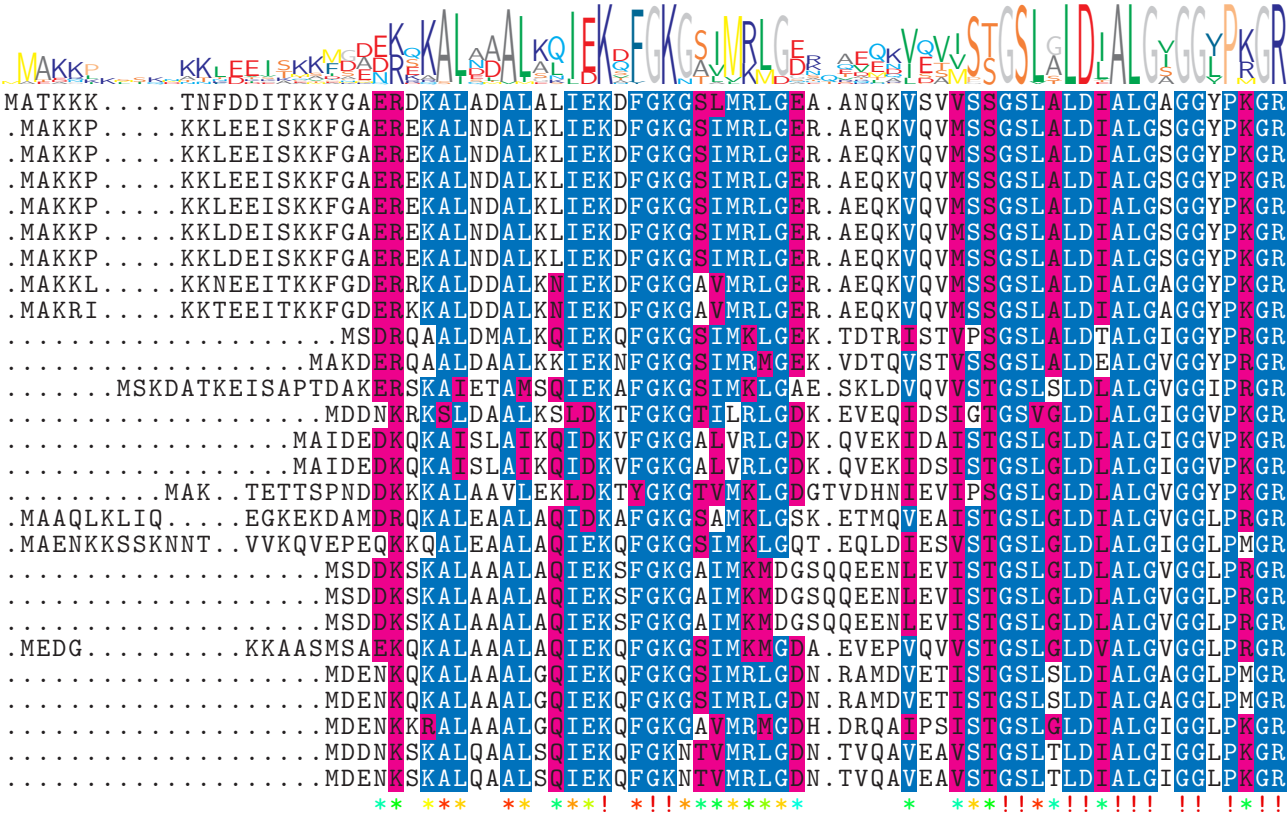


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consensus



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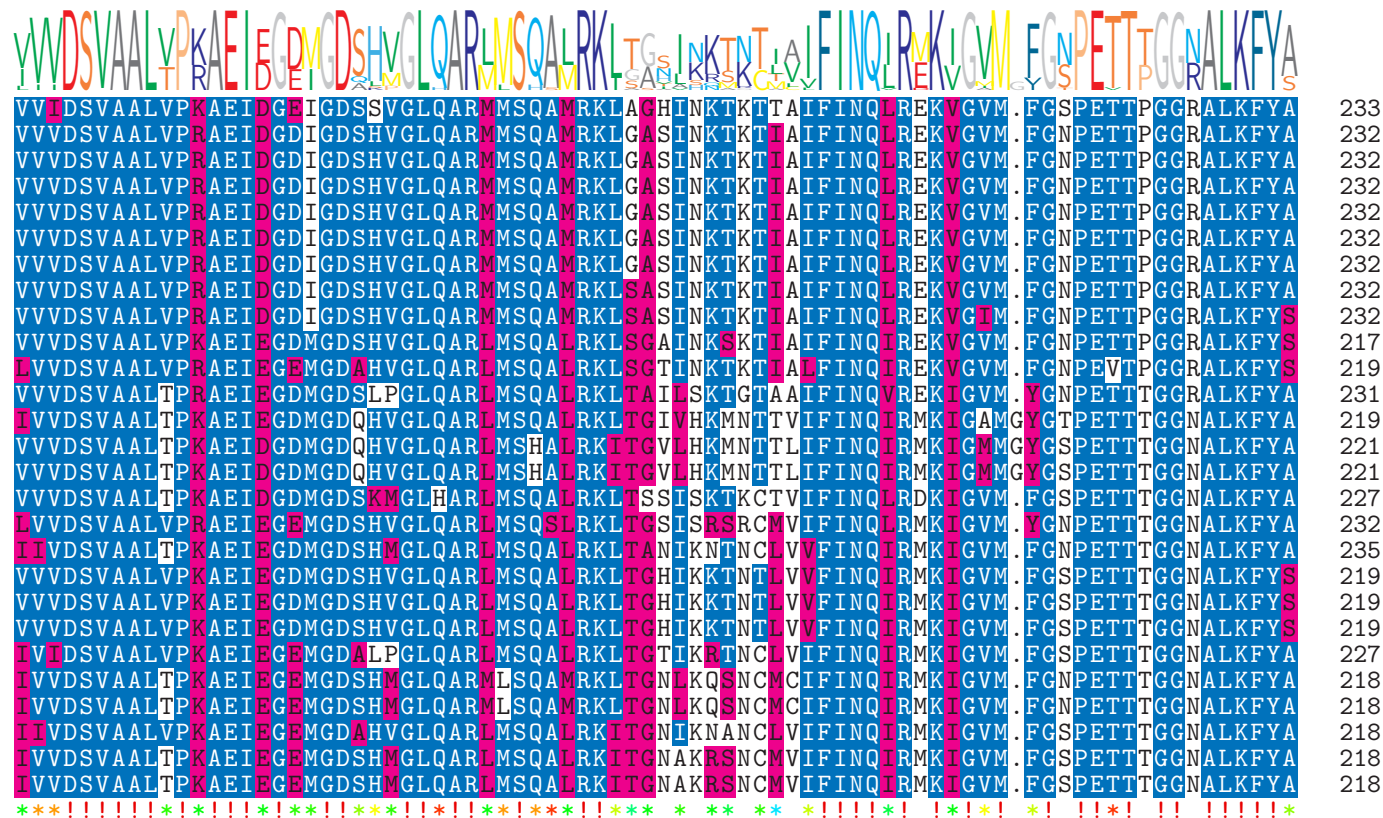
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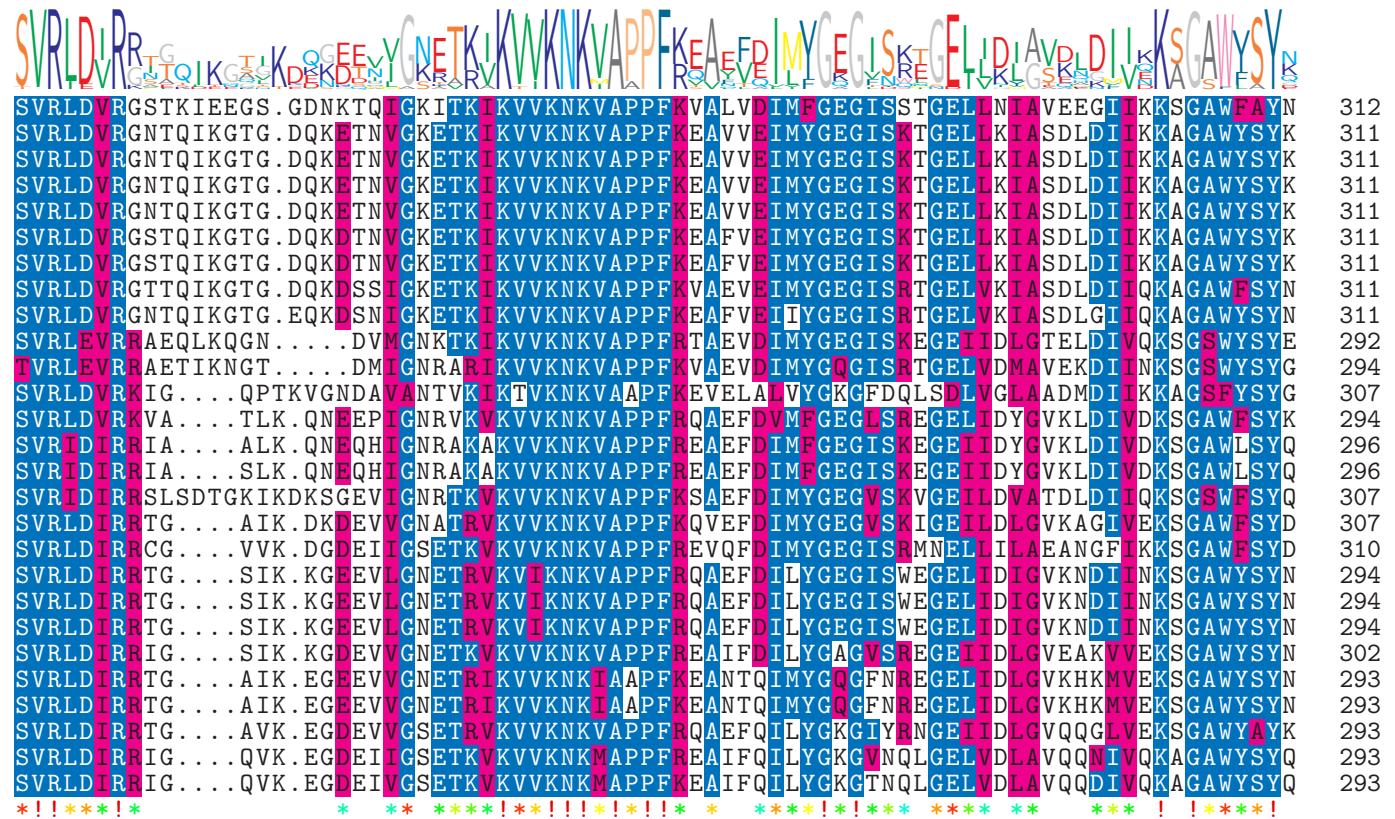
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
consensus

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DEKIGQGS ¹¹ ENAKKYLA ¹² EHPEIF ¹³ DEIDKQVRSKFG ¹⁴ LIDGEEVSEQDTEN.KKDE....PKKEEA..VNEEVPLDLGDELEI	384
DEKIGQGS ¹⁵ ENAKKYLA ¹⁶ EHPEIF ¹⁷ DEIDKQVRSKFG ¹⁸ LIDGEEVSEQDTEN.KKDE....PKKEEA..VNEEVPLDLGDELEI	384
DEKIGQGS ¹⁹ ENAKKYLA ²⁰ ADNPEIF ²¹ DEIDHQVRVQFGLIDGEEIAES....KKDE....AAQVDS..VNEEVPLDLGDELEI	380
DEKIGQGS ²² ENAKKYLA ²³ ADNPEIF ²⁴ DEIDHQVRVQFGLIDGEEIAES....KKDE....VAQVDS..VNEEVPLDLGDELEI	380
GEKIGQGS ²⁵ ENAKRYLADHPQL ²⁶ FDEIDRKVRVKFGLLEESEESAMAVA.SE.....E.....TDDLALDLNGIEI	376
GEKIGQGS ²⁷ ENAKKFLADNPEIF ²⁸ DDIDHKVRVQYGLIEEDNTEEKQSSK.EK.....ETDEK..ADKNLVLELDDTIEL	381
EERL ²⁹ GQGR ³⁰ ENAKQFLKENKDIMLMIQEQIREHYGLDNGGVVQQQAET.QEEL.EFE.E.....	348
SERIGQGR ³¹ ENAKNYLADHEDVE ³² DEVRLKVR ³³ AAAYG ³⁴ ISDVPEEDLPTTED.E.QI.NILPDDST....EE.....	355
DERIGQG ³⁵ KEKTIA ³⁶ YIAERP ³⁷ MEQEIT ³⁸ RDVMAAIRAGNAGEAPALAPAPAPEAAEA.....	363
DKKL ³⁹ GQGR ⁴⁰ ENSKAFLKENPEIADEITKAIQNSMG ⁴¹ IEGMIS..GSEDDEGE..E.....	343
DKKL ⁴² GQGR ⁴³ ENAKALLKEDKALADEITLKIKESIGSNEEIM..PLPDEPLE..EME.....	347
DKKL ⁴⁴ GQGR ⁴⁵ ENAKALLKEDKALADEITLKIKESIGSNEEIM..PLPDEPLE..EME.....	347
DTKL ⁴⁶ GQGRD ⁴⁷ TVKELIKDNPELAE ⁴⁸ ELEAKIREKLQAK.....	343
SIRIGQGR ⁴⁹ ENSKAYLKANPEVADRLEKAI ⁵⁰ RTKTEGLAEEM..MVGPS ⁵¹ EGDDDD.....	357
GTKIGQG ⁵² KNN ⁵³ AVKWLKENPEIADKLEQEIRNTLASNPNSV..LKATASDSNDESDVEEFS.....EEEL.....	372
GAKIGQG ⁵⁴ KDNVRVWLKENPEISDEIDAKIRALNGVEMHIT..EGT.....QDETGERP.....EE.....	348
GAKIGQG ⁵⁵ KDNVRVWLKENPEISDEIDAKIRALNGVEMHIT..EGT.....QDETGERP.....EE.....	348
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GDKIGQG ⁶⁰ KANACKYLKENPEIAKTLDKKLREMLLNPNMQ..LIAETSSAADDVEFGAVP.....EEF.....	354
GDKIGQG ⁶¹ KANACKYLKENPEIAKTLDKKLREMLLNPNMQ..LIAETSSAADDVEFGAVP.....EEF.....	354
GSKIGQG ⁶² KANAACKYLEENPEIGQEIEQQIRAKLLVTSANT..KATPVAEDLADADI.....	347
GNKIGQG ⁶³ KNNVIRYLEENPQISTEIEAVIREQLLTKASDQ..TAA.HDETEEEPDLL..E.....S.....	349
GNKIGQG ⁶⁴ KNNVIRIFYEENQIAEIERNI ⁶⁵ REQLLTGTNG..AVQ.I.EDEE ⁶⁶ EPDLL..L.....ES.....	349

logo

		EIEE	
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 499 SM12261 RS08315 WP 078228268.1	EIEE		384
NTDB id 527 SMSK321 RS11065 WP 001085454.1	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 301026 TQ38 RS11800 WP 043980575.1			357
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 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