

logo

NTDB id 601	KW2 RS02615 WP 021036805.1	LVDLGYNPAMGARPLRRTIQENTIEDSTADFYIEHPEYKQ.LVADLIDDKIVISNQTQETAETDDEEVP	AE
NTDB id 378	SMU RS02690 WP 002263569.1	LVDLGYDPKMGARPLRRTIQEHIEDAITDFYLEHPEQKE.LKAIMTSNGKIVIKASNK.VETVTKATSD.	753
NTDB id 241	SP RS04015 WP 000882523.1	LVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKD.LKAVMTSKGNIQIKSAKK.AEVKSSEKEK.	752
NTDB id 281	KZH43 RS03550 WP 000882517.1	LVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKD.LKAVMTSKGNIQIKSAKK.AEVKSSEKEK.	752
NTDB id 245	SPD RS03860 WP 000882517.1	LVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKD.LKAVMTSKGNIQIKSAKK.AEVKSSEKEK.	752
NTDB id 243	SPR RS03635 WP 000882517.1	LVDLGYDPKMGARPLRRTIQDYIEDTITDYYLENPSEKD.LKAVMTSKGNIQIKSAKK.AEVKSSEKEK.	752
NTDB id 1011887	ABUH86 RS00580 WP 007410388.1	VAEEGVLDLEYGARPLRRRAIQKHVEDRLSEELLRGNIDKGGHIVLQDVEDGEFVVKTTAKTN.....	810
NTDB id 85	BSU 00860 NP 387967.1	VAEEGVLDLEYGARPLRRRAIQKHVEDRLSEELLRGNIHKGQHIVLQDVEDGEFVVKTTAKTN.....	810
NTDB id 610	V4T04 RS10165 WP 012897346.1	IAEVGFDPPEYGARPLRKAALQKEVEDLLSEQLLSGETKAGNHVSIIGASNKKTKIAQIV.....	816
NTDB id 290	KZH43 RS10025 WP 001109677.1	LANQGYDPEMGARPLRRTLQTEVEDKLAELLLKGDVLVAGSILKIGVKAGQLKFDIA.....	810
NTDB id 289	SPD RS10700 WP 001109677.1	LANQGYDPEMGARPLRRTLQTEVEDKLAELLLKGDVLVAGSILKIGVKAGQLKFDIA.....	810
NTDB id 287	SP RS11210 WP 001109712.1	LANQGYDPEMGARPLRRTLQTEVEDKLAELLLKGDVLVAGSTLKIIGVKAGQLKFDIA.....	810
NTDB id 377	SMU RS09275 WP 002262344.1	LALAGYDAEMGARPLRRTLQTEVEDKLAELILSGKLASGQTLKIIGIQKEKIKFDIV.....	813
NTDB id 297	STER RS00545 WP 011680614.1	LAKDGYDIEMGARPLRRTIQTVEDKLSLELLGGQVVSQTLKIIGCSKDKLTFTVV.....	816
NTDB id 329	STU RS10020 WP 011225298.1	LAKDGYDIEMGARPLRRTIQTVEDKLSLELLGGQVVSQTLKIIGCSKDKLTFTVV.....	816
consensus		*** !*****!!!!!!***! *!! *** ** * * * * * * * * * * * * * *	

- non conserved
- similar
- ≥ 50% conserved