**Table S4. Sequence variation in *rpoB*, *DNA topoisomerase I, and DNA polymerase III subunits gamma* genes of *L. pneumophila* strains.**

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Gene | No. of analyzed sequences | Sequence length | \*No. of variablesites | % Sequencevariation |
| *rpoB* | 62 | 4107 | 216 | 5.26 |
| *DNA topoisomerase I* | 51 | 2280 | 79 | 3.46 |
| *DNA polymerase* *III subunits gamma* | 62 | 1671 | 135 | 8.08 |

\* Number of variable sites of the gene were obtained by analyzing known sequences in NCBI database, using Dnasp v5 (http://www.ub.edu/dnasp/).