

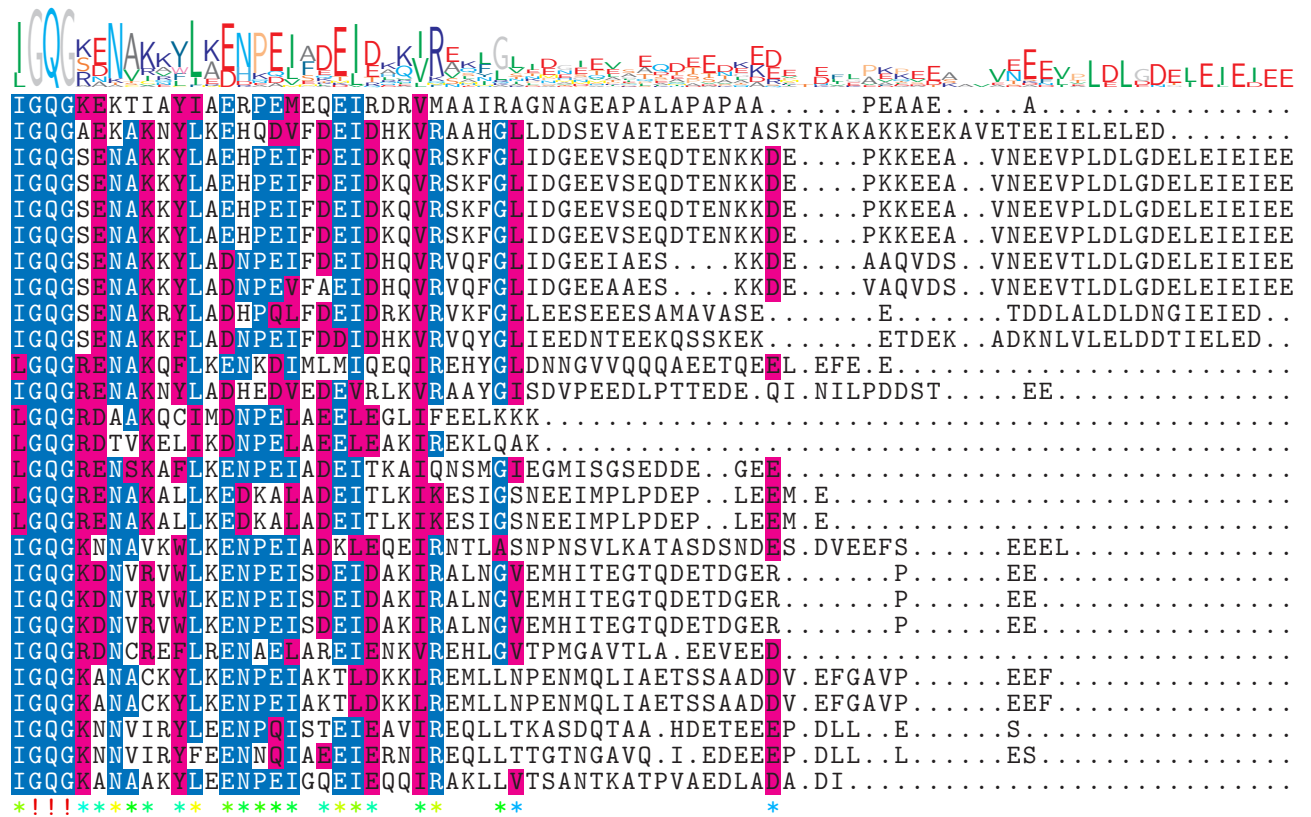
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NTDB id 1313 DR RS12030 WP 010888966.1	MSKD...ATKEISAPTDAKERSKATETAMSQIEKAFGKGSIMKLGAE.SKLDVQVVSTGSLSLDLALGVGGIPRGRITE	75
NTDB id 600 KW2 RS01765 WP 011675413.1	MA..TKKKTNFDDITKKYGAERDKALADALALIEKDFGKGSIMRLGAEA.ANQKVSVVSSGSLALDIALGAGGYPKGRIVE	77
NTDB id 269 KZH43 RS08665 WP 001085462.1	.M..AKKPKKLEEISKKFGEERKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGYPKGRIVE	76
NTDB id 228 SPD RS09265 WP 001085462.1	.M..AKKPKKLEEISKKFGEERKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGYPKGRIVE	76
NTDB id 194 SPR RS08825 WP 001085462.1	.M..AKKPKKLEEISKKFGEERKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGYPKGRIVE	76
NTDB id 159 SP RS09750 WP 001085462.1	.M..AKKPKKLEEISKKFGEERKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGYPKGRIVE	76
NTDB id 499 SM12261 RS08315 WP 078228268.1	.M..AKKPKKLDEISKKFGEERKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGYPKGRIVE	76
NTDB id 527 SMSK321 RS11065 WP 001085454.1	.M..AKKPKKLDEISKKFGEERKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGYPKGRIVE	76
NTDB id 453 Spy49 1753c ACI62003.1	.M..AKKLKKNEEITKKFGDERRKALDDALKNIEKDFGKGAVMRLGER.AEQKVQVMSSGSLALDIALGAGGYPKGRIVE	76
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NTDB id 115 BSU 16940 NP 389576.2MSDRQAALDMALKQIEKQFGKGSIMKLGEEK.TDTRISTVPSGSLALDIALGIGGYPGRGRIVE	61
NTDB id 625 LCA RS02525 WP 011374196.1MAKDERQAALDAALKKIEKNFGKGSIMRMGEK.VDTQVSTVSSGSLALDEALGVGGYPGRGRIVE	63
NTDB id 376588 FQN58 RS12210 WP 004312516.1	MA..KKDELNFETDNKMASSKLLKALQAAMEKIEKSFGKGSIMKMGDDSVQVEVPTGSLALNVALGVGGYPGRGRIVE	77
NTDB id 1334 RAO RS04870 WP 004918236.1MAKTETTSPNDKPKKALAAVLEKLDKTYGKGTVMKLGDTVDHNIIEVIPSGSLGLDLALGVGGYPKGRIVE	71
NTDB id 1245 Cj1673c YP 002345041.1MDDNKRKSLDAALKSLDKTFGKGTILRLGDK.EVEQIDISIGTGSVGLDLALGIGGVPKGRIVE	62
NTDB id 1214 NCTC11637 00359 SQJ03206.1MAIDEDKQKATISLAIKQIDKVFSGKALVRLGDK.QVEKIDAISTGSLGLDLALGIGGVPKGRIVE	64
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NTDB id 1389 A4U84 RS07560 WP 021112712.1	MAENKSSKNNTVVKQVEPEQKKQALEAALAQIEKQFGKGSIMKLGQT.EQLDIESVSTGSLGLDLALGIGGLPMGRIVE	79
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NTDB id 1124 NGFG RS03960 WP 003688695.1MSDDKSKALAAALAQIEKSFSGKAIMKMDGSQEQEENLEVISTGSLGLDLALGVGGLPGRIVE	63
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NTDB id 1427 RS RS02760 WP 011000510.1	MEDG.....KKAASMSAEKQKALAAALAQIEKQFGKGSIMKMGDA.EVEPVQVVSTGSLGLDVALGVGGLPGRIVE	71
NTDB id 1165 A1552VC RS01565 WP 000344154.1MDENKQKALAAALGQIEKQFGKGSIMRLGDN.RAMDVETISTGSLSLDIALGAGGLPMGRIVE	62
NTDB id 1142 GTF74 RS11550 WP 000344154.1MDENKQKALAAALGQIEKQFGKGSIMRLGDN.RAMDVETISTGSLSLDIALGAGGLPMGRIVE	62
NTDB id 1022 ACIAD RS06390 WP 004925639.1MDDNKSALQAALSQIEKQFGKNTVMRLGDN.TVQAVEAVSTGSLTLDIALGIGGLPKGRIVE	62
NTDB id 1074 ABD1 RS10195 WP 000344167.1MDENKSALQAALSQIEKQFGKNTVMRLGDN.TVQAVEAVSTGSLTLDIALGIGGLPKGRIVE	62
NTDB id 1205 PSJM300 05545 AFN77183.1MDENKRALAAALGQIEKQFGKGAVMRMGDH.DRQAIPSIISTGSLGLDIALGIGGLPKGRIVE	62
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NTDB id 1313 DR RS12030 WP 010888966.1
 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
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 consensus



non conserved
 similar
 ≥ 50% conserved