Supplementary Table 6. Nucleotide differences within and between the groups of organisms used for the whole genome SNP analysis.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Organism group | Between group differences | | | Within group differences |
| Number of nucleotides | | | |
| Non-pneumococcal *tts* positive |  |  |  | 5327.3 |
| *Streptococcus pneumoniae* | 6359.3 |  |  | 1545.2 |
| *Streptococcus mitis* | 7661.514 | 6853.7 |  | 7811.4 |
| *Streptococcus pseudopneumoniae* | 7123.5 | 5857.8 | 7870.1 | 3555.7 |