

logo

NTDB id 1193 PAKAF RS26665 WP 003161249.1	MNHYFRQFSMGAVAIAMQAASAQVSMTNIVPMQIAGQGTETIRVMFNGLPPQPQAYQLENPSRLILD	84
NTDB id 1009 ACIAD RS15175 WP 004923730.1	MNHYFRQFSMGAVAIAMQAASAQVSMTNIVPMQIAGQGTETIRVMFNGLPPQPQAYQLENPSRLILD	85
NTDB id 53877 M3Q RS18365 WP 001017033.1	MNHYFRQFSMGAVAIAMQAASAQVSMTNIVPMQIAGQGTETIRVMFNGLPPQPQAYQLENPSRLILD	85
NTDB id 1052 ABD1 RS15850 WP 001017033.1	MNHYFRQFSMGAVAIAMQAASAQVSMTNIVPMQIAGQGTETIRVMFNGLPPQPQAYQLENPSRLILD	85
NTDB id 1039 HON27 RS01460 WP 031993779.1	MNHYFRQFSMGAVAIAMQAASAQVSMTNIVPMQIAGQGTETIRVMFNGLPPQPQAYQLENPSRLILD	85
consensus	!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****	

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NTDB id 1193 PAKAF RS26665 WP 003161249.1	NTRSVTVVEAKDRTRLINLTTLSSYTTRVEGNLFFVGNSPAGASVASTAPVKASIPAPASYAQPIKPKPYVPAGRAIRNIDFQ	169
NTDB id 1009 ACIAD RS15175 WP 004923730.1	NTRSVTVVEAKDRTRLINLTTLSSYTTRVEGNLFFVGNSPAGASVASTAPVKASIPAPASYAQPIKPKPYVPAGRAIRNIDFQ	151
NTDB id 53877 M3Q RS18365 WP 001017033.1	NTRSVTVVEAKDRTRLINLTTLSSYTTRVEGNLFFVGNSPAGASVASTAPVKASIPAPASYAQPIKPKPYVPAGRAIRNIDFQ	150
NTDB id 1052 ABD1 RS15850 WP 001017033.1	NTRSVTVVEAKDRTRLINLTTLSSYTTRVEGNLFFVGNSPAGASVASTAPVKASIPAPASYAQPIKPKPYVPAGRAIRNIDFQ	150
NTDB id 1039 HON27 RS01460 WP 031993779.1	NTRSVTVVEAKDRTRLINLTTLSSYTTRVEGNLFFVGNSPAGASVASTAPVKASIPAPASYAQPIKPKPYVPAGRAIRNIDFQ	150
consensus	***!***!*****!*!*****!*****!*****!*****!*****!*****!*****!*****!*****	

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NTDB id 1193 PAKAF RS26665 WP 003161249.1	RGSQGEGLVVDLLGSNTPVDVQQQGSKVIRTIGTKIPHTLARRLNVNDFATPVSSIDAYNDKGVGVITIQSSGSYEYMAVQAE	254
NTDB id 1009 ACIAD RS15175 WP 004923730.1	RGSQGEGLVVDLLGSNTPVDVQQQGSKVIRTIGTKIPHTLARRLNVNDFATPVSSIDAYNDKGVGVITIQSSGSYEYMAVQAE	236
NTDB id 53877 M3Q RS18365 WP 001017033.1	RGSQGEGLVVDLLGSNTPVDVQQQGSKVIRTIGTKIPHTLARRLNVNDFATPVSSIDAYNDKGVGVITIQSSGSYEYMAVQAE	235
NTDB id 1052 ABD1 RS15850 WP 001017033.1	RGSQGEGLVVDLLGSNTPVDVQQQGSKVIRTIGTKIPHTLARRLNVNDFATPVSSIDAYNDKGVGVITIQSSGSYEYMAVQAE	235
NTDB id 1039 HON27 RS01460 WP 031993779.1	RGSQGEGLVVDLLGSNTPVDVQQQGSKVIRTIGTKIPHTLARRLNVNDFATPVSSIDAYNDKGVGVITIQSSGSYEYMAVQAE	235
consensus	!!*!!!!*!!!!*!!!!*****!*!*****!*****!*****!*****!*****!*****!*****!*****!*****	

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NTDB id 1193 PAKAF RS26665 WP 003161249.1	NKLTISLKRPPQDNVTSLYKTPNYSGNKLSLDFQDIEVRRVLQLLADFTGINMVAADSVQGNITLRLKDVDPWDQALDIILKTKN	339
NTDB id 1009 ACIAD RS15175 WP 004923730.1	NKLTISLKRPPQDNVTSLYKTPNYSGNKLSLDFQDIEVRRVLQLLADFTGINMVAADSVQGNITLRLKDVDPWDQALDIILKTKN	320
NTDB id 53877 M3Q RS18365 WP 001017033.1	NKLTISLKRPPQDNVTSLYKTPNYSGNKLSLDFQDIEVRRVLQLLADFTGINMVAADSVQGNITLRLKDVDPWDQALDIILKTKN	319
NTDB id 1052 ABD1 RS15850 WP 001017033.1	NKLTISLKRPPQDNVTSLYKTPNYSGNKLSLDFQDIEVRRVLQLLADFTGINMVAADSVQGNITLRLKDVDPWDQALDIILKTKN	319
NTDB id 1039 HON27 RS01460 WP 031993779.1	NKLTISLKRPPQDNVTSLYKTPNYSGNKLSLDFQDIEVRRVLQLLADFTGINMVAADSVQGNITLRLKDVDPWDQALDIILKTKN	319
consensus	!!*!!!!*!!!!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****	

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NTDB id 1193 PAKAF RS26665 WP 003161249.1	LDKRRNGNVIWIAPVAELIKAEFEAKAVAQSVKLAPLQTEYIQLKYAKAGDIMGLITQGSNNSNGLQHTSGGGTSTSTNLNTGV	407
NTDB id 1009 ACIAD RS15175 WP 004923730.1	LDKRRNGNVIWIAPVAELIKAEFEAKAVAQSVKLAPLQTEYIQLKYAKAGDIMGLITQGSNNSNGLQHTSGGGTSTSTNLNTGV	396
NTDB id 53877 M3Q RS18365 WP 001017033.1	LDKRRNGNVIWIAPVAELIKAEFEAKAVAQSVKLAPLQTEYIQLKYAKAGDIMGLITQGSNNSNGLQHTSGGGTSTSTNLNTGV	404
NTDB id 1052 ABD1 RS15850 WP 001017033.1	LDKRRNGNVIWIAPVAELIKAEFEAKAVAQSVKLAPLQTEYIQLKYAKAGDIMGLITQGSNNSNGLQHTSGGGTSTSTNLNTGV	404
NTDB id 1039 HON27 RS01460 WP 031993779.1	LDKRRNGNVIWIAPVAELIKAEFEAKAVAQSVKLAPLQTEYIQLKYAKAGDIMGLITQGSNNSNGLQHTSGGGTSTSTNLNTGV	404
consensus	!!!!*!!!!*!!!!*****!*!*****!*****!*****!*****!*****!*****!*****!*****!*****	

