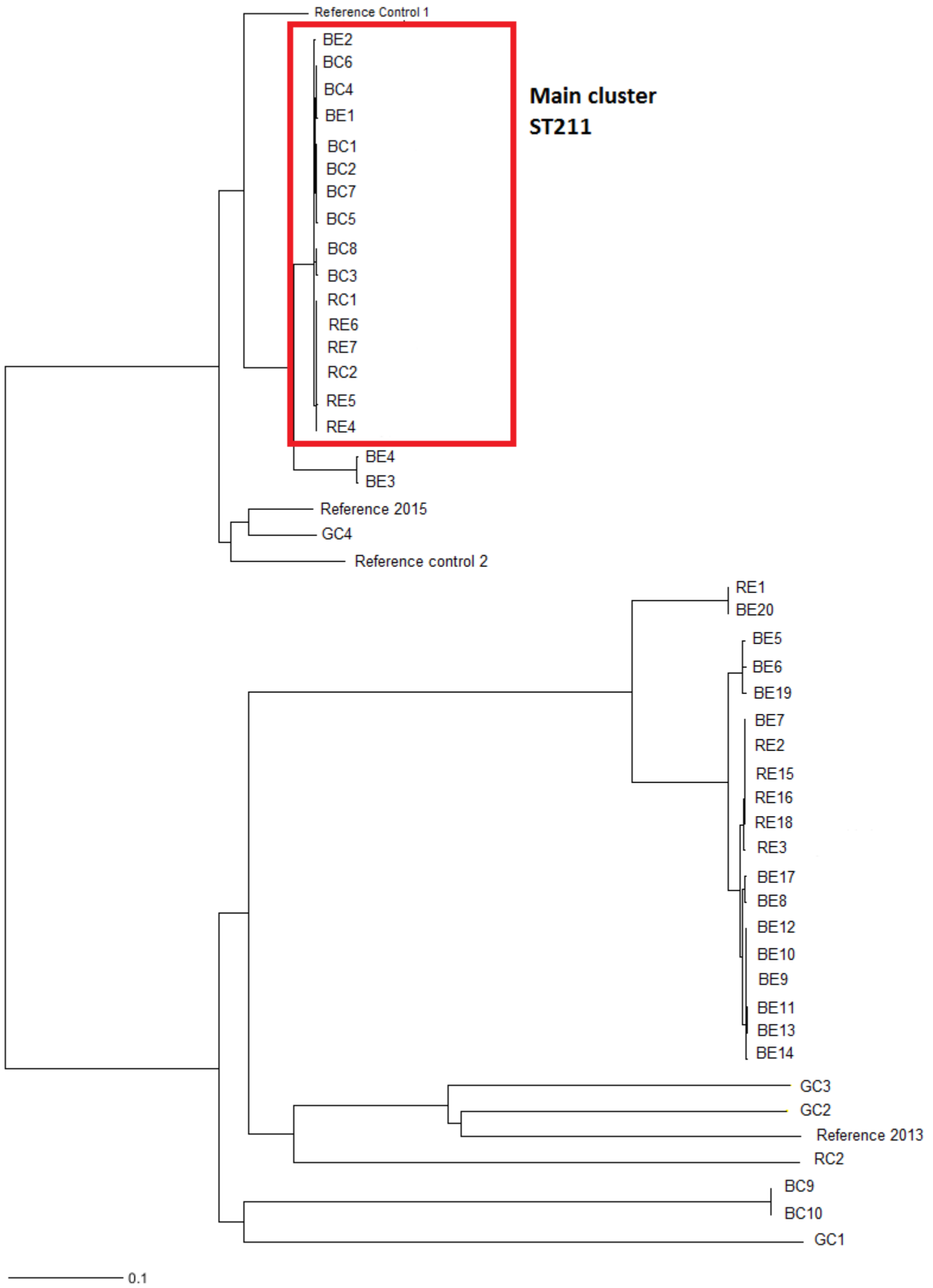


Figure 4: cgMLST phylogeny of all outbreak isolates between February-May 2016 in Sydney.



NJ tree for 48 Samples based on 1530 columns, pairwise ignoring missing values, % columns difference
Distance based on columns from *L. pneumophila* cgMLST 1530 targets Philadelphia 1 (1530)