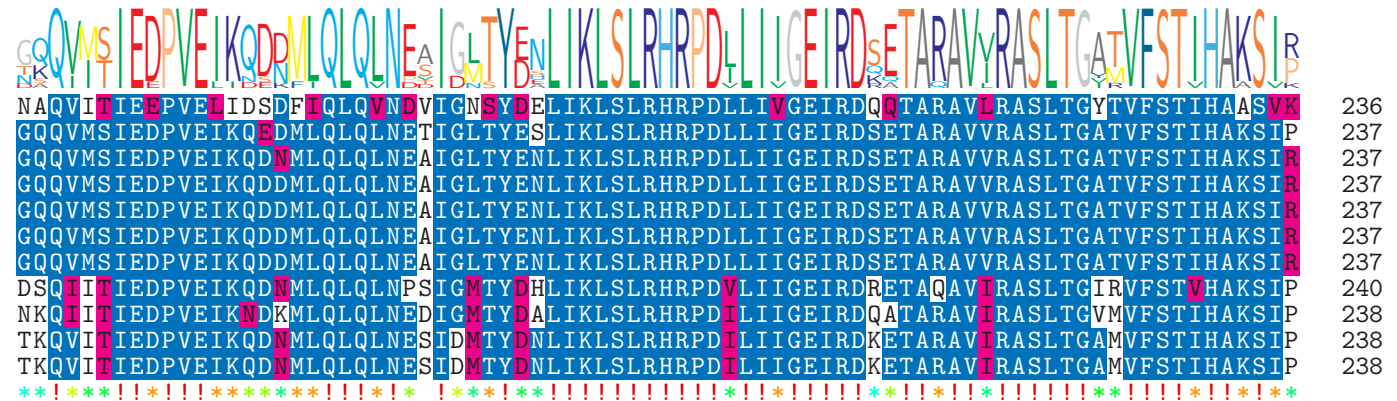
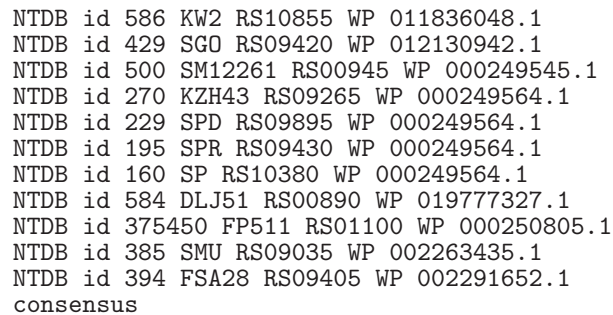
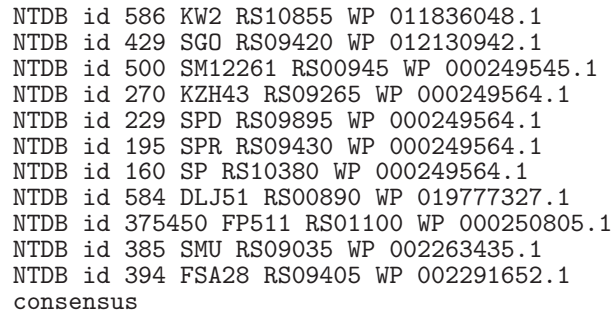


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NTDB id 429 SG0 RS09420 WP 012130942.1
NTDB id 500 SM12261 RS00945 WP 000249545.1
NTDB id 270 KZH43 RS09265 WP 000249564.1
NTDB id 229 SPD RS09895 WP 000249564.1
NTDB id 195 SPR RS09430 WP 000249564.1
NTDB id 160 SP RS10380 WP 000249564.1
NTDB id 584 DLJ51 RS00890 WP 019777327.1
NTDB id 375450 FP511 RS01100 WP 000250805.1
NTDB id 385 SMU RS09035 WP 002263435.1
NTDB id 394 FSA28 RS09405 WP 002291652.1
consensus

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NTDB id	586	KW2 RS10855 WP	011836048.1	G VVQRLLELGLSDWELKNGLEAVVVYQRLLAGKGVLDI A AKSKFDTWSPKKWNEKIENLYANCHLTIAEAEREKISINQASK	316
NTDB id	429	S <i>G</i> O RS09420 WP	012130942.1	G VYERLLELGVSSEELKLIVLQGICYQLRIGGGGVIDFASDNYQEHEPTVWNQQIDQLLAAGHIHPEQAEEA EKIRNQQAKT	317
NTDB id	500	SM12261 RS00945 WP	000249545.1	G VYERLLELGVSSEELA VVLQGVVCYQRLIGGGGI VDFA NKD DYQEHHSTSWNAQIDQLLKDGHITS LQAETEKISYS....	313
NTDB id	270	KZH43 RS09265 WP	000249564.1	G VYERLLELGVSSEELA VVLQGVCYQRLIGGGGI VDFA SRDYQEHAQA KWNEQIDQLLDGHITSLQAETEKISYS....	313
NTDB id	229	SPD RS09895 WP	000249564.1	G VYERLLELGVSSEELA VVLQGVVCYQRLIGGGGI VDFA SRDYQEHAQA KWNEQIDQLLDGHITSLQAETEKISYS....	313
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NTDB id	160	SP RS10380 WP	000249564.1	G VYERLLELGVSSEELA VVLQGVVCYQRLIGGGGI VDFA SRDYQEHAQA KWNEQIDQLLDGHITSLQAETEKISYS....	313
NTDB id	584	DLJ51 RS00890 WP	019777327.1	G VYAARILELVGSQDELDNCLAGIAYQRLIGGGGLIDFAQDHFNQHQAECWNEKIDQLLAAGHLITEE EAAAAEKIKD.....	315
NTDB id	375450	FP511 RS01100 WP	000250805.1	G VYDRRIELGVNYQSLENLKLAIYQRLIGGGSLIDFETGNFKKHSSDKWRNQVIDILAEEGHISKKAQVEKIIPQUETTE	318
NTDB id	385	SMU RS09035 WP	002263435.1	G VYARMLELGTSSSELQNALRGVIYQRLIGGGIIDFAKGDYQNHSAKKWNKKIQIDLFTAGYTEGEAQAAEIED.....	313
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consensus				!!**!*!!!!!*****!! **!*!!!!!!!****!!*** ***** *!!!!!!!!*****!*	

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NTDB id	586	KW2	RS10855	WP 011836048.1	IDTTDGESFE	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
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NTDB id	394	FSA28	RS09405	WP 002291652.1		313
consensus						

- X non conserved
- X similar
- X $\geq 50\%$ conserved