

logo

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
 NTDB id 408 SMU RS09530 WP 002262392.1
 NTDB id 453 Spy49 1753c ACI62003.1
 NTDB id 115 BSU 16940 NP 389576.2
 NTDB id 625 LCA RS02525 WP 011374196.1
 NTDB id 1313 DR RS12030 WP 010888966.1
 NTDB id 1245 Cj1673c YP 002345041.1
 NTDB id 1211 C694 RS00795 WP 000963128.1
 NTDB id 1214 NCTC11637 00359 SQJ03206.1
 NTDB id 1334 RAOC RS04870 WP 004918236.1
 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 173194 PG2T RS09930 WP 068804721.1
 NTDB id 1427 RS RS02760 WP 011000510.1
 NTDB id 1389 A4U84 RS07560 WP 021112712.1
 NTDB id 1165 A1552VC RS01565 WP 000344154.1
 NTDB id 1142 GTF74 RS11550 WP 000344154.1
 NTDB id 1022 ACIAD RS06390 WP 004925639.1
 NTDB id 1074 ABD1 RS10195 WP 000344167.1
 NTDB id 1205 PSJM300 05545 AFN77183.1

consensus



DEKIGQGA EKAKNYLKEHQDVFD EIDHKVRAAHGL LDDSEVAETEEETTASKTKAKAKKEEKAVETEEIELELED 387
 DEKIGQGS ENAKKYLAEHPEIFDE IDKQVRSKFG LIDGEEVSEQD TENKKDE PKKEEA . . VNEEVPLDLGDELEIE 385
 DEKIGQGS ENAKKYLAEHPEIFDE IDKQVRSKFG LIDGEEVSEQD TENKKDE PKKEEA . . VNEEVPLDLGDELEIE 385
 DEKIGQGS ENAKKYLAEHPEIFDE IDKQVRSKFG LIDGEEVSEQD TENKKDE PKKEEA . . VNEEVPLDLGDELEIE 385
 DEKIGQGS ENAKKYLAEHPEIFDE IDKQVRSKFG LIDGEEVSEQD TENKKDE PKKEEA . . VNEEVPLDLGDELEIE 385
 DEKIGQGS ENAKKYLAADNPEIFDE IDHQVVRVQFGL IDGEEIAES KKDE AAQVDS . . VNEEVTLDLGDELEIE 381
 DEKIGQGS ENAKKYLAADNPEVFAE IDHQVVRVQFGL IDGEEAAES KKDE VAQVDS . . VNEEVTLDLGDELEIE 381
 GEKIGQGS ENAKKFLADNPEIFDD IDHKVVRVQYGL IEEDNTEEKQSSKEK ETDEK . . ADKNLVLELDDTIELE 382
 GEKIGQGS ENAKRYLADHPQLFDE IDRKVVRVQFGL LEESEEEESAMAVASE E TDDLALDLDNGIEIE 377
 EERLGGGREN AKQFLKENKDIMLMIQEQ IREHYGL DNNGVVQQAEETQEEL . EFE . E 348
 SERIGGGREN AKNYLADHEDVEDEVRLKVR AAYGISDVPPEEDLPTTEDE . QI . NILPDDST EE 355
 DERIGGGKEK TIAYIAERPMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPEAAEA 363
 DKKLGQGRENSK AFLKENPEIADEITKAIQNSMGI EGMI SGSEDDGEE 343
 DKKLGQGRENAK ALLKEDKALADEITLKI KESIGSNEEI MPLPDEPLEEME 347
 DKKLGQGRENAK ALLKEDKALADEITLKI KESIGSNEEI MPLPDEPLEEME 347
 DTKLGQGRD TVKELIKDNPELAEELEAK IREKLQAK 343
 GAKIGQGGKDNVRVW LKENPEISDEIDAK ITRALNGVEMHI TEGTQDET DGERP EE 348
 GAKIGQGGKDNVRVW LKENPEISDEIDAK ITRALNGVEMHI TEGTQDET DGERP EE 348
 GAKIGQGGKDNVRVW LKENPEISDEIDAK ITRALNGVEMHI TEGTQDET DGERP EE 348
 GERIGQGRDNVRDFLRANPETAE AIEAL IRSRMLPARKV AAEAASPV 341
 GERIGQGRDNCR EFLRENAELAREIENKVREHLG VTPMG AVTLAAEEVEED 352
 GTKIGQGGKNN AVKWLKENPEIADKLEQEIRNTL ASNPN VLKATASDSNDES DVE . . EFSEEEEL 372
 GDKIGQGGKANACKY LKENPEIAKTLDKKLREMLLN PENM QLIAETSSAADDVEFG . . AVPEEF 354
 GDKIGQGGKANACKY LKENPEIAKTLDKKLREMLLN PENM QLIAETSSAADDVEFG . . AVPEEF 354
 GNKIGQGGKNNVIRY LEENPQISTEIEAV IREQLLTKASD QTAA . HDTEEEEPDLL ES 349
 GNKIGQGGKNNVIRY FEENNQIAEIEIERN IREQLLTTGTN GAVQ . I . EDEEEPDLL LES 349
 GSKIGQGGKANAAKY LEENPEIGQEIEQQ IRAKLLVTSAN TKATPVAEDLADADI 347

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NTDB id 269 KZH43 RS08665 WP 001085462.1	IEE	388
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NTDB id 159 SP RS09750 WP 001085462.1	IEE	388
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NTDB id 527 SMSK321 RS11065 WP 001085454.1	IEE	384
NTDB id 408 SMU RS09530 WP 002262392.1	D..	383
NTDB id 453 Spy49 1753c ACI62003.1	D..	378
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 1211 C694 RS00795 WP 000963128.1	...	347
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NTDB id 1124 NGFG RS03960 WP 003688695.1	...	348
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	...	349
NTDB id 1205 PSJM300 05545 AFN77183.1	...	347

consensus

 non conserved
 similar
 ≥ 50% conserved