

logo

NTDB id 586 KW2 RS10855 WP 011836048.1
NTDB id 429 SGO RS09420 WP 012130942.1
NTDB id 500 SM12261 RS00945 WP 000249545.1
NTDB id 270 KZH43 RS09265 WP 000249564.1
NTDB id 229 SPD RS09895 WP 000249564.1
NTDB id 195 SPR RS09430 WP 000249564.1
NTDB id 160 SP RS10380 WP 000249564.1
NTDB id 584 DLJ51 RS00890 WP 019777327.1
NTDB id 1119537 EL071 RS11725 WP 115356969.1
NTDB id 385 SMU RS09035 WP 002263435.1
NTDB id 394 FSA28 RS09405 WP 002291652.1
consensus



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logo

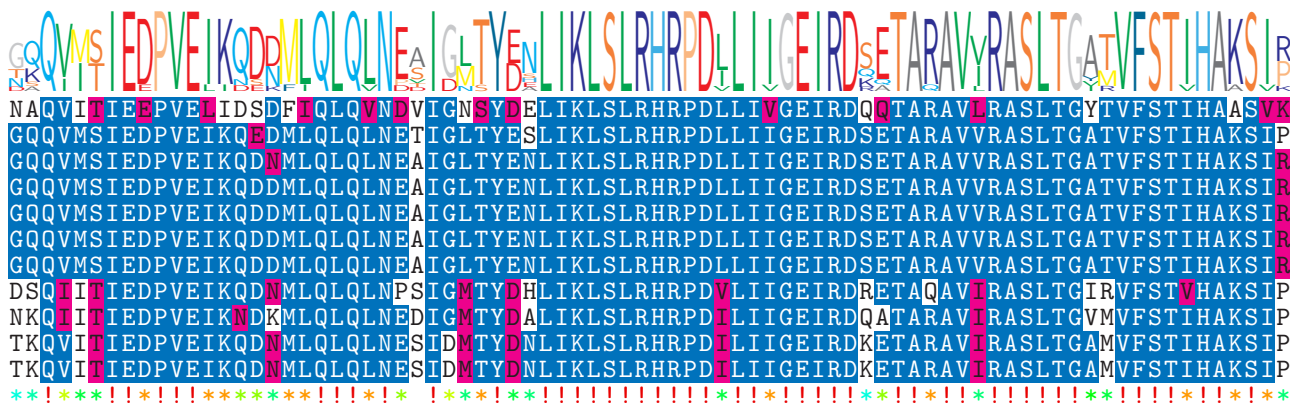
NTDB id 586 KW2 RS10855 WP 011836048.1
NTDB id 429 SGO RS09420 WP 012130942.1
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NTDB id 586 KW2 RS10855 WP 011836048.1
NTDB id 429 SGO RS09420 WP 012130942.1
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NTDB id 394 FSA28 RS09405 WP 002291652.1
consensus



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NTDB id 586 KW2 RS10855 WP 011836048.1	GVVQRLLLELGLSDWELKNGL	EA	VVYQRLI	AGKGVLDIAKSKFD	TWSPK	KWNEK	IENLYANGHITALEAEREKIS	INQASK	316
NTDB id 429 SGO RS09420 WP 012130942.1	GVYERLLELGVSEEEELK	IVLQGT	CYQRLIGGGG	VIDFASD	NYQEHE	PTVWN	QQIDQLLAAGHITHPEQABAEKIRNQQAKT		317
NTDB id 500 SM12261 RS00945 WP 000249545.1	GVYERLLELGVSEEEELAV	VLQGV	CYQRLIGGGG	VIDFANK	DYQEHQ	STSWNA	QIDQLLKDGHITS	LQAETEKISYS....	313
NTDB id 270 KZH43 RS09265 WP 000249564.1	GVYERLLELGVSEEEELAV	VLQGV	CYQRLIGGGG	VIDFASR	DYQEHQA	AKWNE	QIDQLLKDGHITS	LQAETEKISYS....	313
NTDB id 229 SPD RS09895 WP 000249564.1	GVYERLLELGVSEEEELAV	VLQGV	CYQRLIGGGG	VIDFASR	DYQEHQA	AKWNE	QIDQLLKDGHITS	LQAETEKISYS....	313
NTDB id 195 SPR RS09430 WP 000249564.1	GVYERLLELGVSEEEELAV	VLQGV	CYQRLIGGGG	VIDFASR	DYQEHQA	AKWNE	QIDQLLKDGHITS	LQAETEKISYS....	313
NTDB id 160 SP RS10380 WP 000249564.1	GVYERLLELGVSEEEELAV	VLQGV	CYQRLIGGGG	VIDFASR	DYQEHQA	AKWNE	QIDQLLKDGHITS	LQAETEKISYS....	313
NTDB id 584 DLJ51 RS00890 WP 019777327.1	GVYARLLELGVSDLEL	DNCLAG	IAYQRLIGGGG	LIDFAQDHF	QNHQA	ECWNEK	IDQLLAAGHIT	EEEA	315
NTDB id 1119537 EL071 RS11725 WP 115356969.1	GVYDRLLELGVNYQEL	ENSLK	LAYQRLIGGG	LIDFET	SNFKKH	SSDKWN	RQVDILAEE	GHISKKQA	318
NTDB id 385 SMU RS09035 WP 002263435.1	GVYARMLELGTSSSEEL	QNALRG	IYVYQRLIGGGG	LIDFAKGD	YQNHSA	KKWNK	QIYDLFTAGY	ITEGEAQAEE	313
NTDB id 394 FSA28 RS09405 WP 002291652.1	GVYARMLELGTSSSEEL	QNALCG	IYVYQRLIGGGG	LIDFAKGD	YQNHSA	KKWNK	QIYDLFTAGY	ITEGEAQAEE	313
consensus	!!**!*!!**!!**!!	**!******!!!!	!!**!******!!!!	!!**!******!!!!	!!**!******!!!!	!!**!******!!!!	!!**!******!!!!	!!**!******!!!!	

logo

NTDB id 586 KW2 RS10855 WP 011836048.1	IDTTD	GESFE	326
NTDB id 429 SGO RS09420 WP 012130942.1	SY.....		319
NTDB id 500 SM12261 RS00945 WP 000249545.1			313
NTDB id 270 KZH43 RS09265 WP 000249564.1			313
NTDB id 229 SPD RS09895 WP 000249564.1			313
NTDB id 195 SPR RS09430 WP 000249564.1			313
NTDB id 160 SP RS10380 WP 000249564.1			313
NTDB id 584 DLJ51 RS00890 WP 019777327.1			315
NTDB id 1119537 EL071 RS11725 WP 115356969.1	SSPTF.....		323
NTDB id 385 SMU RS09035 WP 002263435.1			313
NTDB id 394 FSA28 RS09405 WP 002291652.1			313
consensus			

- ☒ non conserved
- ☒ similar
- ☒ ≥ 50% conserved