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NTDB id 1306 DR RS10060 WP 027479822.1	MSXLSIGDRRLGAALLDQGYLGNDLQRALERHSEVGGRLADVLDIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	79
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
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NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
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consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLIYVPSGA	157
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
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[illegible]

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consensus

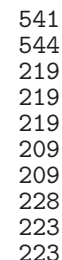
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YDYAN	C	Q	Q	L	G	R	E	L	I	T	R	Y	D	A	L	P	I	A	Q	G	N	L	L	L	L	A	V	S	D	P	T	L	L	Q	A	E	E	F	R	F	A	T	G	...	L	Q	V	E	L	A	L	A	D	H	R	A	L	Q	A	A	I	R	R	L	Y	G	R	141						
YEYAS	C	Q	Q	L	G	R	E	L	I	T	R	H	N	A	L	P	L	H	R	T	P	S	T	L	L	L	A	V	A	D	P	T	N	Q	Q	A	E	D	D	F	R	F	A	T	G	...	L	Q	V	E	L	V	L	A	D	F	R	E	L	S	T	A	I	R	R	L	Y	G	R	141				
YEYAS	C	Q	T	L	G	L	R	D	L	I	T	R	H	N	A	L	P	L	N	T	S	T	L	L	L	A	V	A	D	P	T	N	Q	A	E	D	D	F	R	F	A	T	G	...	L	Q	V	E	L	V	L	A	D	F	R	E	L	S	T	A	A	I	R	R	L	Y	G	R	141					
YPRHR	V	M	G	V	L	T	E	Q	M	V	E	F	H	C	V	P	V	F	R	G	D	K	V	F	F	A	V	S	D	P	T	Q	M	P	Q	I	Q	K	T	V	S	A	A	G	...	I	E	V	E	L	V	I	V	E	D	D	Q	L	A	G	L	D	V	W	G	S	R	140						
YPRHR	V	M	G	V	L	T	E	Q	M	V	E	F	H	C	V	P	V	F	R	G	D	K	V	F	F	A	V	S	D	P	T	Q	M	P	Q	I	Q	K	T	V	S	A	A	G	...	I	E	V	E	L	V	I	V	E	D	D	Q	L	A	G	L	D	V	W	G	S	R	140						
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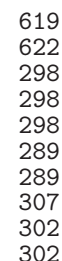
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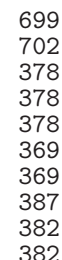
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consensus



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782
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458
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logo

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consensus



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527
527
543
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539

logo

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consensus



887
889
562
561
561
558
558
574
570
570

[X] non conserved
[X] similar
[X] ≥ 50% conserved