100%			Species	Avg. GC (%)	Gram Stain	gDNA Abun. (%)
90%	_		Pseudomonas aeruinosa	66.2	-	12
80%			Escherichia coli	56.8	-	12
70%			Salmonella enterica	52.2	-	12
60%			Lactobacillus fermentum	52.8	+	12
50%			■ Enterococcus faecalis	37.5	+	12
40%			Syaphylococcus aureus	32.7	+	12
30%			■ Listeria monocytogenes	38.0	+	12
20%			Bacillus subtilis	43.8	+	12
10%			Saccharomyces cerevisiae	38.4	Yeast	2
0%	Theoretical	DNA Standard	Cryptococcus neoformans	48.2	Yeast	2