

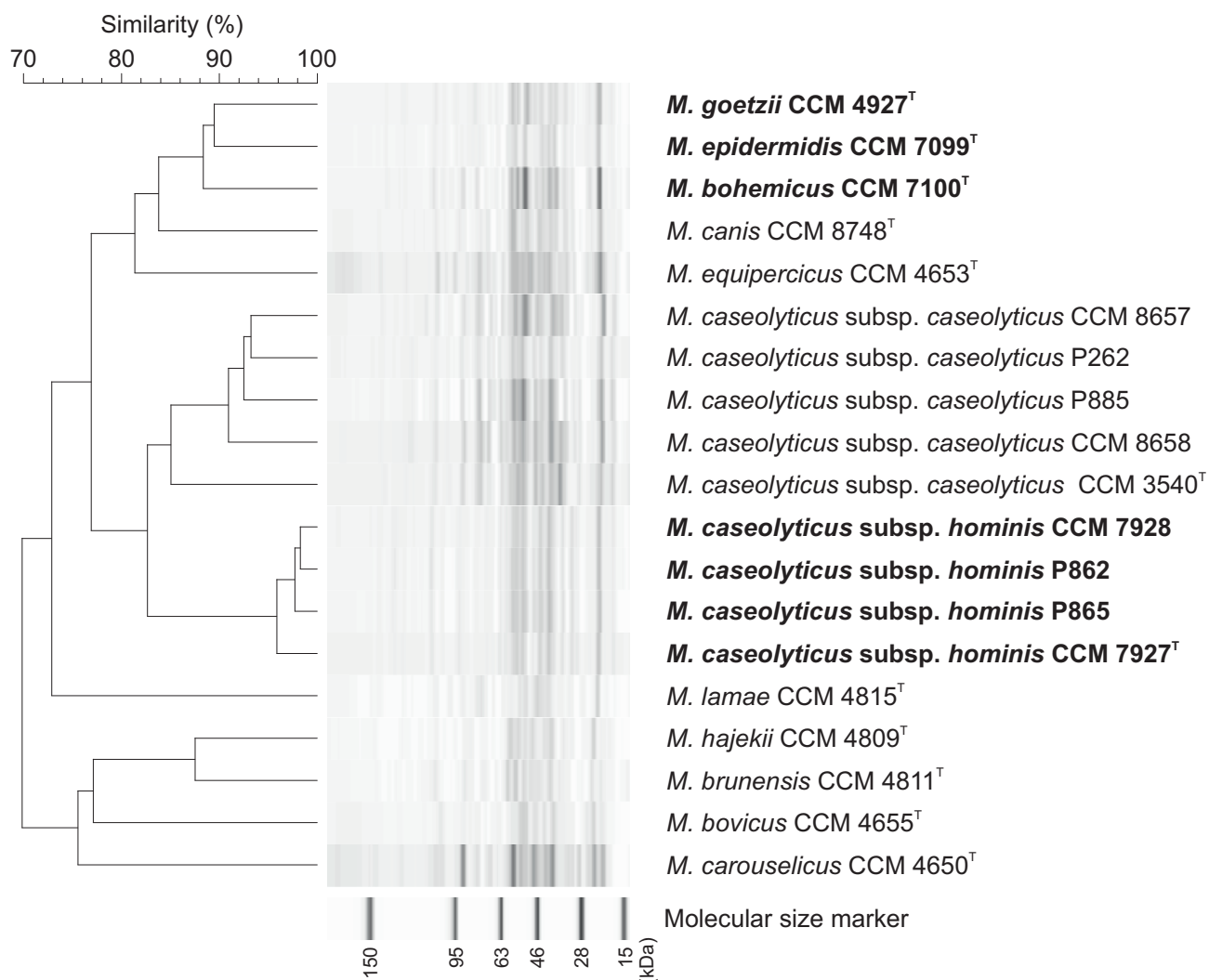


FIGURE S3. Dendrogram based on cluster analysis of whole-cell protein profiles obtained using Agilent 2100 Bioanalyzer system with Protein 230 kit (Agilent Technologies) from investigated strains and representative strains of phylogenetically related *Micrococcus* species. The dendrogram was calculated with Pearson's correlation coefficients by the UPGMA clustering method (r , expressed as percentage similarity values).