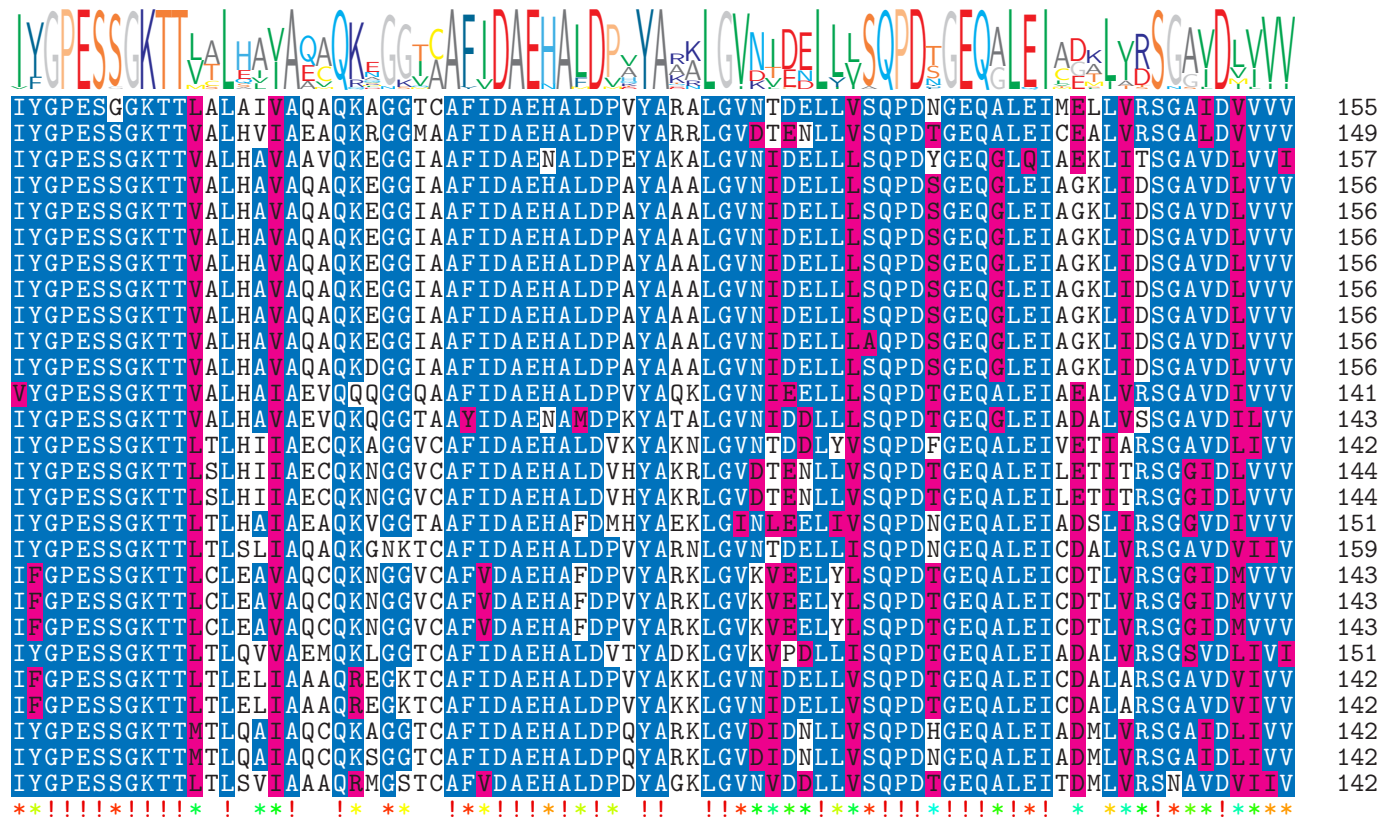


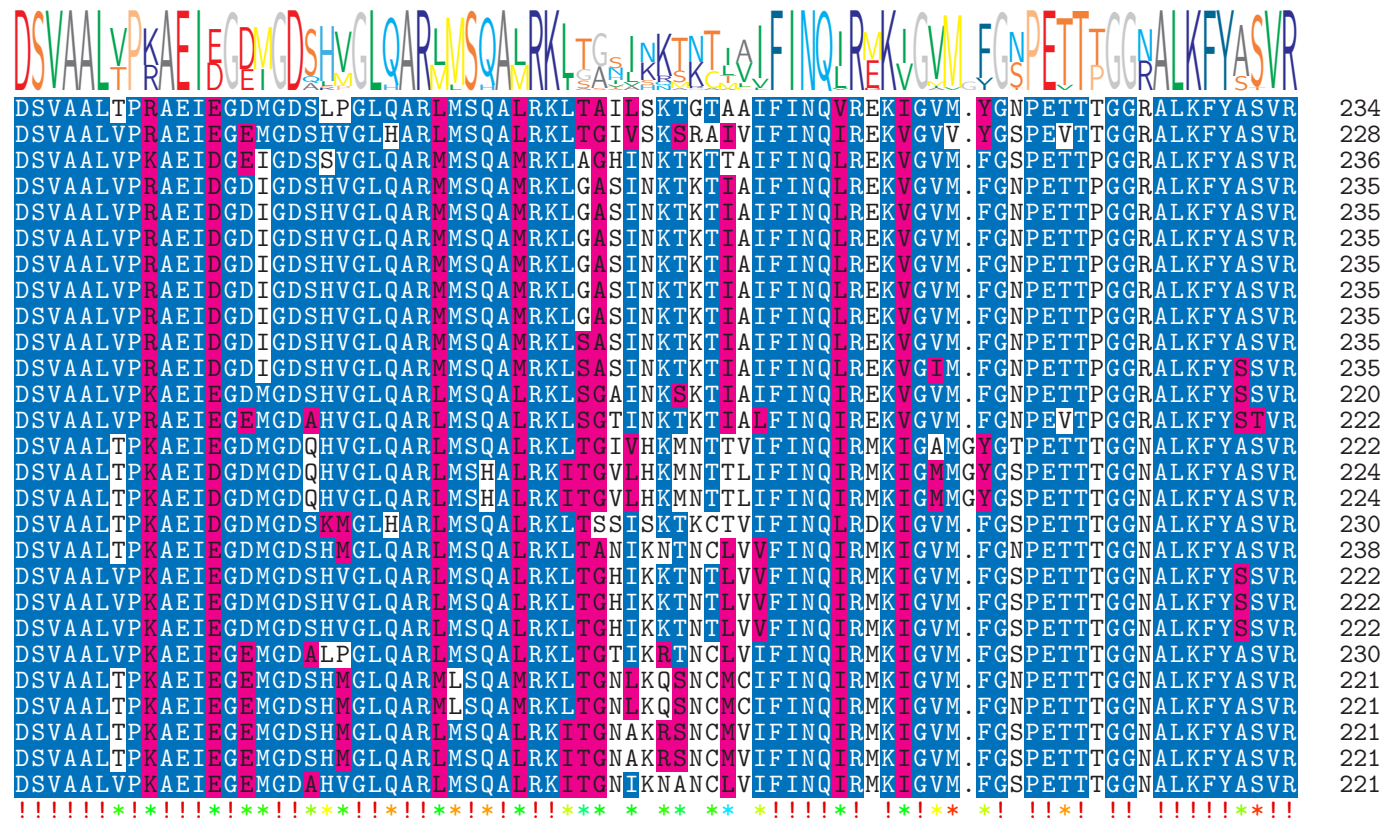
logo

NTDB id 1313 DR RS12030 WP 010888966.1	MSK...DATKEISAPTDAKERSKATETAM	75
NTDB id 217895 B0537 RS09475 WP 077714368.1	MADKTDRLSADKQKALELALANIERNFGKGSIMKLGAS.TKLNIEAISTGALS	69
NTDB id 600 KW2 RS01765 WP 011675413.1	MAT..KKKTNFDDITKKYGAERDKALADALALIEKDFGKGSIMRLGEA.ANQKVS	77
NTDB id 269 KZH43 RS08665 WP 001085462.1	.MA..KKPKKLEEISKKFGAEREKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSS	76
NTDB id 228 SPD RS09265 WP 001085462.1	.MA..KKPKKLEEISKKFGAEREKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSS	76
NTDB id 194 SPR RS08825 WP 001085462.1	.MA..KKPKKLEEISKKFGAEREKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSS	76
NTDB id 159 SP RS09750 WP 001085462.1	.MA..KKPKKLEEISKKFGAEREKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSS	76
NTDB id 499 SM12261 RS08315 WP 078228268.1	.MA..KKPKKLDEISKKFGAEREKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSS	76
NTDB id 527 SMSK321 RS11065 WP 001085454.1	.MA..KKPKKLDEISKKFGAEREKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSS	76
NTDB id 453 Spy49 1753c ACI62003.1	.MA..KKLKKNEEITKKFGDERRKALDDALKNIEKDFGKGAVMRLGER.AEQKVQVMSS	76
NTDB id 408 SMU RS09530 WP 002262392.1	.MA..KRIKKTEEITKKFGDERKKALDDALKNIEKDFGKGAVMRLGER.AEQKVQVMSS	76
NTDB id 115 BSU 16940 NP 389576.2	MSDRQAALDMALKQIEKQFGKGSIMKLGEK.TDTRISTVPSGSLALDTALG	61
NTDB id 625 LCA RS02525 WP 011374196.1	MAKDERRQAALDAALKKIEKNFGKGSIMRMGEK.VDTQVSTVSSGSLALDEAL	63
NTDB id 1245 Cj1673c YP 002345041.1	MDDNKRKSLDAALKSLDKTFGKGITILRLGDK.EVEQIDSIGTGSVGLDLALG	62
NTDB id 1214 NCTC11637 00359 SQJ03206.1	MAIDEDKQKATISLAIKQIDKVFSGKALVRLGDK.QVEKIDAISTGSLGLDLAL	64
NTDB id 1211 C694 RS00795 WP 000963128.1	MAIDEDKQKATISLAIKQIDKVFSGKALVRLGDK.QVEKIDSISTGSLGLDLAL	64
NTDB id 1334 RAOC RS04870 WP 004918236.1	MAKTETTSPNDKKKKALAAVLEKLDKTYGKGTVMKLGDTVDHNIIEVIPSGSL	71
NTDB id 1389 A4U84 RS07560 WP 021112712.1	MAENKKSSKNNTTVVKQVEPEQKKQALEAALAQIEKQFGKGSIMKLGQT.EQLDIESV	79
NTDB id 1130 NGFG RS03960 WP 003688695.1	MSDDKSKALAAALAQIEKSFGKGAIMKMDGSQQEENLEVISTGSLGLDLALG	63
NTDB id 1124 NGFG RS03960 WP 003688695.1	MSDDKSKALAAALAQIEKSFGKGAIMKMDGSQQEENLEVISTGSLGLDLALG	63
NTDB id 1123 OK783 RS03845 WP 003688695.1	MSDDKSKALAAALAQIEKSFGKGAIMKMDGSQQEENLEVISTGSLGLDLALG	63
NTDB id 1427 RS RS02760 WP 011000510.1	MEDG.....KKAASMSAEKQKALAAALAQIEKQFGKGSIMKMGDA.EVEPVQVVSTG	71
NTDB id 1165 A1552VC RS01565 WP 000344154.1	MDENKQKALAAALGQIEKQFGKGSIMRLGDN.RAMDVETISTGSLSLDIALG	62
NTDB id 1142 GTF74 RS11550 WP 000344154.1	MDENKQKALAAALGQIEKQFGKGSIMRLGDN.RAMDVETISTGSLSLDIALG	62
NTDB id 1022 ACIAD RS06390 WP 004925639.1	MDDNKSALQAALSQIEKQFGKNTVMRLGDN.TVQAVEAVSTGSLTLDIALG	62
NTDB id 1074 ABD1 RS10195 WP 000344167.1	MDENKSALQAALSQIEKQFGKNTVMRLGDN.TVQAVEAVSTGSLTLDIALG	62
NTDB id 1205 PSJM300 05545 AFN77183.1	MDENKKRALAAALGQIEKQFGKGAVMRMGDH.DRQAIPSIISTGSLGLDIALG	62
consensus	* * * * *	

NTDB id 1313 DR RS12030 WP 010888966.1
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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
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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
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
NTDB id 1205 PSJM300 05545 AFN77183.1
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NTDB id 1313 DR RS12030 WP 010888966.1
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NTDB id 1124 NGFG RS03960 WP 003688695.1
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NTDB id 1205 PSJM300 05545 AFN77183.1
consensus



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NTDB id 1313 DR RS12030 WP 010888966.1	LDVRRIGQ...PTKVGND	310
NTDB id 217895 B0537 RS09475 WP 077714368.1	MEVRKSENKIQGT...EI	303
NTDB id 600 KW2 RS01765 WP 011675413.1	LDVRGSKIEEGS.GDNKT	315
NTDB id 269 KZH43 RS08665 WP 001085462.1	LDVRGNTQIKGTG.DQK	314
NTDB id 228 SPD RS09265 WP 001085462.1	LDVRGNTQIKGTG.DQK	314
NTDB id 194 SPR RS08825 WP 001085462.1	LDVRGNTQIKGTG.DQK	314
NTDB id 159 SP RS09750 WP 001085462.1	LDVRGNTQIKGTG.DQK	314
NTDB id 499 SM12261 RS08315 WP 078228268.1	LDVRGSTQIKGTG.DQK	314
NTDB id 527 SMSK321 RS11065 WP 001085454.1	LDVRGSTQIKGTG.DQK	314
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NTDB id 408 SMU RS09530 WP 002262392.1	LDVRGNTQIKGTG.EQK	314
NTDB id 115 BSU 16940 NP 389576.2	LEVRRAEQLKQGN....	295
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NTDB id 1245 Cj1673c YP 002345041.1	LDVRKVAT...L.KQN	297
NTDB id 1214 NCTC11637 00359 SQJ03206.1	IDIRRIAA...L.KQN	299
NTDB id 1211 C694 RS00795 WP 000963128.1	IDIRRIAS...L.KQN	299
NTDB id 1334 RAOC RS04870 WP 004918236.1	IDIRRLSDTGKIKDKS	310
NTDB id 1389 A4U84 RS07560 WP 021112712.1	LDIRRCGV...V.KDG	313
NTDB id 1130 NGFG RS03960 WP 003688695.1	LDIRRTGS...I.KKG	297
NTDB id 1124 NGFG RS03960 WP 003688695.1	LDIRRTGS...I.KKG	297
NTDB id 1123 OK783 RS03845 WP 003688695.1	LDIRRTGS...I.KKG	297
NTDB id 1427 RS RS02760 WP 011000510.1	LDIRRIGS...I.KKG	305
NTDB id 1165 A1552VC RS01565 WP 000344154.1	LDIRRTGA...I.KEG	296
NTDB id 1142 GTF74 RS11550 WP 000344154.1	LDIRRTGA...I.KEG	296
NTDB id 1022 ACIAD RS06390 WP 004925639.1	LDIRRIIGQ...V.KEG	296
NTDB id 1074 ABD1 RS10195 WP 000344167.1	LDIRRIIGQ...V.KEG	296
NTDB id 1205 PSJM300 05545 AFN77183.1	LDIRRTGA...V.KEG	296
consensus	***!* * ** *****!!!!!!* * *****!*!*** ** *** ! !*****! *	

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NTDB id 1313 DR RS12030 WP 010888966.1	IGQGKEKTIA Y IA E RP P EME Q EIRDRVMAAIRAGNAGEAPALAPAPA.....APEAAEA.....	363
NTDB id 217895 B0537 RS09475 WP 077714368.1	LGGGREN A KETL K TRP E LA Q EIERLVREKLNQGNVQ...LPSAEGALDQD.....LMDEDMI.....	357
NTDB id 600 KW2 RS01765 WP 011675413.1	IGQGA E KAKN Y LKEH Q DV F DEIDHKVRAAHGILLDDSEVAETEEETTAS K TKAKAKKEEKAVETEEIELELED.....	387
NTDB id 269 KZH43 RS08665 WP 001085462.1	IGQGS E NAKKYLA E HP E IF D EIDKQVRSKFGLIDGEEVSEQDTENKK D E....PKKEEA..VNEEVPLDLGDELEIEIEE	388
NTDB id 228 SPD RS09265 WP 001085462.1	IGQGS E NAKKYLA E HP E IF D EIDKQVRSKFGLIDGEEVSEQDTENKK D E....PKKEEA..VNEEVPLDLGDELEIEIEE	388
NTDB id 194 SPR RS08825 WP 001085462.1	IGQGS E NAKKYLA E HP E IF D EIDKQVRSKFGLIDGEEVSEQDTENKK D E....PKKEEA..VNEEVPLDLGDELEIEIEE	388
NTDB id 159 SP RS09750 WP 001085462.1	IGQGS E NAKKYLA E HP E IF D EIDKQVRSKFGLIDGEEVSEQDTENKK D E....PKKEEA..VNEEVPLDLGDELEIEIEE	388
NTDB id 499 SM12261 RS08315 WP 078228268.1	IGQGS E NAKKYLADN P EIF D EIDH Q VRVQFGLIDGEEIAES...KKDE...AAQVDS..VNEEVTLDLGDELEIEIEE	384
NTDB id 527 SMSK321 RS11065 WP 001085454.1	IGQGS E NAKKYLADN P EV F AEIDH Q VRVQFGLIDGEEAAES...KKDE...VAQVDS..VNEEVTLDLGDELEIEIEE	384
NTDB id 453 Spy49 1753c ACI62003.1	IGQGS E NAKRYLADHP Q LF D EIDR K VRVKFGLLEESEESAMAVASE.....E.....TDDLALDLNNGIEIED..	378
NTDB id 408 SMU RS09530 WP 002262392.1	IGQGS E NAKKFLADN P EIF D DDIDH K VRVQYGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELED..	383
NTDB id 115 BSU 16940 NP 389576.2	LGGGREN A KQFL K EN K DIMLMIQEQIREHYGLDNNGVVQQQAEETQEEL.EFE.E.....	348
NTDB id 625 LCA RS02525 WP 011374196.1	IGQGRENAK N YLADH E D V E D EVRLKVR A AYGISDVPEEDLPTTEDE.QI.NILPDDST.....EE.....	355
NTDB id 1245 Cj1673c YP 002345041.1	LGGGRENSK A FL K EN P EIA D EITKAIQNSMGIEGMISGSEDD..EGEE.....	343
NTDB id 1214 NCTC11637 00359 SQJ03206.1	LGGGRENAKALL K EDKALADEITLKIKESIGSNEEIMPLPDE..PLEEME.....	347
NTDB id 1211 C694 RS00795 WP 000963128.1	LGGGRENAKALL K EDKALADEITLKIKESIGSNEEIMPLPDE..PLEEME.....	347
NTDB id 1334 RAOC RS04870 WP 004918236.1	LGGGRD T V K EL I KDN P E L A E E L EAKIREKLQAK.....	343
NTDB id 1389 A4U84 RS07560 WP 021112712.1	IGQGK N NAV K WLKEN P EIA D KLEQEIRNTLASNPNSVLKATASDSN E SD.VEE.....FSEEEL.....	372
NTDB id 1130 NGFG RS03960 WP 003688695.1	IGQGKDNVRVWLKEN P EIS D EIDAKIRALNGVEMHITEGTQDET D GER.....PEE.....	348
NTDB id 1124 NGFG RS03960 WP 003688695.1	IGQGKDNVRVWLKEN P EIS D EIDAKIRALNGVEMHITEGTQDET D GER.....PEE.....	348
NTDB id 1123 OK783 RS03845 WP 003688695.1	IGQGKDNVRVWLKEN P EIS D EIDAKIRALNGVEMHITEGTQDET D GER.....PEE.....	348
NTDB id 1427 RS RS02760 WP 011000510.1	IGQGRDN C REFLRENAELAREIENK V REHLGVTPMGAVTLA.EEVEED.....	352
NTDB id 1165 A1552VC RS01565 WP 000344154.1	IGQGKANACKY L KEN P EIAKTLDKKLREMLLNPNENMQLIAETSSAADDVE.FGA.....VPEEF.....	354
NTDB id 1142 GTF74 RS11550 WP 000344154.1	IGQGKANACKY L KEN P EIAKTLDKKLREMLLNPNENMQLIAETSSAADDVE.FGA.....VPEEF.....	354
NTDB id 1022 ACIAD RS06390 WP 004925639.1	IGQGKNNVIRY L EEN P QISTEIEAVIREQLLTKASDQTAA.HDETEEEP D .LL.....ES.....	349
NTDB id 1074 ABD1 RS10195 WP 000344167.1	IGQGKNNVIRY L EEN N QIA E EIERNIREQLLTTGTNGAVQ.I.EDEEEP D .LL.....LES.....	349
NTDB id 1205 PSJM300 05545 AFN77183.1	IGQGKANAAKY L EEN P EIGQEIEQQIRAKLLVTSANTKATPVAEDLADAD.I.....	347
consensus	*!!!!***** ** * **** ***** ** *	

- X

 non conserved
- X

 similar
- X

 ≥ 50% conserved