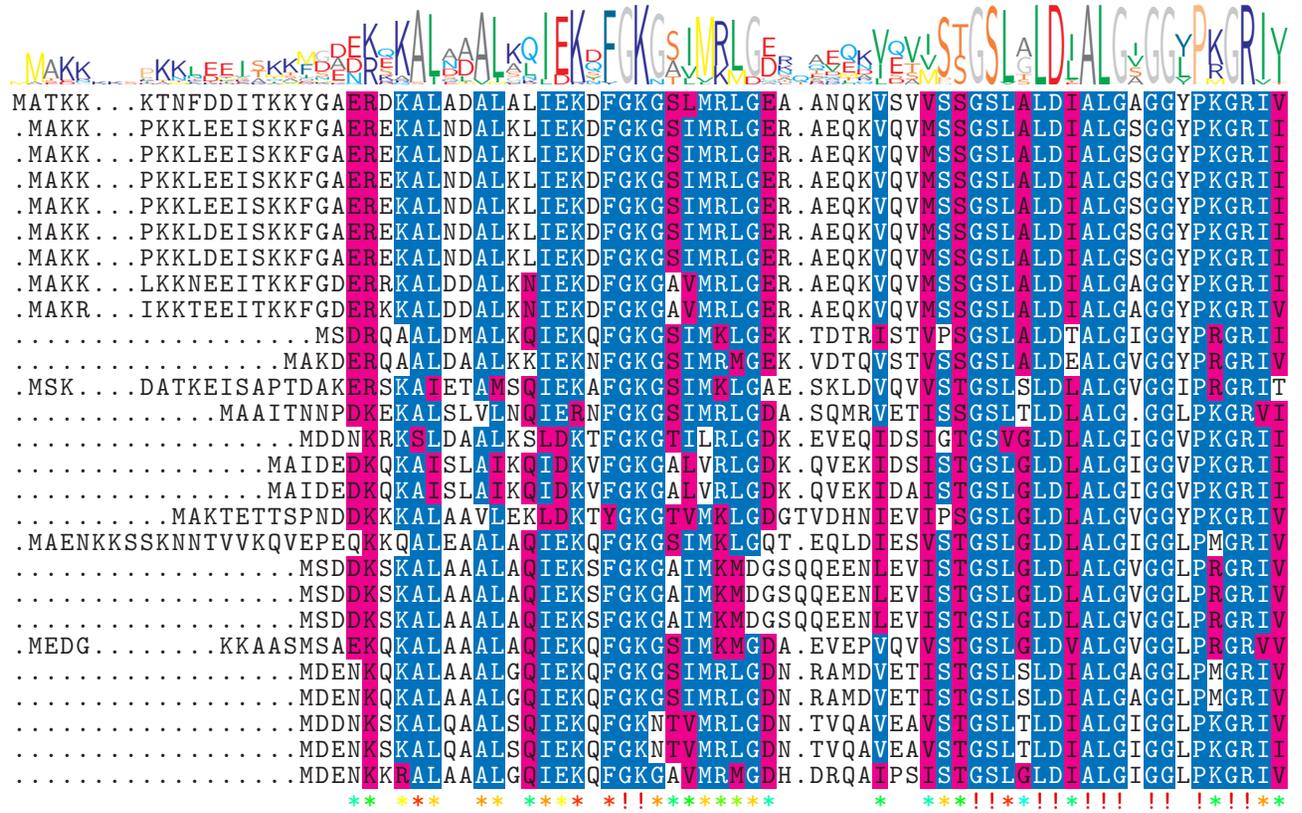


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NTDB id 600 KW2 RS01765 WP 011675413.1
 NTDB id 269 KZH43 RS08665 WP 001085462.1
 NTDB id 228 SPD RS09265 WP 001085462.1
 NTDB id 194 SPR RS08825 WP 001085462.1
 NTDB id 159 SP RS09750 WP 001085462.1
 NTDB id 499 SM12261 RS08315 WP 078228268.1
 NTDB id 527 SMSK321 RS11065 WP 001085454.1
 NTDB id 453 Spy49 1753c ACI62003.1
 NTDB id 408 SMU RS09530 WP 002262392.1
 NTDB id 115 BSU 16940 NP 389576.2
 NTDB id 625 LCA RS02525 WP 011374196.1
 NTDB id 1313 DR RS12030 WP 010888966.1
 NTDB id 54755 PLE7327 RS03840 WP 015142548.1
 NTDB id 1245 Cj1673c YP 002345041.1
 NTDB id 1211 C694 RS00795 WP 000963128.1
 NTDB id 1214 NCTC11637 00359 SQJ03206.1
 NTDB id 1334 RA0C RS04870 WP 004918236.1
 NTDB id 1389 A4U84 RS07560 WP 021112712.1
 NTDB id 1130 NGFG RS03960 WP 003688695.1
 NTDB id 1124 NGFG RS03960 WP 003688695.1
 NTDB id 1123 OK783 RS03845 WP 003688695.1
 NTDB id 1427 RS RS02760 WP 011000510.1
 NTDB id 1165 A1552VC RS01565 WP 000344154.1
 NTDB id 1142 GTF74 RS11550 WP 000344154.1
 NTDB id 1022 ACIAD RS06390 WP 004925639.1
 NTDB id 1074 ABD1 RS10195 WP 000344167.1
 NTDB id 1205 PSJM300 05545 AFN77183.1
 consensus



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NTDB id 600 KW2 RS01765 WP 011675413.1	KIGQGAEKAKNYLKEHQDVDFEIDHKVRAAHCLDDSEVAET EETTASKTAKAKKKEEKAVETEEIELELED . .	387
NTDB id 269 KZH43 RS08665 WP 001085462.1	KIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFCIDGEEVSEQ DTENKKDE PKKEEA . . VNEEVPLDLGDEL	382
NTDB id 228 SPD RS09265 WP 001085462.1	KIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFCIDGEEVSEQ DTENKKDE PKKEEA . . VNEEVPLDLGDEL	382
NTDB id 194 SPR RS08825 WP 001085462.1	KIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFCIDGEEVSEQ DTENKKDE PKKEEA . . VNEEVPLDLGDEL	382
NTDB id 159 SP RS09750 WP 001085462.1	KIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFCIDGEEVSEQ DTENKKDE PKKEEA . . VNEEVPLDLGDEL	382
NTDB id 499 SM12261 RS08315 WP 078228268.1	KIGQGSSENAKKYLADNPEIFDEIDHQVVRVQFCIDGEEIAES KKDE AAQVDS . . VNEEVTLDLGDEL	378
NTDB id 527 SMSK321 RS11065 WP 001085454.1	KIGQGSSENAKKYLADNPEVFAEIDHQVVRVQFCIDGEEAAES KKDE VAQVDS . . VNEEVTLDLGDDEL	378
NTDB id 453 Spy49 1753c ACI62003.1	KIGQGSSENAKRYLADHPQLFDEIDRKVRVKFCILEESEEEESA MAVASE E TDDLALDLNNGI	374
NTDB id 408 SMU RS09530 WP 002262392.1	KIGQGSSENAKKFLADNPEIFDDIDHKVVRVQFCILEEDNTEEK QSSKEK ETDEK ADKNLVLELDDTI	379
NTDB id 115 BSU 16940 NP 389576.2	RIGQGRENAAKQFLKENKDIMLMIQEQRREHYGLDNNGVVQQQ AEETQEEL . EFE . E	348
NTDB id 625 LCA RS02525 WP 011374196.1	RIGQGRENAKNYLADHEDVEDEVRLKVRVAAAYGISDVPEEDLP TTEDE . QI . NILPDDST EE	355
NTDB id 1313 DR RS12030 WP 010888966.1	RIGQCKEKTIAIYIAERPEMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPEAAE . A	363
NTDB id 54755 PLE7327 RS03840 WP 015142548.1	NIAGGRDNAAVKYLEENPEIANKVEQQVREKLDI GAVSFA SSTS . . DDE . EISLEDE	353
NTDB id 1245 Cj1673c YP 002345041.1	KIGQGRENSKAFLKENPEIADAITKATQNSMCI EGMISG SEDD . EGE . E	343
NTDB id 1211 C694 RS00795 WP 000963128.1	KIGQGRENAKALLKEDKALADEITLKIRESICSNEEIMP LPDE . . PLE . EME	347
NTDB id 1214 NCTC11637 00359 SQJ03206.1	KIGQGRENAKALLKEDKALADEITLKIRESICSNEEIMP LPDE . . PLE . EME	347
NTDB id 1334 RAOC RS04870 WP 004918236.1	KIGQGRDTRVKEITKDNPEIAEELAKIREKLQAK	343
NTDB id 1389 A4U84 RS07560 WP 021112712.1	KIGQGKNNAVKWLKENPEIADKLEQEIRNTLASNPNSVL KATASDSND . ESDVEEFSE EEL	372
NTDB id 1130 NGFG RS03960 WP 003688695.1	KIGQKDNVRVWLKENPEISDEIDAKIRALNCEMHITE GTQDETGE . R PE E	348
NTDB id 1124 NGFG RS03960 WP 003688695.1	KIGQKDNVRVWLKENPEISDEIDAKIRALNCEMHITE GTQDETGE . R PE E	348
NTDB id 1123 OK783 RS03845 WP 003688695.1	KIGQKDNVRVWLKENPEISDEIDAKIRALNCEMHITE GTQDETGE . R PE E	348
NTDB id 1427 RS RS02760 WP 011000510.1	RIGQGRDNCREFLRENAREIENKVRREHLGVTPMGAV TLA . EEVEE . D	352
NTDB id 1165 A1552VC RS01565 WP 000344154.1	KIGQKANAACKYLKENPEIAKTLDDKKIREMLLNPNMQL IAETSSAAD . DVEFGAVPE EF	354
NTDB id 1142 GTF74 RS11550 WP 000344154.1	KIGQKANAACKYLKENPEIAKTLDDKKIREMLLNPNMQL IAETSSAAD . DVEFGAVPE EF	354
NTDB id 1022 ACIAD RS06390 WP 004925639.1	KIGQKNNVIRYLEENPCISTEIEAVIREQLLTKASDQT AA . HDETEE . EPDLL . ES	349
NTDB id 1074 ABD1 RS10195 WP 000344167.1	KIGQKNNVIRYFEENNCIAEIEIRNIREQLLTTGTNGA VQ . I . EDEE . EPDLL . LE S	349
NTDB id 1205 PSJM300 05545 AFN77183.1	KIGQKANAACKYLEENPEIQQEIEQQIRAKLLVTSANTK ATPVAEDLA . DADI	347
consensus	***! !***** * * ***** ***** * * * * *	

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	E	I	E	L	E	E	
NTDB id 600 KW2 RS01765 WP 011675413.1	387
NTDB id 269 KZH43 RS08665 WP 001085462.1	E	I	E	I	E	E	388
NTDB id 228 SPD RS09265 WP 001085462.1	E	I	E	I	E	E	388
NTDB id 194 SPR RS08825 WP 001085462.1	E	I	E	I	E	E	388
NTDB id 159 SP RS09750 WP 001085462.1	E	I	E	I	E	E	388
NTDB id 499 SM12261 RS08315 WP 078228268.1	E	I	E	I	E	E	384
NTDB id 527 SMSK321 RS11065 WP 001085454.1	E	I	E	I	E	E	384
NTDB id 453 Spy49 1753c ACI62003.1	E	I	E	D	.	.	378
NTDB id 408 SMU RS09530 WP 002262392.1	E	L	E	D	.	.	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 54755 PLE7327 RS03840 WP 015142548.1	353
NTDB id 1245 Cj1673c YP 002345041.1	343
NTDB id 1211 C694 RS00795 WP 000963128.1	347
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NTDB id 1334 RA0C RS04870 WP 004918236.1	343
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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 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 1022 ACIAD RS06390 WP 004925639.1	349
NTDB id 1074 ABD1 RS10195 WP 000344167.1	349
NTDB id 1205 PSJM300 05545 AFN77183.1	347
consensus							

non conserved
 similar
 ≥ 50% conserved