

logo



NTDB id 293421 LPEG9 RS14365 WP 016387612.1	SVRLDVRGQVVKDGGK...DVGFA	SVRLDVRGQVVKDGGK...DVGFA	307
NTDB id 600 KW2 RS01765 WP 011675413.1	SVRLDVRGSTKIEEGS.GDNKTQ	SVRLDVRGSTKIEEGS.GDNKTQ	308
NTDB id 269 KZH43 RS08665 WP 001085462.1	SVRLDVRGNTQIKGTG.DQKETN	SVRLDVRGNTQIKGTG.DQKETN	307
NTDB id 228 SPD RS09265 WP 001085462.1	SVRLDVRGNTQIKGTG.DQKETN	SVRLDVRGNTQIKGTG.DQKETN	307
NTDB id 194 SPR RS08825 WP 001085462.1	SVRLDVRGNTQIKGTG.DQKETN	SVRLDVRGNTQIKGTG.DQKETN	307
NTDB id 159 SP RS09750 WP 001085462.1	SVRLDVRGNTQIKGTG.DQKETN	SVRLDVRGNTQIKGTG.DQKETN	307
NTDB id 499 SM12261 RS08315 WP 078228268.1	SVRLDVRGNTQIKGTG.DQKETN	SVRLDVRGNTQIKGTG.DQKETN	307
NTDB id 527 SMSK321 RS11065 WP 001085454.1	SVRLDVRGNTQIKGTG.DQKETN	SVRLDVRGNTQIKGTG.DQKETN	307
NTDB id 453 Spy49 1753c ACI62003.1	SVRLDVRGNTQIKGTG.DQKDS	SVRLDVRGNTQIKGTG.DQKDS	307
NTDB id 408 SMU RS09530 WP 002262392.1	SVRLDVRGNTQIKGTG.EQKDS	SVRLDVRGNTQIKGTG.EQKDS	307
NTDB id 625 LCA RS02525 WP 011374196.1	TVRLEVRRAETIKNGT...DM	TVRLEVRRAETIKNGT...DM	290
NTDB id 115 BSU 16940 NP 389576.2	SVRLEVRRAEQLKQGN...D	SVRLEVRRAEQLKQGN...D	288
NTDB id 1313 DR RS12030 WP 010888966.1	SVRLDVRKIGQ...PTKVGND	SVRLDVRKIGQ...PTKVGND	303
NTDB id 1245 Cj1673c YP 002345041.1	SVRLDVRKIVAT...L.KQNE	SVRLDVRKIVAT...L.KQNE	290
NTDB id 1214 NCTC11637 00359 SQJ03206.1	SVRIDIRRIAA...L.KQNE	SVRIDIRRIAA...L.KQNE	292
NTDB id 1211 C694 RS00795 WP 000963128.1	SVRIDIRRIAS...L.KQNE	SVRIDIRRIAS...L.KQNE	292
NTDB id 1334 RAOC RS04870 WP 004918236.1	SVRIDIRRLSDTGKIKDKSGE	SVRIDIRRLSDTGKIKDKSGE	303
NTDB id 1389 A4U84 RS07560 WP 021112712.1	SVRLDIRRCGV...V.KGD	SVRLDIRRCGV...V.KGD	306
NTDB id 1130 NGFG RS03960 WP 003688695.1	SVRLDIRRTGS...I.KKG	SVRLDIRRTGS...I.KKG	290
NTDB id 1124 NGFG RS03960 WP 003688695.1	SVRLDIRRTGS...I.KKG	SVRLDIRRTGS...I.KKG	290
NTDB id 1123 OK783 RS03845 WP 003688695.1	SVRLDIRRTGS...I.KKG	SVRLDIRRTGS...I.KKG	290
NTDB id 1427 RS RS02760 WP 011000510.1	SVRLDIRRIGS...I.KKG	SVRLDIRRIGS...I.KKG	298
NTDB id 1165 A1552VC RS01565 WP 000344154.1	SVRLDIRRTGA...I.KEG	SVRLDIRRTGA...I.KEG	289
NTDB id 1142 GTF74 RS11550 WP 000344154.1	SVRLDIRRTGA...I.KEG	SVRLDIRRTGA...I.KEG	289
NTDB id 1205 PSJM300 05545 AFN77183.1	SVRLDIRRTGA...V.KEG	SVRLDIRRTGA...V.KEG	289
NTDB id 1022 ACIAD RS06390 WP 004925639.1	SVRLDIRRIGQ...V.KEG	SVRLDIRRIGQ...V.KEG	289
NTDB id 1074 ABD1 RS10195 WP 000344167.1	SVRLDIRRIGQ...V.KEG	SVRLDIRRIGQ...V.KEG	289
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NTDB id 293421 LPEG9 RS14365 WP 016387612.1	YNYGKERLIGQGLVNASHYLD EHPEISKEIEEK TREQVAPKKPEETDKQDDKKDDQASGDKKTEKTTTTDTEKKADKKDD	387
NTDB id 600 KW2 RS01765 WP 011675413.1	FAYNDEKIGQGA EKA KNYLKEHQDV FDEIDHKVRAAHG LDDSEVAETEETETASKTKAKAKKEEKAVETEEIELE.LED	387
NTDB id 269 KZH43 RS08665 WP 001085462.1	YSYKDEKIGQGS ENAKKYLAEHPEIF DEIDKQVRSKFG LIDGEEVSEQDTENKKDE ...PKKEEA..VNEEVPLD.LGD	380
NTDB id 228 SPD RS09265 WP 001085462.1	YSYKDEKIGQGS ENAKKYLAEHPEIF DEIDKQVRSKFG LIDGEEVSEQDTENKKDE ...PKKEEA..VNEEVPLD.LGD	380
NTDB id 194 SPR RS08825 WP 001085462.1	YSYKDEKIGQGS ENAKKYLAEHPEIF DEIDKQVRSKFG LIDGEEVSEQDTENKKDE ...PKKEEA..VNEEVPLD.LGD	380
NTDB id 159 SP RS09750 WP 001085462.1	YSYKDEKIGQGS ENAKKYLAEHPEIF DEIDKQVRSKFG LIDGEEVSEQDTENKKDE ...PKKEEA..VNEEVPLD.LGD	380
NTDB id 499 SM12261 RS08315 WP 078228268.1	YSYKDEKIGQGS ENAKKYLADNPEIF DEIDHQV RVVQFGLIDGEEIAES ...KKDE...AAQVDS..VNEEVTLD.LGD	376
NTDB id 527 SMSK321 RS11065 WP 001085454.1	YSYKDEKIGQGS ENAKKYLADNPEVFAEIDHQV RVVQFGLIDGEEAAES...KKDE...VAQVDS..VNEEVTLD.LGD	376
NTDB id 453 Spy49 1753c ACI62003.1	FSYNGEKIGQGS ENAKRYLADHPQLF DEIDR KVRVKFGLLEEESEESAMAVASE ...E...TDDLALD.LDN	372
NTDB id 408 SMU RS09530 WP 002262392.1	YSYNGEKIGQGS ENAKKFLADNPEIF DDIDHKV RVVQYGLIEEDNTEEKQSSKEK ...ETDEK..ADKNLVLE.LDD	377
NTDB id 625 LCA RS02525 WP 011374196.1	YSYGSERIGQGR ENAKNYLADHEDVE DEVRLK VRAAYGISDVP PEEDLPTTEDE.QI.NILPDDST...EE.....	355
NTDB id 115 BSU 16940 NP 389576.2	YSYEEERLIGQGR ENAKQFLKENK DIMLMIQEQ TREHYGLDNNGV VVQQAEETQEEL.EFE.E.....	348
NTDB id 1313 DR RS12030 WP 010888966.1	YSYGDERIGQG KEKTIA YIAERP EME QEI RD RVMAAIRAGNAGEAPALAPAPAAPEAAEA.....	363
NTDB id 1245 Cj1673c YP 002345041.1	FSYKDKKLGQGR ENSKAFLKENPEI ADEITKAI QNSMGT EGMI...SGSEDDE..GEE.....	343
NTDB id 1214 NCTC11637 00359 SQJ03206.1	LSYQDKKLGQGR ENAKALLKEDKAL ADEITL KIKESIGS NEEI...MPLPDEP..LEEME.....	347
NTDB id 1211 C694 RS00795 WP 000963128.1	LSYQDKKLGQGR ENAKALLKEDKAL ADEITL KIKESIGS NEEI...MPLPDEP..LEEME.....	347
NTDB id 1334 RAOC RS04870 WP 004918236.1	FSYQDTKLGQGR DTVKELIKDNPEL AEELEAK TREKLQAK	343
NTDB id 1389 A4U84 RS07560 WP 021112712.1	FSYDGTKIGQGG KNNAVKWLKENPEI ADKLEQE IRNTLASNPNS ...VLKATASDSNDESVDVEE..FSEEEL.....	372
NTDB id 1130 NGFG RS03960 WP 003688695.1	YSYNGAKIGQGD NVRVWLKENPEIS DEIDAK IRALNGVEMHI ...TEGT...QDETGE..RPEE.....	348
NTDB id 1124 NGFG RS03960 WP 003688695.1	YSYNGAKIGQGD NVRVWLKENPEIS DEIDAK IRALNGVEMHI ...TEGT...QDETGE..RPEE.....	348
NTDB id 1123 OK783 RS03845 WP 003688695.1	YSYNGAKIGQGD NVRVWLKENPEIS DEIDAK IRALNGVEMHI ...TEGT...QDETGE..RPEE.....	348
NTDB id 1427 RS RS02760 WP 011000510.1	YSYNGERIGQGR DNCREFLRENAELAREI ENK VREHLGV TPMG...AV.TLAEVEED.....	352
NTDB id 1165 A1552VC RS01565 WP 000344154.1	YSYNGDKIGQGG KANACKYLKENPEI AKTLD KKLREMLLN PENM...QLIAETSSAADDVEFGA..VPEEF.....	354
NTDB id 1142 GTF74 RS11550 WP 000344154.1	YSYNGDKIGQGG KANACKYLKENPEI AKTLD KKLREMLLN PENM...QLIAETSSAADDVEFGA..VPEEF.....	354
NTDB id 1205 PSJM300 05545 AFN77183.1	YAYKGSKIGQGG KANAACKYLEENPEI GQIE EQ TRAKLL V TSAN...TKATPVAEDLADADI.....	347
NTDB id 1022 ACIAD RS06390 WP 004925639.1	YSYQGNKIGQGG KNNVIRYLEENPQ ISTEIE AV TREQLLTKASD...QTAA.HDETEEEPDLL...ES.....	349
NTDB id 1074 ABD1 RS10195 WP 000344167.1	YSYQGNKIGQGG KNNVIRYFEENNQ IAE E IERNIREQLLTTGTN...GAVQ.I.EDEEEPDLL...LES.....	349
consensus	***! **!!!***** ** * *** ***** ** **	

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NTDB id	Accession	Sequence	Score
293421	LPEG9 RS14365 WP 016387612.1	PASFGVDRLV	397
600	KW2 RS01765 WP 011675413.1	387
269	KZH43 RS08665 WP 001085462.1	ELEIEIEE..	388
228	SPD RS09265 WP 001085462.1	ELEIEIEE..	388
194	SPR RS08825 WP 001085462.1	ELEIEIEE..	388
159	SP RS09750 WP 001085462.1	ELEIEIEE..	388
499	SM12261 RS08315 WP 078228268.1	ELEIEIEE..	384
527	SMSK321 RS11065 WP 001085454.1	ELEIEIEE..	384
453	Spy49 1753c ACI62003.1	GIEIED....	378
408	SMU RS09530 WP 002262392.1	TIELED....	383
625	LCA RS02525 WP 011374196.1	355
115	BSU 16940 NP 389576.2	348
1313	DR RS12030 WP 010888966.1	363
1245	Cj1673c YP 002345041.1	343
1214	NCTC11637 00359 SQJ03206.1	347
1211	C694 RS00795 WP 000963128.1	347
1334	RAOC RS04870 WP 004918236.1	343
1389	A4U84 RS07560 WP 021112712.1	372
1130	NGFG RS03960 WP 003688695.1	348
1124	NGFG RS03960 WP 003688695.1	348
1123	OK783 RS03845 WP 003688695.1	348
1427	RS RS02760 WP 011000510.1	352
1165	A1552VC RS01565 WP 000344154.1	354
1142	GTF74 RS11550 WP 000344154.1	354
1205	PSJM300 05545 AFN77183.1	347
1022	ACIAD RS06390 WP 004925639.1	349
1074	ABD1 RS10195 WP 000344167.1	349

consensus

- X non conserved
- X similar
- X ≥ 50% conserved