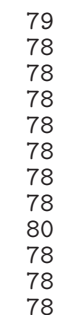


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NTDB id 586 KW2 RS10855 WP 011836048.1
NTDB id 429 SGO RS09420 WP 012130942.1
NTDB id 500 SM12261 RS00945 WP 000249545.1
NTDB id 270 KZH43 RS09265 WP 000249564.1
NTDB id 229 SPD RS09895 WP 000249564.1
NTDB id 195 SPR RS09430 WP 000249564.1
NTDB id 160 SP RS10380 WP 000249564.1
NTDB id 584 DLJ51 RS00890 WP 019777327.1
NTDB id 40020 SGBBAA2069 RS00755 WP 012961327.1
NTDB id 385 SMU RS09035 WP 002263435.1
NTDB id 394 FSA28 RS09405 WP 002291652.1
consensus

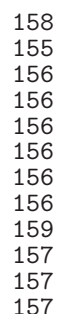
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NTDB id 195 SPR RS09430 WP 000249564.1
NTDB id 160 SP RS10380 WP 000249564.1
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NTDB id 40020 SGGBA2069 RS00755 WP 012961327.1
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NTDB id 394 FSA28 RS09405 WP 002291652.1
consensus

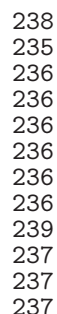
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NTDB id 385 SMU RS09035 WP 002263435.1
NTDB id 394 FSA28 RS09405 WP 002291652.1
consensus

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logo

NTDB id 617	LCA	RS06480	WP 011375001.1	QGVVARLLDLGISELQLRACLTGIAVQSLVTKDDVVQAH...YELLTGSDYFSKKEGG.....CDV.....	296
NTDB id 586	KW2	RS10855	WP 011836048.1	KGVVQRLLLELGLSDWELKNGLEAVVYQRLTAGKGVLDIAKSKFDTWSPKKWNEKTENLYANGHLETAIEAEREKISINQAS	315
NTDB id 429	SG0	RS09420	WP 012130942.1	PGVYERLLELGVSEEEELKIVLQGICYQRLIGGGGVIDFASDNYQEHEPTVWNQQIDQLLAAGHIEPEQAEAEKIRNQQAK	316
NTDB id 500	SM12261	RS00945	WP 000249545.1	RGVYERLLELGVSEEEELAVVLQGVICYQRLIGGGGVIDFANKDYQEHQSTSWNAQIDQLLKDGHITSLQAETEKISYS...	313
NTDB id 270	KZH43	RS09265	WP 000249564.1	RGVYERLLELGVSEEEELAVVLQGVICYQRLIGGGGVIDFASRDYQEHQAQWNEQIDQLLKDGHITSLQAETEKISYS...	313
NTDB id 229	SPD	RS09895	WP 000249564.1	RGVYERLLELGVSEEEELAVVLQGVICYQRLIGGGGVIDFASRDYQEHQAQWNEQIDQLLKDGHITSLQAETEKISYS...	313
NTDB id 195	SPR	RS09430	WP 000249564.1	RGVYERLLELGVSEEEELAVVLQGVICYQRLIGGGGVIDFASRDYQEHQAQWNEQIDQLLKDGHITSLQAETEKISYS...	313
NTDB id 160	SP	RS10380	WP 000249564.1	RGVYERLLELGVSEEEELAVVLQGVICYQRLIGGGGVIDFASRDYQEHQAQWNEQIDQLLKDGHITSLQAETEKISYS...	313
NTDB id 584	DLJ51	RS00890	WP 019777327.1	PGVYARLLELGVSDDELNCLAGIAYQRLIGGGGLIDFAQDHFQNHQAECWNEKIDQLLAAGHLEEEBAAAEEKIKD....	315
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NTDB id 385	SMU	RS09035	WP 002263435.1	PGVYARLLELGVSDDELNCLRGIVYQRLIGGGGVIDFAKGDYQNHSAKKWNKQIYDLFTAGYITEGEAQAEETIED....	313
NTDB id 394	FSA28	RS09405	WP 002291652.1	PGVYARLLELGVSDDELNCLCGIVYQRLIGGGGVIDFAKGDYQNHSAKKWNKQIYDLFTAGYITEGEAQAEETIED....	313
consensus				!!*!*!*!!*!***!*!*!!*!*!***************	

logo

NTDB id 617	LCA	RS06480	WP 011375001.1	KIDTTDGSFE	296
NTDB id 586	KW2	RS10855	WP 011836048.1	KIDTTDGSFE	326
NTDB id 429	SG0	RS09420	WP 012130942.1	TSY.....	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
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NTDB id 40020	SGGBAA2069	RS00755	WP 012961327.1		313
NTDB id 385	SMU	RS09035	WP 002263435.1		313
NTDB id 394	FSA28	RS09405	WP 002291652.1		313
consensus					

- X non conserved
- X similar
- X ≥ 50% conserved