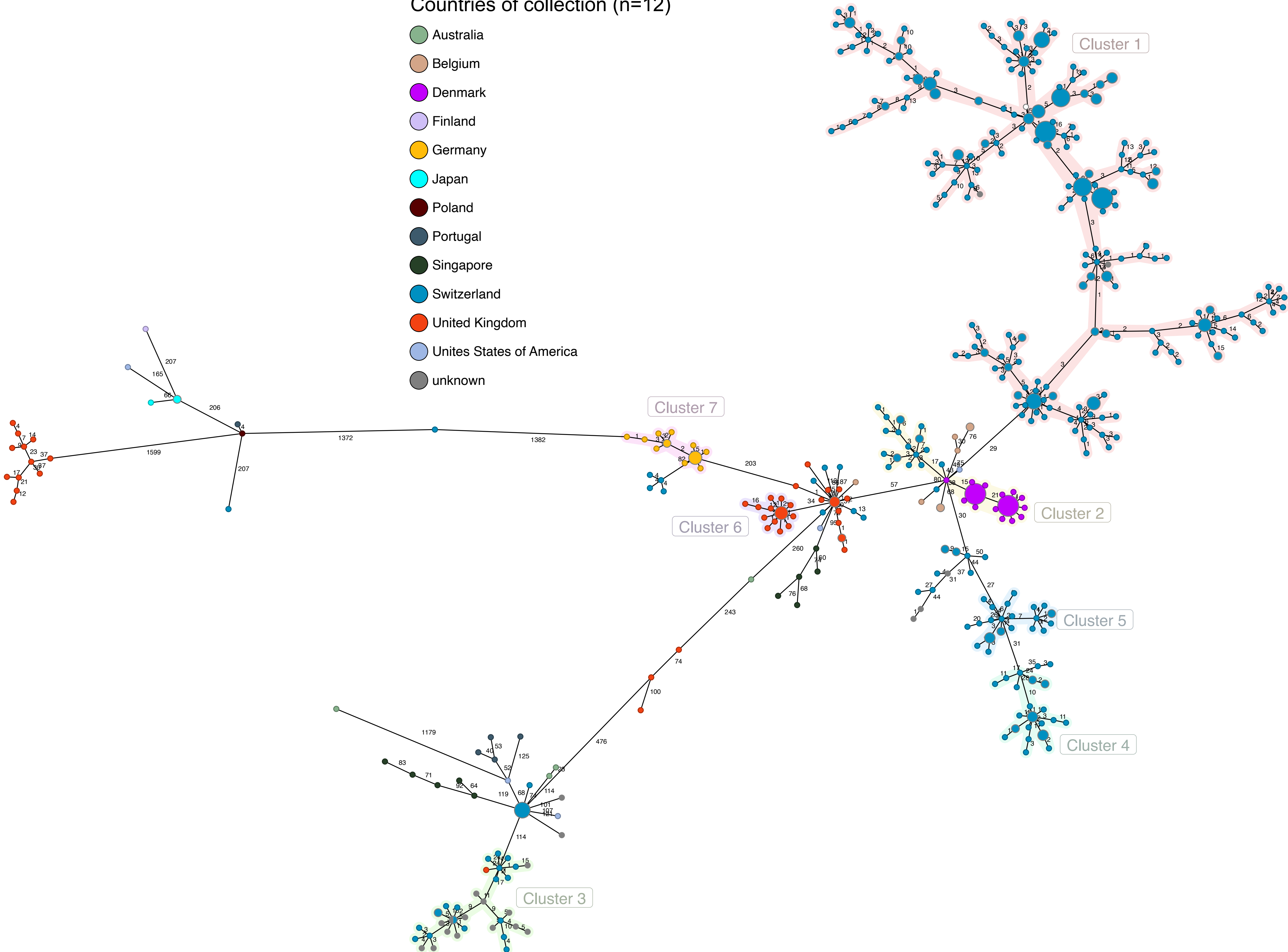


# Countries of collection (n=12)

- Australia
- Belgium
- Denmark
- Finland
- Germany
- Japan
- Poland
- Portugal
- Singapore
- Switzerland
- United Kingdom
- Unites States of America
- unknown



B. cgMLST MST of CCoS and public repository genomes of MRSA (1960-1992, n = 554), nodes colored by collection country. Clusters are shown with a maximal cluster distance of 24 allele differences and a minimal cluster size of 15.