIDLAIVISIIASSYKDKPTNPQECFVGELGLTGEIRRVNRIEQRINEAAKLG	416
consensus	*****!!!!*****!!!!*****!!!!*****!!!!*****!!!!*****!!!!*****!!!!*****!!!!*****!!!!	

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NTDB id 133058 RY29 RS08595 WP 004193568.1	FTKIYVPKNSLTGITLPKEIQVIGVTTIQEVLKKVFA	458
NTDB id 125 BSU 00870 NP 387968.1	FTTALIPKANAPKQPIDGLRVHHAVERIEQAIDQIRVLE	458
NTDB id 279 KZH43 RS00140 WP 074017595.1	FKRMIIPAANLDGWTKPKGIEVIGVANVAEALRTSLGG	453
NTDB id 238 SPD RS00140 WP 074017595.1	FTKIYVPKNSLTGITLPKEIQVIGVTTIQEVLKKVFA	453
NTDB id 204 SPR RS00140 WP 074017595.1	FTKIYVPKNSLTGITLPKEIQVIGVTTIQEVLKKVFA	453
NTDB id 169 SP RS00155 WP 074017595.1	FTKIYVPKNSLTGITLPKEIQVIGVTTIQEVLKKVFA	453
NTDB id 537 SMSK321 RS07120 WP 080550752.1	FTKIYVPKNSLTGITPPMEIQVIGVTTIQEVLKKVFA	453
NTDB id 509 SM12261 RS00130 WP 078228442.1	FTKIYVPKNSLTGITPPKEIQVIGVTTIQEVLKKVFA	453
consensus	!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****	

- non conserved
- similar
- ≥ 50% conserved