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Label	Sequence	Pos
NTDB id 600 KW2 RS01765 WP 011675413.1	MATKK.....KTNFDDITKKYGAERDKALADALALIEKDFGKGSLMRLGEA.ANQKVSVVSSGSLALDIALGAGGYPKGR	74
NTDB id 269 KZH43 RS08665 WP 001085462.1	.MAKK.....PKKLEEISKKFGAEREKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGYPKGR	73
NTDB id 228 SPD RS09265 WP 001085462.1	.MAKK.....PKKLEEISKKFGAEREKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGYPKGR	73
NTDB id 194 SPR RS08825 WP 001085462.1	.MAKK.....PKKLEEISKKFGAEREKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGYPKGR	73
NTDB id 159 SP RS09750 WP 001085462.1	.MAKK.....PKKLEEISKKFGAEREKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGYPKGR	73
NTDB id 527 SMSK321 RS11065 WP 001085454.1	.MAKK.....PKKLDEISKKFGAEREKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGYPKGR	73
NTDB id 499 SM12261 RS08315 WP 078228268.1	.MAKK.....PKKLDEISKKFGAEREKALNDALKLIEKDFGKGSIMRLGER.AEQKVQVMSSGSLALDIALGSGGYPKGR	73
NTDB id 453 Spy49 1753c ACI62003.1	.MAKK.....LKKNEEITKKFGDERRKALDDALKNIEKDFGKGAVMRLGER.AEQKVQVMSSGSLALDIALGAGGYPKGR	73
NTDB id 408 SMU RS09530 WP 002262392.1	.MAKR.....IKKTEEITKKFGDERRKALDDALKNIEKDFGKGAVMRLGER.AEQKVQVMSSGSLALDIALGSGGYPKGR	73
NTDB id 115 BSU 16940 NP 389576.2	...................MSDRQAALDMALKQIEKDFGKGSIMKLGEK.TDTRISTVPSGSLALDIALGIGGYPRGR	58
NTDB id 625 LCA RS02525 WP 011374196.1	...................MAKDERQAALDAALKKIEKNFGKGSIMRMGEK.VDTQVSTVSSGSLALDEALGVGGYPRGR	60
NTDB id 1313 DR RS12030 WP 010888966.1	.......MSKDATKEISAPTDAKERSKAIETAMSQIEKAFGKGSIMKLGAE.SKLDVQVVSTGSLSLDIALGVGGIPRGR	72
NTDB id 1245 Cj1673c YP 002345041.1	..................MDDNKRKSLDAALKSLDKTFGKGTILRLGDK.EVEQIDSIGTGSVGLDIALGIGGVPKGR	59
NTDB id 1214 NCTC11637 00359 SQJ03206.1	..................MAIDEDKQKAISLAIKQIDKVFGKGALVRLGDK.QVEKIDAISTGSVGLDIALGIGGVPKGR	61
NTDB id 1211 C694 RS00795 WP 000963128.1	..................MAIDEDKQKAISLAIKQIDKVFGKGALVRLGDK.QVEKIDSISTGSLGLDIALGIGGVPKGR	61
NTDB id 1334 RAOC RS04870 WP 004918236.1	..........MAKT.ETTSPNDDKKKALAAVLEKLDKTYGKGTVMKLGDGTVDHNIEVIPSGSLGLDIALGVGGYPKGR	68
NTDB id 1389 A4U84 RS07560 WP 021112712.1	.MAENKKSSKNNTV..VKQVEPEQKKQALEAALAQIEKQFGKGSIMKLGQT.EQLDIESVSTGSLGLDIALGIGGLPMGR	76
NTDB id 1130 NGFG RS03960 WP 003688695.1	...................MSDDKSKALAAALAQIEKSFGKGAIMKMDGSQQEENLEVISTGSLGLDIALGVGGLPRGR	60
NTDB id 1124 NGFG RS03960 WP 003688695.1	...................MSDDKSKALAAALAQIEKSFGKGAIMKMDGSQQEENLEVISTGSLGLDIALGVGGLPRGR	60
NTDB id 1123 OK783 RS03845 WP 003688695.1	...................MSDDKSKALAAALAQIEKSFGKGAIMKMDGSQQEENLEVISTGSLGLDIALGVGGLPRGR	60
NTDB id 1427 RS RS02760 WP 011000510.1	.MEDG........K..KAASMSAEKQKALAAALAQIEKQFGKGSIMKMGDA.EVEPVQVVSTGSLGLDVALGVGGLPRGR	68
NTDB id 1165 A1552VC RS01565 WP 000344154.1	...................MDENKQKALAAALGQIEKQFGKGSIMRLGDN.RAMDVETISTGSLSLDIALGAGGLPMGR	59
NTDB id 1142 GTF74 RS11550 WP 000344154.1	...................MDENKQKALAAALGQIEKQFGKGSIMRLGDN.RAMDVETISTGSLSLDIALGAGGLPMGR	59
NTDB id 1205 PSJM300 05545 AFN77183.1	...................MDENKKRALAAALGQIEKQFGKGAVMRMGDH.DRQAIPSISTGSLGLDIALGIGGLPKGR	59
NTDB id 313049 AhaeAN43 RS07035 WP 005090091.1	...................MDDNKSKALQAALSQIEKQFGKNTVMRLGDN.TVQAVEAVSTGSLTLDIALGIGGLPKGR	59
NTDB id 1074 ABD1 RS10195 WP 000344167.1	...................MDENKSKALQAALSQIEKQFGKNTVMRLGDN.TVQAVEAVSTGSLTLDIALGIGGLPKGR	59
NTDB id 1022 ACIAD RS06390 WP 004925639.1	...................MDDNKSKALQAALSQIEKQFGKNTVMRLGDN.TVQAVEAVSTGSLTLDIALGIGGLPKGR	59

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Label	Sequence	Number
NTDB id 600 KW2 RS01765 WP 011675413.1	IVEIYGPESSGKTTVALHAVAAVQKEGGIAAFIDAENALDPEYAKALGVNIDELLLSQPDYGEQGLQIAEKLITSGAVDL	154
NTDB id 269 KZH43 RS08665 WP 001085462.1	IIEIYGPESSGKTTVALHAVAQAQKEGGIAAFIDAEHALDPAYAAALGVNIDELLLSQPDSGEQGLEIAGKLIDSGAVDL	153
NTDB id 228 SPD RS09265 WP 001085462.1	IIEIYGPESSGKTTVALHAVAQAQKEGGIAAFIDAEHALDPAYAAALGVNIDELLLSQPDSGEQGLEIAGKLIDSGAVDL	153
NTDB id 194 SPR RS08825 WP 001085462.1	IIEIYGPESSGKTTVALHAVAQAQKEGGIAAFIDAEHALDPAYAAALGVNIDELLLSQPDSGEQGLEIAGKLIDSGAVDL	153
NTDB id 159 SP RS09750 WP 001085462.1	IIEIYGPESSGKTTVALHAVAQAQKEGGIAAFIDAEHALDPAYAAALGVNIDELLLSQPDSGEQGLEIAGKLIDSGAVDL	153
NTDB id 527 SMSK321 RS11065 WP 001085454.1	IIEIYGPESSGKTTVALHAVAQAQKEGGIAAFIDAEHALDPAYAAALGVNIDELLLSQPDSGEQGLEIAGKLIDSGAVDL	153
NTDB id 499 SM12261 RS08315 WP 078228268.1	IIEIYGPESSGKTTVALHAVAQAQKEGGIAAFIDAEHALDPAYAAALGVNIDELLLSQPDSGEQGLEIAGKLIDSGAVDL	153
NTDB id 453 Spy49 1753c ACI62003.1	IIEIYGPESSGKTTVALHAVAQAQKEGGIAAFIDAEHALDPAYAAALGVNIDELLLAQPDSGEQGLEIAGKLIDSGAVDL	153
NTDB id 408 SMU RS09530 WP 002262392.1	IIEIYGPESSGKTTVALHAVAQAQKDGGIAAFIDAEHALDPAYAAALGVNIDELLLSQPDSGEQGLEIAGKLIDSGAVDL	153
NTDB id 115 BSU 16940 NP 389576.2	IIEVYGPESSGKTTVALHAIAEVQQQGGQAAFIDAEHALDPVYAQKLGVNIEELLLSQPDTGEQALEIAEALVRSGAVDI	138
NTDB id 625 LCA RS02525 WP 011374196.1	IEIYGPESSGKTTVALHAVAEVQKQGGTAAYIDAENAMDPKYATALGVNIDDLLLSQPDTGEQALEIADALVSSGAVDI	140
NTDB id 1313 DR RS12030 WP 010888966.1	ITEIYGPESGGKTTLALAIVAQAQKAGGTCAFIDAEHALDPVYARALGVNTDELLVSQPDNGEQALEIMELLVRSGAIDV	152
NTDB id 1245 Cj1673c YP 002345041.1	IIEIYGPESSGKTTLTLHIIAECQKAGGVCAFIDAEHALDVKYAKNLGVNTDDLYVSQPDFGEQALEIVETIARSGAVDL	139
NTDB id 1214 NCTC11637 00359 SQJ03206.1	IIEIYGPESSGKTTLSLHIIAECQKNGGVCAFIDAEHALDVHYAKRLGVDTENLLVSQPDTGEQALEILETITRSGGIDL	141
NTDB id 1211 C694 RS00795 WP 000963128.1	IIEIYGPESSGKTTLSLHILAECQKNGGVCAFIDAEHALDVHYAKRLGVDTENLLVSQPDTGEQALEILETITRSGGIDL	141
NTDB id 1334 RAOC RS04870 WP 004918236.1	IVEIYGPESSGKTTLTLHAIAEAQKVGGTAAFIDAEHAFDMHYAEKLGHNLEELIVSQPDNGEQALEIADSLIRSGGVDI	148
NTDB id 1389 A4U84 RS07560 WP 021112712.1	IVEIYGPESSGKTTLTLSLIAQAQKGNKTCAFIDAEHALDPVYARNLGVNTDELLISQPDNGEQALEICDALVRSGAVDV	156
NTDB id 1130 NGFG RS03960 WP 003688695.1	IVEIFGPESSGKTTLCLEAVAQCQKNGGVCAFVDAEHAFDPVYARKLGVKVEELYLSQPDTGEQALEICDTLVRSGGIDM	140
NTDB id 1124 NGFG RS03960 WP 003688695.1	IVEIFGPESSGKTTLCLEAVAQCQKNGGVCAFVDAEHAFDPVYARKLGVKVEELYLSQPDTGEQALEICDTLVRSGGIDM	140
NTDB id 1123 OK783 RS03845 WP 003688695.1	IVEIFGPESSGKTTLCLEAVAQCQKNGGVCAFVDAEHAFDPVYARKLGVKVEELYLSQPDTGEQALEICDTLVRSGGIDM	140
NTDB id 1427 RS RS02760 WP 011000510.1	VVEIYGPESSGKTTLTLQVVAEMQKLGGTCAFIDAEHALDVTYADKLGVKVPDLLISQPDTGEQALEIADALVRSGSVDL	148
NTDB id 1165 A1552VC RS01565 WP 000344154.1	IVEIFGPESSGKTTLTLELIAAAQREGKTCAFIDAEHALDPVYAKKLGVNIDELLVSQPDTGEQALEICDALARSGAVDV	139
NTDB id 1142 GTF74 RS11550 WP 000344154.1	IVEIFGPESSGKTTLTLELIAAAQREGKTCAFIDAEHALDPVYAKKLGVNIDELLVSQPDTGEQALEICDALARSGAVDV	139
NTDB id 1205 PSJM300 05545 AFN77183.1	IVEIYGPESSGKTTLTLSVIAAAQRMGSTCAFVDAEHALDPDYAGKLGVNVDDLLVSQPDTGEQALEITDMLVRSNAVDV	139
NTDB id 313049 AhaeAN43 RS07035 WP 005090091.1	ITEIYGPESSGKTTMTLQAIAQCQKTGGTCAFIDAEHALDPEYARKLGVDIDNLLVSQPDNGEQALEIADMLVRSGAIDL	139
NTDB id 1074 ABD1 RS10195 WP 000344167.1	IIEIYGPESSGKTTMTLQAIAQCQKSGGTCAFIDAEHALDPQYARKLGVDIDNLLVSQPDNGEQALEIADMLVRSGAIDL	139
NTDB id 1022 ACIAD RS06390 WP 004925639.1	IVEIYGPESSGKTTMTLQAIAQCQKAGGTCAFIDAEHALDPQYARKLGVDIDNLLVSQPDHGEQALEIADMLVRSGAIDL	139

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NTDB id 600 KW2 RS01765 WP 011675413.1	VVIDSVAALVPKAEIDGEIGDSSVGLQARMMSQAMRKLAGHINKTKTTAIFINQLREKVGVM.FGSPETTPGGRALKFYA	233
NTDB id 269 KZH43 RS08665 WP 001085462.1	VVVDSVAALVPRAEIDGDIGDSHVGLQARMMSQAMRKLGASINKTKTIAIFINQLREKVGVM.FGNPETTPGGRALKFYA	232
NTDB id 228 SPD RS09265 WP 001085462.1	VVVDSVAALVPRAEIDGDIGDSHVGLQARMMSQAMRKLGASINKTKTIAIFINQLREKVGVM.FGNPETTPGGRALKFYA	232
NTDB id 194 SPR RS08825 WP 001085462.1	VVVDSVAALVPRAEIDGDIGDSHVGLQARMMSQAMRKLGASINKTKTIAIFINQLREKVGVM.FGNPETTPGGRALKFYA	232
NTDB id 159 SP RS09750 WP 001085462.1	VVVDSVAALVPRAEIDGDIGDSHVGLQARMMSQAMRKLGASINKTKTIAIFINQLREKVGVM.FGNPETTPGGRALKFYA	232
NTDB id 527 SMSK321 RS11065 WP 001085454.1	VVVDSVAALVPRAEIDGDIGDSHVGLQARMMSQAMRKLGASINKTKTIAIFINQLREKVGVM.FGNPETTPGGRALKFYA	232
NTDB id 499 SM12261 RS08315 WP 078228268.1	VVVDSVAALVPRAEIDGDIGDSHVGLQARMMSQAMRKLGASINKTKTIAIFINQLREKVGVM.FGNPETTPGGRALKFYA	232
NTDB id 453 Spy49 1753c ACI62003.1	VVVDSVAALVPRAEIDGDIGDSHVGLQARMMSQAMRKLSASINKTKTIAIFINQLREKVGVM.FGNPETTPGGRALKFYA	232
NTDB id 408 SMU RS09530 WP 002262392.1	VVVDSVAALVPRAEIDGDIGDSHVGLQARMMSQAMRKLSASINKTKTIAIFINQLREKVGIM.FGNPETTPGGRALKFYS	232
NTDB id 115 BSU 16940 NP 389576.2	VVVDSVAALVPKAEIEGDMGDSHVGLQARLMSQALRKLSGAINKSKTIAIFINQIREKVGVM.FGNPETTPGGRALKFYS	217
NTDB id 625 LCA RS02525 WP 011374196.1	LVVDSVAALVPRAEIEGEMGDAHVGLQARLMSQALRKLSGTINKTKTIALFINQLREKVGVM.FGNPEVTPGGRALKFYS	219
NTDB id 1313 DR RS12030 WP 010888966.1	VVVDSVAALTPRAEIEGDMGDSLPGLQARLMSQALRKLTAILSKTGTAAIFINQVREKIGVM.YGNPETTPGGRALKFYA	231
NTDB id 1245 Cj1673c YP 002345041.1	IVVDSVAALTPKAEIEGDMGDQHVGLQARLMSQALRKLTGIVHKMNTTVIFINQIRMKIGAMGYGTPETTPGGNALKFYA	219
NTDB id 1214 NCTC11637 00359 SQJ03206.1	VVVDSVAALTPKAEIDGDMGDQHVGLQARLMSHALRKITGVLHKMNTTLIFINQIRMKIGMMGYGSPETTPGGNALKFYA	221
NTDB id 1211 C694 RS00795 WP 000963128.1	VVVDSVAALTPKAEIDGDMGDQHVGLQARLMSHALRKITGVLHKMNTTLIFINQIRMKIGMMGYGSPETTPGGNALKFYA	221
NTDB id 1334 RAOC RS04870 WP 004918236.1	VVVDSVAALTPKAEIDGDMGDSKMGLHARLMSQALRKLTSSISKTKCTVIFINQLRDKIGVM.FGSPETTPGGNALKFYA	227
NTDB id 1389 A4U84 RS07560 WP 021112712.1	IIVDSVAALTPKAEIEGDMGDSHMGLQARLMSQALRKLTANIKNTNCIVVFINQIRMKIGVM.FGNPETTPGGNALKFYA	235
NTDB id 1130 NGFG RS03960 WP 003688695.1	VVVDSVAALVPKAEIEGDMGDSHMGLQARLMSQALRKLTGHIKKTNTLVVFINQIRMKIGVM.FGSPETTPGGNALKFYS	219
NTDB id 1124 NGFG RS03960 WP 003688695.1	VVVDSVAALVPKAEIEGDMGDSHVGLQARLMSQALRKLTGHIKKTNTLVVFINQIRMKIGVM.FGSPETTPGGNALKFYS	219
NTDB id 1123 OK783 RS03845 WP 003688695.1	VVVDSVAALVPKAEIEGDMGDSHVGLQARLMSQALRKLTGHIKKTNTLVVFINQIRMKIGVM.FGSPETTPGGNALKFYS	219
NTDB id 1427 RS RS02760 WP 011000510.1	IVIDSVAALVPKAEIEGEMGDALPGLQARLMSQALRKLTGTIKRTNCLVIFINQIRMKIGVM.FGSPETTPGGNALKFYA	227
NTDB id 1165 A1552VC RS01565 WP 000344154.1	IVVDSVAALTPKAEIEGEMGDSHMGLQARMLSQAMRKLTGNLKQSNCMCIFINQIRMKIGVM.FGNPETTPGGNALKFYA	218
NTDB id 1142 GTF74 RS11550 WP 000344154.1	IVVDSVAALTPKAEIEGEMGDSHMGLQARMLSQAMRKLTGNLKQSNCMCIFINQIRMKIGVM.FGNPETTPGGNALKFYA	218
NTDB id 1205 PSJM300 05545 AFN77183.1	IVVDSVAALVPKAEIEGEMGDAHVGLQARLMSQALRKITGNIKNANCLVIFINQIRMKIGVM.FGSPETTPGGNALKFYA	218
NTDB id 313049 AhaeAN43 RS07035 WP 005090091.1	IVVDSVAALTPKAEIEGEMGDSHMGLQARLMSQALRKITGNAKRSNCMVIFINQIRMKIGVM.FGSPETTPGGNALKFYA	218
NTDB id 1074 ABD1 RS10195 WP 000344167.1	IVVDSVAALTPKAEIEGEMGDSHMGLQARLMSQALRKITGNAKRSNCMVIFINQIRMKIGVM.FGSPETTPGGNALKFYA	218
NTDB id 1022 ACIAD RS06390 WP 004925639.1	IVVDSVAALTPKAEIEGEMGDSHMGLQARLMSQALRKITGNAKRSNCMVIFINQIRMKIGVM.FGSPETTPGGNALKFYA	218

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logo                                            SVRLDVRR...KQ.EE..VGNETKIKVWKNKVAPPFKEAEFDIMYGEGISKTGELIDIAVDIJKSGAWYSY

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NTDB id 269 KZH43 RS08665 WP 001085462.1        SVRLDVRGNTQIKGTG.DQKETNVGKETKIKVVKNKVAPPFKEAVVEIMYGEGISKTGELLKIASDLDIIKKAGAWYSYK   311
NTDB id 228 SPD RS09265 WP 001085462.1          SVRLDVRGNTQIKGTG.DQKETNVGKETKIKVVKNKVAPPFKEAVVEIMYGEGISKTGELLKIASDLDIIKKAGAWYSYK   311
NTDB id 194 SPR RS08825 WP 001085462.1          SVRLDVRGNTQIKGTG.DQKETNVGKETKIKVVKNKVAPPFKEAVVEIMYGEGISKTGELLKIASDLDIIKKAGAWYSYK   311
NTDB id 159 SP RS09750 WP 001085462.1           SVRLDVRGNTQIKGTG.DQKETNVGKETKIKVVKNKVAPPFKEAVVEIMYGEGISKTGELLKIASDLDIIKKAGAWYSYK   311
NTDB id 527 SMSK321 RS11065 WP 001085454.1      SVRLDVRGSTQIKGTG.DQKDTNVGKETKIKVVKNKVAPPFKEAFVEIMYGEGISKTGELLKIASDLDIIKKAGAWYSYK   311
NTDB id 499 SM12261 RS08315 WP 078228268.1      SVRLDVRGSTQIKGTG.DQKDTNVGKETKIKVVKNKVAPPFKEAFVEIMYGEGISKTGELLKIASDLDIIKKAGAWYSYK   311
NTDB id 453 Spy49 1753c ACI62003.1              SVRLDVRGTTQIKGTG.DQKDSSIGKETKIKVVKNKVAPPFKVAEVEIMYGEGISRTGELVKIASDLDIIQKAGAWFSYN   311
NTDB id 408 SMU RS09530 WP 002262392.1          SVRLDVRGNTQIKGTG.EQKDSNIGKETKIKVVKNKVAPPFKEAFVEITYGEGISRTGELLKIASDLGIIQKAGAWYSYK   311
NTDB id 115 BSU 16940 NP 389576.2               SVRLEVRRAEQLKQGN.....DVMGNKTKIKVVKNKVAPPFRTAEVDIMYGEGISKEGIIDLGTELDIVQKSGSWYSYE   292
NTDB id 625 LCA RS02525 WP 011374196.1          TVRLEVRRAETIKNGT.....DMIGNRARIKVVKNKVAPPFKVAEVDIMYGQGISRTGELVDMAVEKDIIKSGSWYSYG   294
NTDB id 1313 DR RS12030 WP 010888966.1          SVRLDVRKIGQ....PTKVGNDAVANTVKIKTVKNKVAAPKEVELALVYGKGFDQLSDLVGLAADMDIIKKAGSFYSYG   307
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NTDB id 1214 NCTC11637 00359 SQJ03206.1         SVRIDIRRIAA....L.KQNEQHIGNRAKAKVVKNKVAPPFREAEFDIMFGEGISKEGEIIDYGVKLDIVDKSGAWLSYQ   296
NTDB id 1211 C694 RS00795 WP 000963128.1        SVRIDIRRIAS....L.KQNEQHIGNRAKAKVVKNKVAPPFREAEFDIMFGEGISKEGEIIDYGVKLDIVDKSGAWLSYQ   296
NTDB id 1334 RAOC RS04870 WP 004918236.1        SVRIDIRRSLSDTGKIKDKSGEVIGNRTKVKVVKNKVAPPFKSAEFDIMYGEGVSKVGEILDVATDLDIIQKSSWFSYQ   307
NTDB id 1389 A4U84 RS07560 WP 021112712.1       SVRLDIRRCGV....V.KDGDEIIGSETKVKVVKNKVAPPFREVQFDIMYGEGISRMNELLILAEANGFIKKSGAWFSYD   310
NTDB id 1130 NGFG RS03960 WP 003688695.1        SVRLDIRRTGS....I.KKGEEVLGNETRVKVIKNKVAPPFRQAEFDILYGEGISWEGELIDIGVKNDIINKSGAWYSYN   294
NTDB id 1124 NGFG RS03960 WP 003688695.1        SVRLDIRRTGS....I.KKGEEVLGNETRVKVIKNKVAPPFRQAEFDILYGEGISWEGELIDIGVKNDIINKSGAWYSYN   294
NTDB id 1123 OK783 RS03845 WP 003688695.1       SVRLDIRRTGS....I.KKGEEVLGNETRVKVIKNKVAPPFRQAEFDILYGEGISWEGELIDIGVKNDIINKSGAWYSYN   294
NTDB id 1427 RS RS02760 WP 011000510.1          SVRLDIRRIGS....I.KKGDEVVGNETKVKVVKNKVAPPFREAIFDILYGAGVSREGEIIDLGVEAKVVEKSGAWYSYN   302
NTDB id 1165 A1552VC RS01565 WP 000344154.1     SVRLDIRRTGA....I.KEGDEVVGNETRIKVVKNKIAAPFKEANTQIMYGQGFNREGELIDLGVKHKMVEKSGAWYSYQ   293
NTDB id 1142 GTF74 RS11550 WP 000344154.1       SVRLDIRRTGA....I.KEGDEVVGNETRIKVVKNKIAAPFKEANTQIMYGQGFNREGELIDLGVKHKMVEKSGAWYSYQ   293
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NTDB id 313049 AhaeAN43 RS07035 WP 005090091.1  SVRLDIRRIGA....V.KEGDEITGNETRVKVVKNKMAPPFKEAVFQIMYGKGVDLAVQQDIVQKAGAWYSYQ   293
NTDB id 1074 ABD1 RS10195 WP 000344167.1        SVRLDIRRIGQ....V.KEGDEIVGSETKVKVVKNKMAPPFKEAIFQILYGKGTNQLGELVDLAVQQDIVQKAGAWYSYQ   293
NTDB id 1022 ACIAD RS06390 WP 004925639.1       SVRLDIRRIGQ....V.KEGDEIIGSETKVKVVKNKMAPPFREAIFQILYGKGVNQLGELVDLAVQQNIVQKAGAWYSYQ   293
consensus                                       *!!***!*       *  **  *****!**!!!!*!*!!***  ****!*!*** **** ** *** ! !****!
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NTDB id 600 KW2 RS01765 WP 011675413.1      DEKIGQGAEKAKNYLKEHQDVFDEIDHKVRAAHGLLDDSEVAETEEETTASKTKAKAKKEEKAVETEEIELELED.....   387
NTDB id 269 KZH43 RS08665 WP 001085462.1     DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFGLIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE   385
NTDB id 228 SPD RS09265 WP 001085462.1       DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFGLIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE   385
NTDB id 194 SPR RS08825 WP 001085462.1       DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFGLIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE   385
NTDB id 159 SP RS09750 WP 001085462.1        DEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFGLIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE   385
NTDB id 527 SMSK321 RS11065 WP 001085454.1   DEKIGQGSENAKKYLADNPEVFAEIDHQVRVQFGLIDGEEAAE....SKKDE....VAQVDS..VNEEVTLDLGDELEIE   381
NTDB id 499 SM12261 RS08315 WP 078228268.1   DEKIGQGSENAKKYLADNPEIFDEIDHQVRVQFGLIDGEEIAE....SKKDE....AAQVDS..VNEEVTLDLGDELEIE   381
NTDB id 453 Spy49 1753c ACI62003.1           GEKIGQGSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEEESAMAVASE.......E.......TDDLALDLDNGIEIE   377
NTDB id 408 SMU RS09530 WP 002262392.1       GEKIGQGSENAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEKQSSKEK.......ETDEK..ADKNLVLELDDTIELE   382
NTDB id 115 BSU 16940 NP 389576.2            EERLGQGRENAKQFLKENKDIMLMIQEQIREHYGLDNNGVVQQQAEETQEEL.EFE.E.......................   348
NTDB id 625 LCA RS02525 WP 011374196.1       SERIGQGRENAKNYLADHEDVEDEVRLKVRAAYGISDVPEEDLPTTEDE.QI.NILPDDST......EE............   355
NTDB id 1313 DR RS12030 WP 010888966.1       DERIGQGKEKTIAYIAERPEMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPEAAEA.....................   363
NTDB id 1245 Cj1673c YP 002345041.1          DKKLGQGRENSKAFLKENPEIADEITKAIQNSMGIEG.....MISGSEDDE..GEE.....................   343
NTDB id 1214 NCTC11637 00359 SQJ03206.1      DKKLGQGRENAKALLKEDKALADEITLKIKESIGSNE....EIMPLPDEP..LEEME....................   347
NTDB id 1211 C694 RS00795 WP 000963128.1     DKKLGQGRENAKALLKEDKALADEITLKIKESIGSNE....EIMPLPDEP..LEEME....................   347
NTDB id 1334 RAOC RS04870 WP 004918236.1     DTKLGQGRDTVKELIKDNPELAEELEAKIREKLQAK.....................................   343
NTDB id 1389 A4U84 RS07560 WP 021112712.1    GTKIGQGKNNAVKWLKENPEIADKLEQEIRNTLASNP.....NSVLKATASDSNDESDV..EEFSEEEL.........   372
NTDB id 1130 NGFG RS03960 WP 003688695.1     GAKIGQGKDNVRVWLKENPEISDEIDAKIRALNGVEM.....HITEGTQDE......TD..GERPEE..........   348
NTDB id 1124 NGFG RS03960 WP 003688695.1     GAKIGQGKDNVRVWLKENPEISDEIDAKIRALNGVEM.....HITEGTQDE......TD..GERPEE..........   348
NTDB id 1123 OK783 RS03845 WP 003688695.1    GAKIGQGKDNVRVWLKENPEISDEIDAKIRALNGVEM.....HITEGTQDE......TD..GERPEE..........   348
NTDB id 1427 RS RS02760 WP 011000510.1       GERIGQGRDNCREFLRENAELAREIENKVREHLGVTP.....MGAVTLAE.EVEED....................   352
NTDB id 1165 A1552VC RS01565 WP 000344154.1  GDKIGQGKANACKYLKENPEIAKTLDKKLREMLLNPE...NMQLIAETSSAADDVEF..GAVPEEF..........   354
NTDB id 1142 GTF74 RS11550 WP 000344154.1    GDKIGQGKANACKYLKENPEIAKTLDKKLREMLLNPE...NMQLIAETSSAADDVEF..GAVPEEF..........   354
NTDB id 1205 PSJM300 05545 AFN77183.1        GSKIGQGKANAAKYLEENPEIGQEIEQQIRAKLLVTS....ANTKATPVAEDLADAD....I.............   347
NTDB id 313049 AhaeAN43 RS07035 WP 005090091.1  GNKIGQGKNNVIRHFEENPQMAQEIEKLIREQLLTKN.....NNEAA.AVEED....ELD....LLES.........   347
NTDB id 1074 ABD1 RS10195 WP 000344167.1     GNKIGQGKNNVIRYFEENNQIAEEIERNIREQLLTTG.....TNGAV.QI.EDEEEPD....LLLES.........   349
NTDB id 1022 ACIAD RS06390 WP 004925639.1    GNKIGQGKNNVIRYLEENPQISTEIEAVIREQLLTKA.....SDQTA.AHDETEEEPD....LLES..........   349

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logo

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NTDB id 600 KW2 RS01765 WP 011675413.1          ...    387
NTDB id 269 KZH43 RS08665 WP 001085462.1        IEE    388
NTDB id 228 SPD RS09265 WP 001085462.1          IEE    388
NTDB id 194 SPR RS08825 WP 001085462.1          IEE    388
NTDB id 159 SP RS09750 WP 001085462.1           IEE    388
NTDB id 527 SMSK321 RS11065 WP 001085454.1      IEE    384
NTDB id 499 SM12261 RS08315 WP 078228268.1      IEE    384
NTDB id 453 Spy49 1753c ACI62003.1              D..    378
NTDB id 408 SMU RS09530 WP 002262392.1          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 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 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 313049 AhaeAN43 RS07035 WP 005090091.1  ...    347
NTDB id 1074 ABD1 RS10195 WP 000344167.1        ...    349
NTDB id 1022 ACIAD RS06390 WP 004925639.1       ...    349
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consensus

X non conserved
X similar
X $\geq 50\%$ conserved