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NTDB id 1028 TT RS08215 WP 011173991.1	MSXLSIGDRRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 229254 CA930 RS07410 WP 027227022.1		0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	RLTPEERKRLAALPAFALDEEAGVVRVAFLNPLDPLSTSELEEDLGHDYEPYQATIRKAYAAAKHYPELGLLEYYPSGK	158
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consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	LEMLRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSR	317
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NTDB id 1251 GC085 RS07725 WP 011946523.1	MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVKNKILSAEQIALTAAQNFQVPMMDINC	74
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQALEKAKKFKQDTPYPLIDNFSISPLTIAEIIISLEFGEPLLDLGV	74
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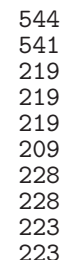
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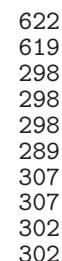
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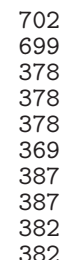
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consensus



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consensus



889
887
562
561
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558
574
574
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[X] non conserved
[X] similar
[X] ≥ 50% conserved