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NTDB id 1306 DR RS10060 WP 027479822.1	MSXLSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLDIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	79
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
NTDB id 360543 FAZ30 RS08540 WP 205676668.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	157
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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NTDB id 1306 DR RS10060 WP 027479822.1	SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
NTDB id 1028 TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKGWISREALEEALVEQKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA	237
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[illegible]

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DPPDPGAPIL	P	E	D	I	C	R	R	Y	G	F	P	H	R	L	E	G	N	R	L	V	L	M	K	D	P	R	N	I	L	A	D	D	V	R	L	A	L	K	R	G	L	N	Y	E	V	A	P	A	V	A	T	E	A	A	I	K	L	I	E	R	F	Y	G	477					
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	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	141	
Y EYAS	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	141
Y EYAS	C	Q	T	L	G	R	L	I	T	R	H	N	A	L	P	L	N	T	S	T	S	T	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	I	R	R	L	Y	G	141			
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YPRHRVLMGV	L	T																																																																			

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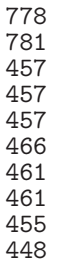
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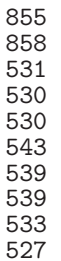
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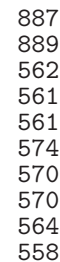
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- X non conserved
- X similar
- X > 50% conserved