









logo



NTDB id 92852 MWL86 RS00230 WP 279612170.1	KGETIAQGGKDRLAFLDKNPEMLAEIRKTVLDQAIEGLGF...YTAPEKSDDESDVDSGHGVDIDEEVIDLDISDESM	376
NTDB id 600 KW2 RS01765 WP 011675413.1	NDEKIGQGAEKAKNYLKEHQDVFDEIDHKVRAAHGLDDSEVAETEETETASKTKAKAKKE.EKAVETEEIELELED...	387
NTDB id 269 KZH43 RS08665 WP 001085462.1	KDEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFGIDGEEVSEQDTENKKDE...PKKE.EA..VNEEVPLDLGDELE	383
NTDB id 228 SPD RS09265 WP 001085462.1	KDEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFGIDGEEVSEQDTENKKDE...PKKE.EA..VNEEVPLDLGDELE	383
NTDB id 194 SPR RS08825 WP 001085462.1	KDEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFGIDGEEVSEQDTENKKDE...PKKE.EA..VNEEVPLDLGDELE	383
NTDB id 159 SP RS09750 WP 001085462.1	KDEKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFGIDGEEVSEQDTENKKDE...PKKE.EA..VNEEVPLDLGDELE	383
NTDB id 527 SMSK321 RS11065 WP 001085454.1	KDEKIGQGSENAKKYLDNPEVFAEIDHQVVRVQFGLIDGEEAAE...SKKDE...VAQV.DS..VNEEVTLDLGDELE	379
NTDB id 499 SM12261 RS08315 WP 078228268.1	KDEKIGQGSENAKKYLDNPEIFDEIDHQVVRVQFGLIDGEEIAE...SKKDE...AAQV.DS..VNEEVTLDLGDELE	379
NTDB id 408 SMU RS09530 WP 002262392.1	NGEKIGQGSENAKKFLADNPEIFDDIDHKVVRVQYGLIEEDNTEEKQSSKEK...ETD.EK..ADKNLVLELDDTIE	380
NTDB id 453 Spy49 1753c ACI62003.1	NGEKIGQGSENAKRYLADHPQLFDEIDRKVVRVQFGLLEESEESAMAVASE...E...TDDLALDLNGIE	375
NTDB id 115 BSU 16940 NP 389576.2	EEERLGGRENAKQFLKENKDIMLMIQEIQIREHYGLDNNGVVQQAEETQEEL.EFE.E...	348
NTDB id 625 LCA RS02525 WP 011374196.1	GSERIGQGRENAKNYLADHEDVEDEVRLKVRAAAGTISDVPEEDLPTTEDE.QI.NILPDDS.T...EE...	355
NTDB id 1313 DR RS12030 WP 010888966.1	GDERIGQGGKEKTIAYIAERPEMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPEAAE..A...	363
NTDB id 1245 Cj1673c YP 002345041.1	KDKKLGQGRENSKAFLENPEIADKLEQEIIRDRVMAAIRAGNAGEAPALAPAPAAPEAAE..A...	343
NTDB id 1214 NCTC11637 00359 SQJ03206.1	QDKKLGQGRENAKALLKEDKALADEITLTKESIGSNEEI...MPLPDE..PLE..EME...	347
NTDB id 1211 C694 RS00795 WP 000963128.1	QDKKLGQGRENAKALLKEDKALADEITLTKESIGSNEEI...MPLPDE..PLE..EME...	347
NTDB id 1334 RAOC RS04870 WP 004918236.1	QDTKLGQGRDRTVKELTKDNPELAEELEAKIREKLQAK...	343
NTDB id 1389 A4U84 RS07560 WP 021112712.1	DGTKIGQGGKNNAVKWLKENPEIADKLEQEIIRNTLASNPNS...VLKATASDSND..ESD.VEEFSEEEL...	372
NTDB id 1130 NGFG RS03960 WP 003688695.1	NGAKIGQGGKDNVVRVWLKENPEISDEIDAKIRALNGVEMHI...TEGTQDETDEGE..R...PEE...	348
NTDB id 1124 NGFG RS03960 WP 003688695.1	NGAKIGQGGKDNVVRVWLKENPEISDEIDAKIRALNGVEMHI...TEGTQDETDEGE..R...PEE...	348
NTDB id 1123 OK783 RS03845 WP 003688695.1	NGAKIGQGGKDNVVRVWLKENPEISDEIDAKIRALNGVEMHI...TEGTQDETDEGE..R...PEE...	348
NTDB id 1427 RS RS02760 WP 011000510.1	NGERIGQGRDNCREFLRENAELAREIENKVRHLGVTPMG...AVTLA.EEVEE..D...	352
NTDB id 1165 A1552VC RS01565 WP 000344154.1	NGDKIGQGGKANACKYLNKENPEIAKTLDKKLRREMLLNPNM...QLIAETSSAAD..DVE.FGAVPEEF...	354
NTDB id 1142 GTF74 RS11550 WP 000344154.1	NGDKIGQGGKANACKYLNKENPEIAKTLDKKLRREMLLNPNM...QLIAETSSAAD..DVE.FGAVPEEF...	354
NTDB id 1022 ACIAD RS06390 WP 004925639.1	QGNKIGQGGKNNVIRYLEENPQISTEIEAVIREQLLTKASD...QTAA.HDETEE..EPD.LL..ES...	349
NTDB id 1074 ABD1 RS10195 WP 000344167.1	QGNKIGQGGKNNVIRYFEENNQIAEEIERNIREQLLTTGTN...GAVQ.I.EDEE..EPD.LL..LES...	349
NTDB id 1205 PSJM300 05545 AFN77183.1	KGSKIGQGGKANAAKYLEENPEIGQEIEQQIRAKLLVTSAN...TKATPVAEDLA..DAD.I...	347
consensus	***!!***** ** ***** ***** ** **	

logo

NTDB id	Accession	Sequence	Position
92852	MWL86	QAEPEKEGKR	386
600	KW2	.....	387
269	KZH43	IEIEE.....	388
228	SPD	IEIEE.....	388
194	SPR	IEIEE.....	388
159	SP	IEIEE.....	388
527	SMSK321	IEIEE.....	384
499	SM12261	IEIEE.....	384
408	SMU	LED.....	383
453	Spy49	IED.....	378
115	BSU	.....	348
625	LCA	.....	355
1313	DR	.....	363
1245	Cj1673c	.....	343
1214	NCTC11637	.....	347
1211	C694	.....	347
1334	RAOC	.....	343
1389	A4U84	.....	372
1130	NGFG	.....	348
1124	NGFG	.....	348
1123	OK783	.....	348
1427	RS	.....	352
1165	A1552VC	.....	354
1142	GTF74	.....	354
1022	ACIAD	.....	349
1074	ABD1	.....	349
1205	PSJM300	.....	347

consensus

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