

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 288712 DC017 RS09210 WP 173956424.1
 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 ***!***!* * ** *****!*****!*****!***** *****!***** ***** ** ***** ! !*****!

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consensus



DEKIGQGA EKAKNYLKEHQDVFD EIDHKVRAAHG LLDSEVAETEEETT . ASKTKAKAKKEEKAVETEEIELELED 387
 DEKIGQGS ENAKKYLAEHPEIFDE IDKQVRSKFG LIDGEEVSEQDTENK . KDE PKKEEA . . VNEEVPLDLGDELEI 384
 DEKIGQGS ENAKKYLAEHPEIFDE IDKQVRSKFG LIDGEEVSEQDTENK . KDE PKKEEA . . VNEEVPLDLGDELEI 384
 DEKIGQGS ENAKKYLAEHPEIFDE IDKQVRSKFG LIDGEEVSEQDTENK . KDE PKKEEA . . VNEEVPLDLGDELEI 384
 DEKIGQGS ENAKKYLAEHPEIFDE IDKQVRSKFG LIDGEEVSEQDTENK . KDE PKKEEA . . VNEEVPLDLGDELEI 384
 DEKIGQGS ENAKKYLAADNPEIFDE IDHQVVRVQFGL IDGEEIAES K . KDE AAQVDS . . VNEEVTLDLGDELEI 380
 DEKIGQGS ENAKKYLAADNPEVFAE IDHQVVRVQFGL IDGEEAAES K . KDE VAQVDS . . VNEEVTLDLGDELEI 380
 GEKIGQGS ENAKKFLADNPEIFDD IDHKVVRVQYGL IEEDNTEEKQSSKE . K ETDEK . . ADKNLVLELDDTIEL 381
 GEKIGQGS ENAKRYLADHPQLFDE IDRKVVRVQFGL LEESEEEESAMAVAS . E E TDDLALDLNGIEI 376
 EERLGGGREN AKQFLKENKDIMLMIQEQ IREHYGL DNNGVVQQAEETQ . EEL . EFE . E 348
 SERIGGGREN AKNYLADHEDVEDEVRLKVR AAYGISDVPPEEDLPTTEDE . . QI . NILPDDST EE 355
 DERIGGGKEK TIAYIAERPMEQEIRDRVMAA IIRAGNAGEAPALAPAPAAP EAAEA 363
 DKKLGQGRENSK AFLKENPEIADEITKA IQNSMG IEGMISG . . SEDDEGEE 343
 DKKLGQGRENAK ALLKEDKALADEITLKI KESIGSNEEIMP . . LPDEPLEEME 347
 DKKLGQGRENAK ALLKEDKALADEITLKI KESIGSNEEIMP . . LPDEPLEEME 347
 DTKLGQGRD TVKELIKDNPELAELEAK IREKLQAK 343
 GAKIGQGGKDNVRVW LKENPEISDEIDAK ITRALNGVEMHITE . . GTQDET DGERPEE 348
 GAKIGQGGKDNVRVW LKENPEISDEIDAK ITRALNGVEMHITE . . GTQDET DGERPEE 348
 GAKIGQGGKDNVRVW LKENPEISDEIDAK ITRALNGVEMHITE . . GTQDET DGERPEE 348
 GDRIGQGGKDNVREFLKENPEIAKD IEAK IREKLGVKAGSAV . . VTDVLGEE EEEVE 363
 GERIGQGRDNCR EFLRENAELAREI ENKVREHLG VTPMGAV . . TLAE EEEV EED 352
 GTKIGQGGKNNAVKWLKENPEIADKLEQE IIRNTL ASNPN SVL . . KATASDSNDES DVE EFSE EEL 372
 GDKIGQGGKANACKYLKENPEIAKTL DKKLREMLLN PENMQL . . IAETSSAAD DVEFG AVPEEF 354
 GDKIGQGGKANACKYLKENPEIAKTL DKKLREMLLN PENMQL . . IAETSSAAD DVEFG AVPEEF 354
 GNKIGQGGKNNVIRY LEENPQISTE IEAV IREQLLTKASDQT . . AA . HDETE EEPDLL ES 349
 GNKIGQGGKNNVIRY FEENNQIAE IEIERN IREQLLTTGTNGA . . VQ . I . EDEE EEPDLL LES 349
 GSKIGQGGKANAAKY LEENPEIGQEIEQQ IIRAKLLVTSANTK . . ATPVAEDLADADI 347

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NTDB id 600 KW2 RS01765 WP 011675413.1	387
NTDB id 269 KZH43 RS08665 WP 001085462.1	EIEE	388
NTDB id 228 SPD RS09265 WP 001085462.1	EIEE	388
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NTDB id 408 SMU RS09530 WP 002262392.1	ED..	383
NTDB id 453 Spy49 1753c ACI62003.1	ED..	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
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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