UPGMA Tree

		Salmonella virus Fels2 (AE006468) #4
0.0184		Salmonella virus Fels2 (AE006468) #6
0.0131		Salmonella virus Fels2 (AE006468) #5
	0.0011	Salmonella virus Fels2 (AE006468) #1

- 0.000713 — 0.000492 — 0.000328 Salmonella virus Fels2 (AE006468) #3
- 0.000328 Salmonella virus Fels2 (AE006468) #7
- 0.00277 — 0.000822 Salmonella virus Fels2 (AE006468) #2

0.065

- 0.00463 Escherichia coli str. K-12 substr. MG1655 (U00096) #1
- 0.0103 — 0.00328 Escherichia coli str. K-12 substr. MG1655 (U00096) #6
- 0.0023 — 0.00164 Escherichia coli str. K-12 substr. MG1655 (U00096) #7
- 0.00328 Escherichia coli str. K-12 substr. MG1655 (U00096) #5
- 0.00195
- 0.000495 — 0.00164 Escherichia coli str. K-12 substr. MG1655 (U00096) #4
- 0.00496
- 0.0214 Escherichia coli str. K-12 substr. MG1655 (U00096) #3
- 0.0214 Escherichia coli str. K-12 substr. MG1655 (U00096) #2

0.0519

- 0.00033 — 0 Pseudomonas aeruginosa PAO1 (AE004091) #3
- 0 Pseudomonas aeruginosa PAO1 (AE004091) #4
- 0.0796
- 0.00033 — 0 Pseudomonas aeruginosa PAO1 (AE004091) #2
- 0 Pseudomonas aeruginosa PAO1 (AE004091) #1

- 0.000326 Staphylococcus aureus subsp. aureus NCTC 8325 (CP000253) #4
- 0.000436 — 0 Staphylococcus aureus subsp. aureus NCTC 8325 (CP000253) #2
- 0.000326 — 0 Staphylococcus aureus subsp. aureus NCTC 8325 (CP000253) #5
- 0.000463
- 0.000762 Staphylococcus aureus subsp. aureus NCTC 8325 (CP000253) #3
- 0.0409
- 0.0122 Staphylococcus aureus subsp. aureus NCTC 8325 (CP000253) #1
- 0.0896

0.0421 Bacillus subtilis (AJ276351)

0.416667