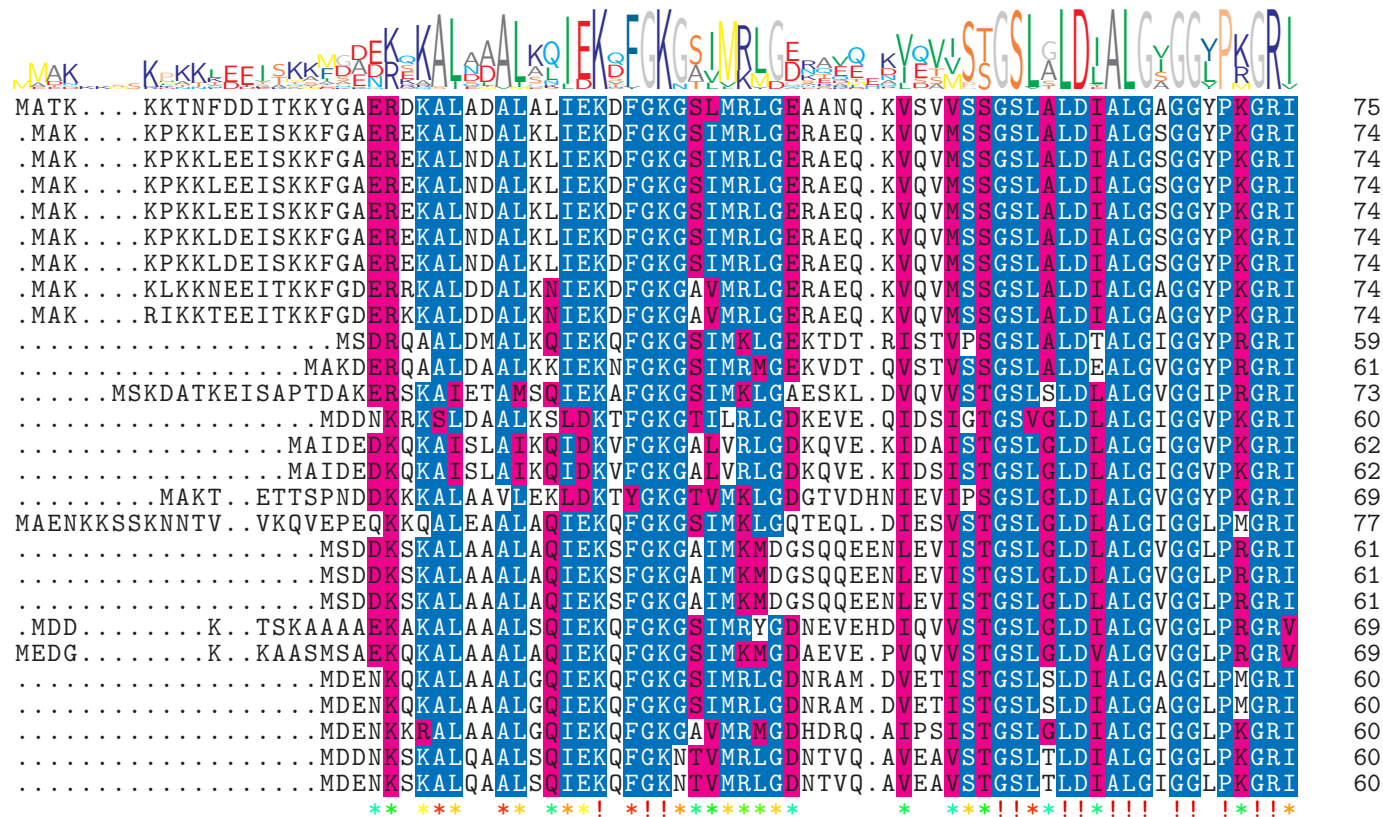


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logo

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NTDB id 1427 RS RS02760 WP 011000510.1	VR	L	D	I	R	R	I	G	S	.	.	.	.	.	.	.	.	.	.	I	.	K	K	G	.	D	E	V	V	G	N	E	T	K	V	K	V	V	K	N	K	V	A	P	P	F	R	E	A	I	F	D	I	L	Y	G	A	G	V	S	R	E	G	E	I	I	D	L	G	V	E	A	K	V	V	E	K	S	G	A	W	Y	S	Y	N	G	303
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NTDB id 1142 GTF74 RS11550 WP 000344154.1	VR	L	D	I	R	R	T	G	A	.	.	.	.	.	.	.	.	.	.	I	.	K	E	G	.	E	E	V	V	G	N	E	T	R	I	K	V	V	K	N	K	I	A	A	P	F	K	E	A	N	T	Q	I	M	Y	G	Q	G	F	N	R	E	G	E	L	I	D	L	G	V	K	H	K	M	V	E	K	S	G	A	W	Y	S	Y	N	G	294
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NTDB id 1022 ACIAD RS06390 WP 004925639.1	VR	L	D	I	R	R	I	G	Q	.	.	.	.	.	.	.	.	.	.	V	.	K	E	G	.	D	E	I	I	G	S	E	T	K	V	K	V	V	K	N	K	M	A	P	P	F	R	E	A	I	F	Q	I	L	Y	G	K	G	V	N	Q	L	G	E	L	V	D	L	A	V	Q	Q	N	I	V	Q	K	A	G	A	W	Y	S	Y	Q	G	294
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

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Sequence	Score
EKIGQGA EKAKNYLKEHQDV FDEIDHKVRAAHGLDDSEVAE..TEEETTASKTAKAKKKEEKAVETEEIELELED...	387
EKIGQGSENAKKYLA EHP E I FDEIDKQVRSKFG LIDGEEVSE..QDTENKKDE...PKKEEA..VNEEVPLDLGDELEI	384
EKIGQGSENAKKYLA EHP E I FDEIDKQVRSKFG LIDGEEVSE..QDTENKKDE...PKKEEA..VNEEVPLDLGDELEI	384
EKIGQGSENAKKYLA EHP E I FDEIDKQVRSKFG LIDGEEVSE..QDTENKKDE...PKKEEA..VNEEVPLDLGDELEI	384
EKIGQGSENAKKYLA EHP E I FDEIDKQVRSKFG LIDGEEVSE..QDTENKKDE...PKKEEA..VNEEVPLDLGDELEI	384
EKIGQGSENAKKYLA DNPEVFAEIDHQVRVQFGLIDGEEAAE.....SKKDE...VAQVDS..VNEEVPLDLGDELEI	380
EKIGQGSENAKKYLA DNPEI FDEIDHQVRVQFGLIDGEEIAE.....SKKDE...AAQVDS..VNEEVPLDLGDELEI	380
EKIGQGSENAKRYLA DHPQLFDEIDRKVRVKFGLLEESEES..AMAVASE.....ETD...TDDLALDLNGIEI	376
EKIGQGSENAKKFLADNPEI FDDIDHKVRVQYGLIEEDNTEE..KQSSKEK.....ETDEK..ADKNLVLELDDTIEL	381
ERLGGGR ENAKQFLKENKDIMLMIQEQIREHYGLDNGGVVQQ..QAETQEEL.EFE.E.....	348
ERIGGGRENAKNYLADHEDVEDEVRLKVR AAYGISDVPEEDL..PTTEDE.QI.NILPDDST.....EE.....	355
ERIGGGKEKTIAYIAERPMEQEITDRVM AAI RAGNAGEAPALAPAPAAPEAA.EA.....	363
KKLGGGRENKAF LKENPEIADEITKAIQNSMGIEGMIS..GSE...DDEGE.E.....	343
KKLGGGRENAKALLKEDKALADEITLKIKESIGSNEEIM..PLP...DEPLE.EME.....	347
KKLGGGRENAKALLKEDKALADEITLKIKESIGSNEEIM..PLP...DEPLE.EME.....	347
TKLGGGRD TVKELIKDNPELAEELEAKIREKLQAK.....	343
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AKIGQKDNVRVWLKENPEISDEIDAKIRALNGVMHIT..E.....GTQDE.TDGER.....PEE.....	348
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AKIGQKDNVRVWLKENPEISDEIDAKIRALNGVMHIT..E.....GTQDE.TDGER.....PEE.....	348
NRIGQKDNVREYLKEHKEMAEIENKVR ENQGI VSRAA..T.FPASEAEDGE.....	353
ERIGGGRDNCREFLR ENAELARIENKVR EHLGVTPMGA..V.TLAEVEED.....	352
DKIGQKANAACKYLKENPEIAKTLDDKKLREMLLNPNMQ..LIAETSSAADV.EFGAV.....PEEF.....	354
DKIGQKANAACKYLKENPEIAKTLDDKKLREMLLNPNMQ..LIAETSSAADV.EFGAV.....PEEF.....	354
SKIGQKANAACKYLEENPEIGQEIEQQIRAKLLVTSANT..KATPVAEDLADA.DI.....	347
NKIGQGNNVIRYLEENPQISTEIEAVIREQLLTKASDQ..TAA.HDETEEEP.DLL.....ES.....	349
NKIGQGNNVIRYFEENNQIAEIERNI REQLLTGTNG..AVQ.I.EDEEEP.DLL.....LES.....	349

logo

				E.I.E.E	
NTDB id 600 KW2 RS01765 WP 011675413.1		387			
NTDB id 269 KZH43 RS08665 WP 001085462.1	EIEE	388			
NTDB id 228 SPD RS09265 WP 001085462.1	EIEE	388			
NTDB id 194 SPR RS08825 WP 001085462.1	EIEE	388			
NTDB id 159 SP RS09750 WP 001085462.1	EIEE	388			
NTDB id 527 SMSK321 RS11065 WP 001085454.1	EIEE	384			
NTDB id 499 SM12261 RS08315 WP 078228268.1	EIEE	384			
NTDB id 453 Spy49 1753c ACI62003.1	ED..	378			
NTDB id 408 SMU RS09530 WP 002262392.1	ED..	383			
NTDB id 115 BSU 16940 NP 389576.2		348			
NTDB id 625 LCA RS02525 WP 011374196.1		355			
NTDB id 1313 DR RS12030 WP 010888966.1		363			
NTDB id 1245 Cj1673c YP 002345041.1		343			
NTDB id 1214 NCTC11637 00359 SQJ03206.1		347			
NTDB id 1211 C694 RS00795 WP 000963128.1		347			
NTDB id 1334 RAOC RS04870 WP 004918236.1		343			
NTDB id 1389 A4U84 RS07560 WP 021112712.1		372			
NTDB id 1130 NGFG RS03960 WP 003688695.1		348			
NTDB id 1124 NGFG RS03960 WP 003688695.1		348			
NTDB id 1123 OK783 RS03845 WP 003688695.1		348			
NTDB id 143789 XA16 RS06660 WP 003812760.1		353			
NTDB id 1427 RS RS02760 WP 011000510.1		352			
NTDB id 1165 A1552VC RS01565 WP 000344154.1		354			
NTDB id 1142 GTF74 RS11550 WP 000344154.1		354			
NTDB id 1205 PSJM300 05545 AFN77183.1		347			
NTDB id 1022 ACIAD RS06390 WP 004925639.1		349			
NTDB id 1074 ABD1 RS10195 WP 000344167.1		349			
consensus					

-  non conserved
-  similar
-  ≥ 50% conserved