pMLST-2.0 Server - Results

pMLST profile: *IncF RST*

Sequence Type: *[S1:A-:B17]*

Locus	Identity	Coverage	Alignment Length	Allele Length	Gaps	Allele
FIA						No hit found
FIB	100.0	100.0	373	373	0	FIB_17
FIC						No hit found
FII						No hit found
FIIK						No hit found
FIIS	100.0	100.0	215	215	0	FIIS_1
FIIY						No hit found

[extended output]

Input Files: *unknown_sample.fasta.txt*

Please download your results using the buttons below.

[Results as text] [Results tsv] [Hit in genome sequences] [pMLST allele sequences]