#Sequences	2,000		4,114		6,718		15,011	
Method	Sec	Log-Lk	Sec	Log Lk	Sec	Log Lk	Sec	Log Lk
FastTree, no ML	84	-368,567	210	-330,774	348	-499,732	1,082	-3,981,266
FastTree with ML	265	-366,517	585	-328,062	955	-493,841	2,441	-3,961,198
no $ML + 1$ round $SPRs$	576	-366,128	1,154	-327,193	2,085	-492,522	4,499	-3,955,028
no ML $+ 2$ rounds SPRs	973	-365,766	2,003	-326,786	3,253	-489,991	6,245	-3,948,824
RAxML Parsimony	286	-366,605	740	-328,069	1,264	-484,909	7,764	-3,951,842
+1 round SPRs	943	-365,745	1,832	-326,854	3,559	-482,927	12,467	-3,940,585
+2 rounds SPRs	1,307	-365,617	2,576	-326,584	4,685	-482,530	14,126	-3,937,613
RAxML, Final	12,618	-364,912	23,637	-325,581	44,905	-481,259	106,855	-3,924,702

This is the same analysis as in Figure 2, but in tabular form, and with data for additional 16S rRNA alignments.