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NTDB id 600 KW2 RS01765 WP 011675413.1	DEKIGGQAEKAKNYLKEHQDVFEIDHKVRAAHGILDDSEVAETEETTASKTKAKAKKEEKAVETEEIELELED.....	387
NTDB id 269 KZH43 RS08665 WP 001085462.1	DEKIGGQSENAKKYLAEHPEIFDEIDKQVRSKFCIDGEEVSEQDTEENKKDEPKK.....EEAVNEEVPLDLGDELEIE	385
NTDB id 228 SPD RS09265 WP 001085462.1	DEKIGGQSENAKKYLAEHPEIFDEIDKQVRSKFCIDGEEVSEQDTEENKKDEPKK.....EEAVNEEVPLDLGDELEIE	385
NTDB id 194 SPR RS08825 WP 001085462.1	DEKIGGQSENAKKYLAEHPEIFDEIDKQVRSKFCIDGEEVSEQDTEENKKDEPKK.....EEAVNEEVPLDLGDELEIE	385
NTDB id 159 SP RS09750 WP 001085462.1	DEKIGGQSENAKKYLAEHPEIFDEIDKQVRSKFCIDGEEVSEQDTEENKKDEPKK.....EEAVNEEVPLDLGDELEIE	385
NTDB id 499 SM12261 RS08315 WP 078228268.1	DEKIGGQSENAKKYLADNPEIFDEIDHQVVRVQFCIDGEEIAES...KKDEAAQ.....VDSVNNEEVTLDLGDELEIE	381
NTDB id 527 SMSK321 RS11065 WP 001085454.1	DEKIGGQSENAKKYLADNPEVFAEIDHQVVRVQFCIDGEEAAES...KKDEVAQ.....VDSVNNEEVTLDLGDELEIE	381
NTDB id 1147517 CKV85 RS10955 WP 029691479.1	GEKIGGQSENAKKYLAENPEVFEIDQKVRVHFCIEAPEEADKADAADKDEQAS.....ENSQADELVLDDLDTIEIE	385
NTDB id 408 SMU RS09530 WP 002262392.1	GEKIGGQSENAKKFLADNPEIFDDIDHKVVRVQFCIEEDNTEEKQSSKE...KET.....DEKADKNLVLELDDTIEIE	382
NTDB id 453 Spy49 1753c ACI62003.1	GEKIGGQSENAKRYLADHPQLFDEIDRKVRVKFCILEESEESAMAVAS...EE.....TDDLALDLNNGIEIE	377
NTDB id 625 LCA RS02525 WP 011374196.1	SERIGGQRENAKNYLADHEDVEDEVRLKVRVAAAYGIVSDVPEEDLPTTEDE.QINILPDDS.....TEE.....	355
NTDB id 115 BSU 16940 NP 389576.2	EERIGGQRENAKQFLKENKDIIMLMITQEQIREHYCIDNNGVVQQAEETQEELEFE.E.....	348
NTDB id 1313 DR RS12030 WP 010888966.1	DERIGGQKEKTIAYIAERPMEQEIRDRVMAATRAGNAGEAPALAPAPAAPEAA..EA.....	363
NTDB id 1245 Cj1673c YP 002345041.1	DKKIGGQRENSKAEIKENPEIADAITKATIQNSMCI EGMI...SGSEDD..EG..EE.....	343
NTDB id 1211 C694 RS00795 WP 000963128.1	DKKIGGQRENAKALLKEDKALADEITLKIRESICSNEEI...MPLPDE..PL..EEME.....	347
NTDB id 1214 NCTC11637 00359 SQJ03206.1	DKKIGGQRENAKALLKEDKALADEITLKIRESICSNEEI...MPLPDE..PL..EEME.....	347
NTDB id 1334 RAOC RS04870 WP 004918236.1	DTKIGGQRDITVKELTKDNPEIAEELAKIREKLQAK.....	343
NTDB id 1389 A4U84 RS07560 WP 021112712.1	GTKIGGQKNNAVKWLKENPEIADKLEQEIIRNTLASNPNS...VLKATASDSN..DESDVEEFSEEL.....	372
NTDB id 1130 NGFG RS03960 WP 003688695.1	GAKIGGQKDNVRVWLKENPEISDEIDAKIRALNCVEMHI...TEGTQDETGDG..ER.....PEE.....	348
NTDB id 1124 NGFG RS03960 WP 003688695.1	GAKIGGQKDNVRVWLKENPEISDEIDAKIRALNCVEMHI...TEGTQDETGDG..ER.....PEE.....	348
NTDB id 1123 OK783 RS03845 WP 003688695.1	GAKIGGQKDNVRVWLKENPEISDEIDAKIRALNCVEMHI...TEGTQDETGDG..ER.....PEE.....	348
NTDB id 1427 RS RS02760 WP 011000510.1	GERIGGQRDNCREELRENAEIAREIENKVRHLGVTPMG...AVTLA..EEVE..ED.....	352
NTDB id 1165 A1552VC RS01565 WP 000344154.1	GDKIGGQKANACKYLKENPEIAKTLDDKKLREMLLNPNEM...QLIAETSSAA..DDVEFGAVPEEF.....	354
NTDB id 1142 GTF74 RS11550 WP 000344154.1	GDKIGGQKANACKYLKENPEIAKTLDDKKLREMLLNPNEM...QLIAETSSAA..DDVEFGAVPEEF.....	354
NTDB id 1022 ACIAD RS06390 WP 004925639.1	GNKIGGQKNNVIRYLEENPCISTEIEAVIREQLLTKASD...QTAA..HDETE..EEPDLL..ES.....	349
NTDB id 1074 ABD1 RS10195 WP 000344167.1	GNKIGGQKNNVIRYFEENPCIAEIEIRNIREQLLTTGTN...GAVQ..I..EDE..EEPDLL..LES.....	349
NTDB id 1205 PSJM300 05545 AFN77183.1	GSKIGGQKANAAKYLEENPEIQEIEQQIRAKLLVTSAN...TKATPVAEDL..ADADI.....	347
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consensus

↓EE

⊠ non conserved
⊠ similar
⊠ ≥ 50% conserved