

logo

NTDB id 1306 DR RS10060 WP 027479822.1	MSXLSIGDRLGAALLDQGYLGDNDLQRALERHSEVGGRLADVLDIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	79
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	80
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
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NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIYVPSGA	157
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDRTLSEEEVDTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..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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NPPDPKVAL	M	P	E	A	T	A	R	I	V	V	P	V	P	L	R	L	Q	N	S	L	V	A	M	K	D	P	R	N	F	A	L	D	D	L	K	L	I	T	G	...	K	E	L	P	A	V	A	M	A	E	K	D	I	I	R	L	T	E	R	F	Y	G	E	472										
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	T	K	L	I	E	R	F	Y	G	K	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	F	R	F	A	T	G	...	L	Q	V	E	L	A	D	A	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	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	A	D	F	R	E	L	S	T	A	A	I	R	R	L	Y	G	R	141						
YPRHR	V	L	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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I	D	V	G	T	I	P	V	N	L	V	N	E	K	L	I	K	R	H	A	M	V	P	L	F	S	R	G	T	N	L	Y	L	A	T	D	D	P	S	K	Q	A	S	L	K	E	I	Q	F	H	T	G	...	L	N	T	H	A	I	V	E	T	D	K	L	S	A	L	I	D	N	L	T	A	149
F	D	P	A	L	F	L	K	D	K	I	D	E	K	L	I	Q	K	Y	R	I	M	P	L	V	H	R	G	H	V	L	Y	V	A	T	S	P	T	N	I	E	A	M	D	A	I	R	F	N	S	K	...	L																						

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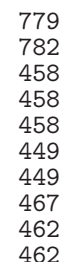
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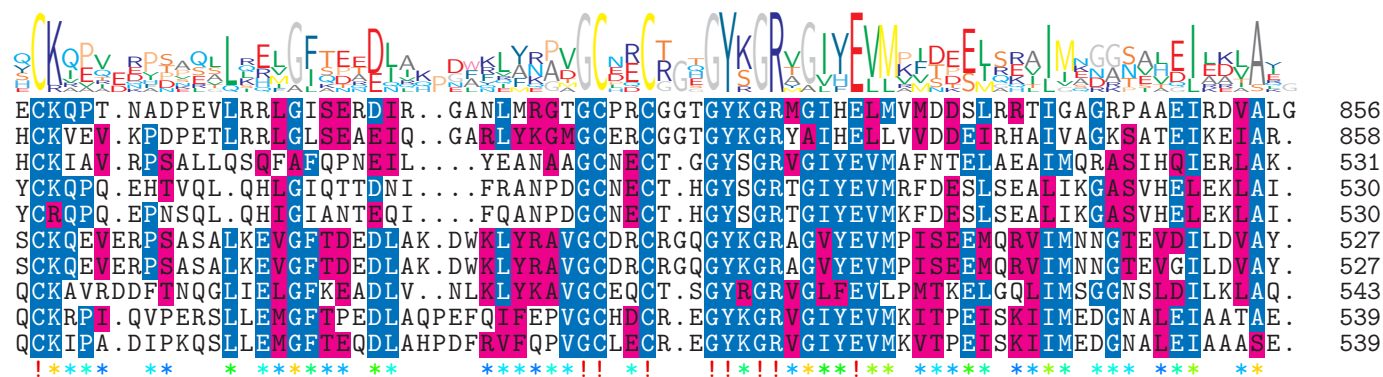
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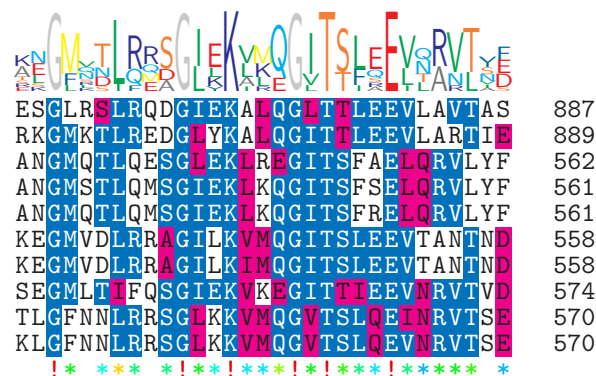
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```



-  non conserved
-  similar
-  > 50% conserved