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NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
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consensus	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0

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NTDB id 1293	VP RS12240	WP 005479695.1	MHSNLSLILRQKGLLTFSQEESLIEQVKASGISMPEALLSSGFFTSSELAEHLSSIFGLSQPELSQYEE	69
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NTDB id 1110	NGFG RS09235	WP 003689817.1	...MSVGLLR....IL....VQNNQ.VVTVEQAEHYYNESQAGKEVLPMFLSDGVISPKSLAALIARVFSYSILDLRHYPR	68
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consensus			* * * * *	

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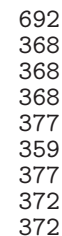
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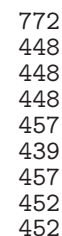
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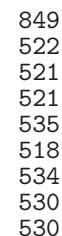
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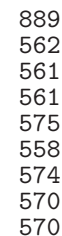
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consensus

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 non conserved
 similar
 $\geq 50\%$ conserved