Supplementary Table S1. *Non-CMRSA2/7/10 clones by PFGE and* spa *type*

|  |  |  |
| --- | --- | --- |
| PFGE type | *spa* type | Frequency |
| CMRSA 1/USA600 (n=2) | t065 | 2 |
| CMRSA 3/6  (n=11) | t037 | 11 |
| CMRSA 4/USA200 (n=2) | t012  t840 | 1  1 |
| CMRSA 5/USA500 (n=1) | t064 | 1 |
| CMRSA 8  (n=31) | t022  t032  t11427  t12130  t3490  t5983  t6443  t7982 | 10  2  1  1  1  1  14  1 |
| CMRSA not assigned\*  (n=15) | t019 (USA1100, SWP/Oceania)  t044  t1346 (USA700)  t148 (USA700)  t1752 (USA1100, SWP/Oceania)  t1839  t216 (USA1000, China/Taiwan)  t314  t437 (USA1000, China/Taiwan)  t4549  t657  t7747 | 1  1  1  1  1  1  3  2  1  1  1  1 |
| Total |  | 62 |

\* Isolates were designated as ‘not assigned’ if their PFGE patterns did not cluster with the profiles of the ten Canadian epidemic clones [1].

**REFERENCE**

1. **Li V, *et al*.** Cost-effectiveness and efficacy of *spa*, SCC*mec*, and PVL genotyping of methicillin-resistant *Staphylococcus aureus* as compared to pulsed-field gel electrophoresis. *PLoS ONE.* Published online: 14 November2013. doi:10.1371/journal.pone.0079149.