

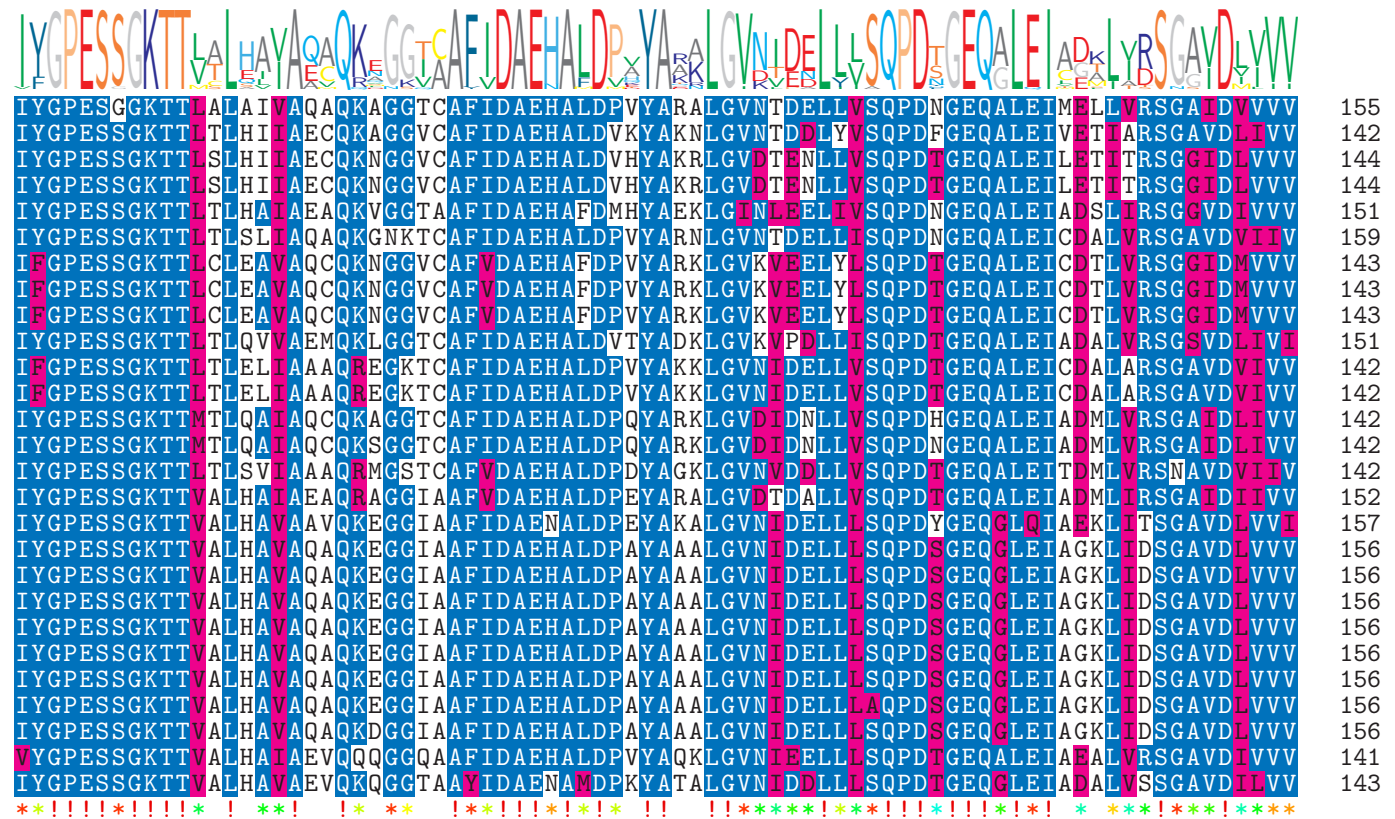
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NTDB id 1142 GTF74 RS11550 WP 000344154.1	MDENK	KKALAAALG	QIEKQFGKGSIMRLGDN.RAMD	VETISTGSLSLD	IALGAGGLPMGRIVE	62	
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NTDB id 625 LCA RS02525 WP 011374196.1	MAKD	ERQAALDAALKK	IEKNFGKGSIMRMGEK.VDTQ	VSTVSSGSLALDE	ALGVGGYP	PRGRIVE	63
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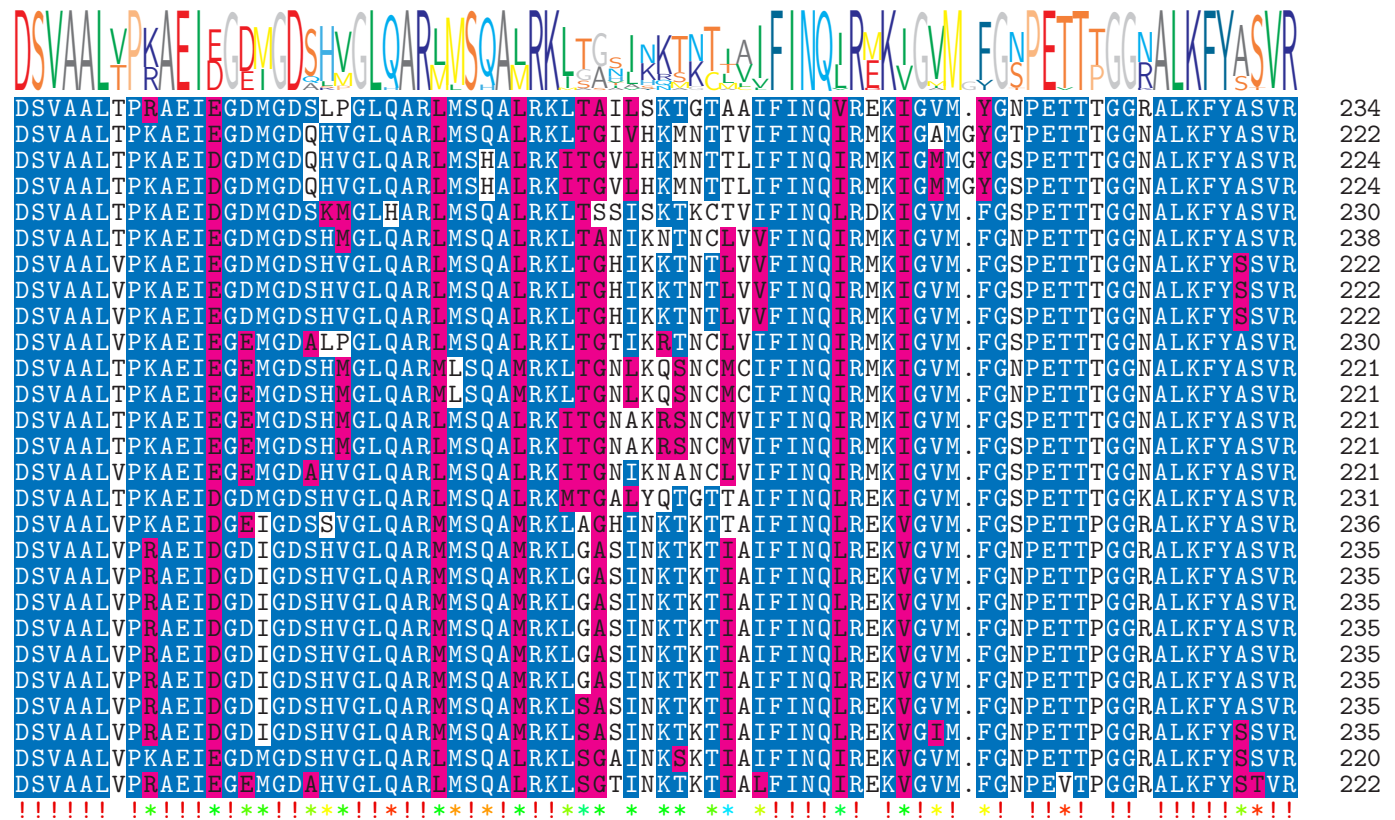
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logo

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consensus	***	!	*										*																																																																										

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NTDB id 1245 Cj1673c YP 002345041.1	LGQGRENSKAF LKENPEIADEITKAIQNSMGI EGMISGSEDD..EGEE.....	343
NTDB id 1214 NCTC11637 00359 SQJ03206.1	LGQGRENAKALLKEDKALADEITLKI KESIGSNEEIIMPLPDE..PLEEM.E.....	347
NTDB id 1211 C694 RS00795 WP 000963128.1	LGQGRENAKALLKEDKALADEITLKI KESIGSNEEIIMPLPDE..PLEEM.E.....	347
NTDB id 1334 RAOC RS04870 WP 004918236.1	LGQGRD TVKELIKDNPELAE EEL EAKIREKLQAK.....	343
NTDB id 1389 A4U84 RS07560 WP 021112712.1	IGQGGKNNAVKWLKENPEIADKLEQEIRNTLASNPNSVLKATASDSNDES.DVE.....EFSEEEL.....	372
NTDB id 1130 NGFG RS03960 WP 003688695.1	IGQGGKDNVRVWLKENPEISDEIDAKIRALNGVEMHITEGTQDET DGER.....PEE.....	348
NTDB id 1124 NGFG RS03960 WP 003688695.1	IGQGGKDNVRVWLKENPEISDEIDAKIRALNGVEMHITEGTQDET DGER.....PEE.....	348
NTDB id 1123 OK783 RS03845 WP 003688695.1	IGQGGKDNVRVWLKENPEISDEIDAKIRALNGVEMHITEGTQDET DGER.....PEE.....	348
NTDB id 1427 RS RS02760 WP 011000510.1	IGQGRDNCR EFLRENAELAREIENK VREHLGVTPMGAVTLA.EEVEED.....	352
NTDB id 1165 A1552VC RS01565 WP 000344154.1	IGQGGKANACKYLLKENPEIAKTLDKKLREMLLNPNENMQLIAETSSAAD DV.EFG.....AVPEEF.....	354
NTDB id 1142 GTF74 RS11550 WP 000344154.1	IGQGGKANACKYLLKENPEIAKTLDKKLREMLLNPNENMQLIAETSSAAD DV.EFG.....AVPEEF.....	354
NTDB id 1022 ACIAD RS06390 WP 004925639.1	IGQGGKNNVIRYLEENPQISTEIEAVIREQLLTKASDQTA A.HDETEEEP.DLL.....ES.....	349
NTDB id 1074 ABD1 RS10195 WP 000344167.1	IGQGGKNNVIRYFEENNQIAEEIERNIREQLLTTGTNGAVQ.I.EDEEEP.DLL.....LES.....	349
NTDB id 1205 PSJM300 05545 AFN77183.1	IGQGGKANA AKYLEENPEIGQEIEQQIRAKLLVTSANTKATPVAEDLADA.DI.....	347
NTDB id 24390 JK RS05785 WP 011273662.1	LGQGGKEKAREFLKNNRDL SAELEDR IARELKIGPYAKMKDEQTEEA A.....GDQMDEDKPIDLSPNFDDDDAN	375
NTDB id 600 KW2 RS01765 WP 011675413.1	IGQGA EKAKNYLKEHQDVFDEIDHKVRAAHGLLDDSEVAETEEETTAS KTKAKAKKEEKAVETEEIELELED.....	387
NTDB id 269 KZH43 RS08665 WP 001085462.1	IGQGS ENAKKYLA EHPEIFDEIDKQVRSKFGLIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIEIEE	388
NTDB id 228 SPD RS09265 WP 001085462.1	IGQGS ENAKKYLA EHPEIFDEIDKQVRSKFGLIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIEIEE	388
NTDB id 194 SPR RS08825 WP 001085462.1	IGQGS ENAKKYLA EHPEIFDEIDKQVRSKFGLIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIEIEE	388
NTDB id 159 SP RS09750 WP 001085462.1	IGQGS ENAKKYLA EHPEIFDEIDKQVRSKFGLIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIEIEE	388
NTDB id 499 SM12261 RS08315 WP 078228268.1	IGQGS ENAKKYLA DNPEIFDEIDHQVRVQFGLIDGEEIAES....KKDE....AAQVDS..VNEEVTLDLGDELEIEIEE	384
NTDB id 527 SMSK321 RS11065 WP 001085454.1	IGQGS ENAKKYLA DNPEVFAEIDHQVRVQFGLIDGEEAAES....KKDE....VAQVDS..VNEEVTLDLGDELEIEIEE	384
NTDB id 453 Spy49 1753c ACI62003.1	IGQGS ENAKRYLA DHPQLFDEIDRKVVRVKFGLLEESEESAMAVASE.....E.....TDDLALDLNNGIEIED..	378
NTDB id 408 SMU RS09530 WP 002262392.1	IGQGS ENAKKFLADNPEIFDDIDHKVVRVQYGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELED..	383
NTDB id 115 BSU 16940 NP 389576.2	LGQGRENAKQFLKENKDIMLMIQEQIREHYGLDNNGVVQQAEETQEEL.EFE.E.....	348
NTDB id 625 LCA RS02525 WP 011374196.1	IGQGRENAKNYLA DHEDVEDEVRLKVR AAYGISDVPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
consensus	*!!!!***** * * ***** ***** * * * *	

- X

 non conserved
- X

 similar
- X

 ≥ 50% conserved