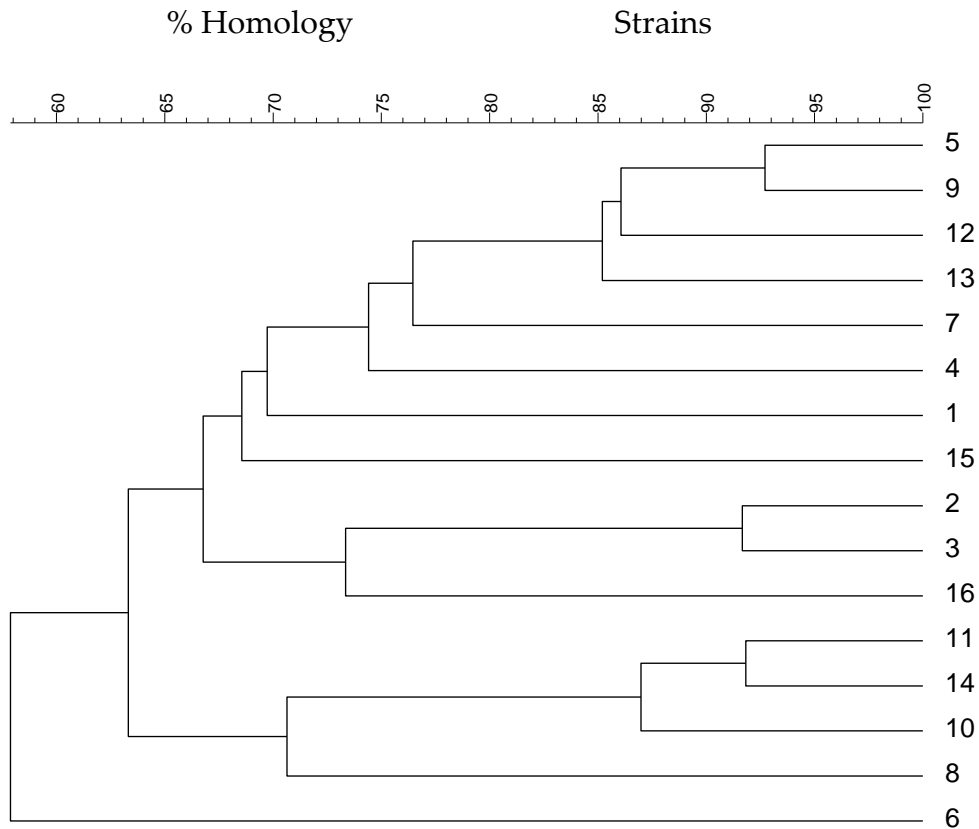


# Supplementary Materials: *Staphylococcus aureus* Isolates from Bovine Mastitis in Eight Countries: Genotypes, Detection of Genes Encoding Different Toxins and Other Virulence Genes

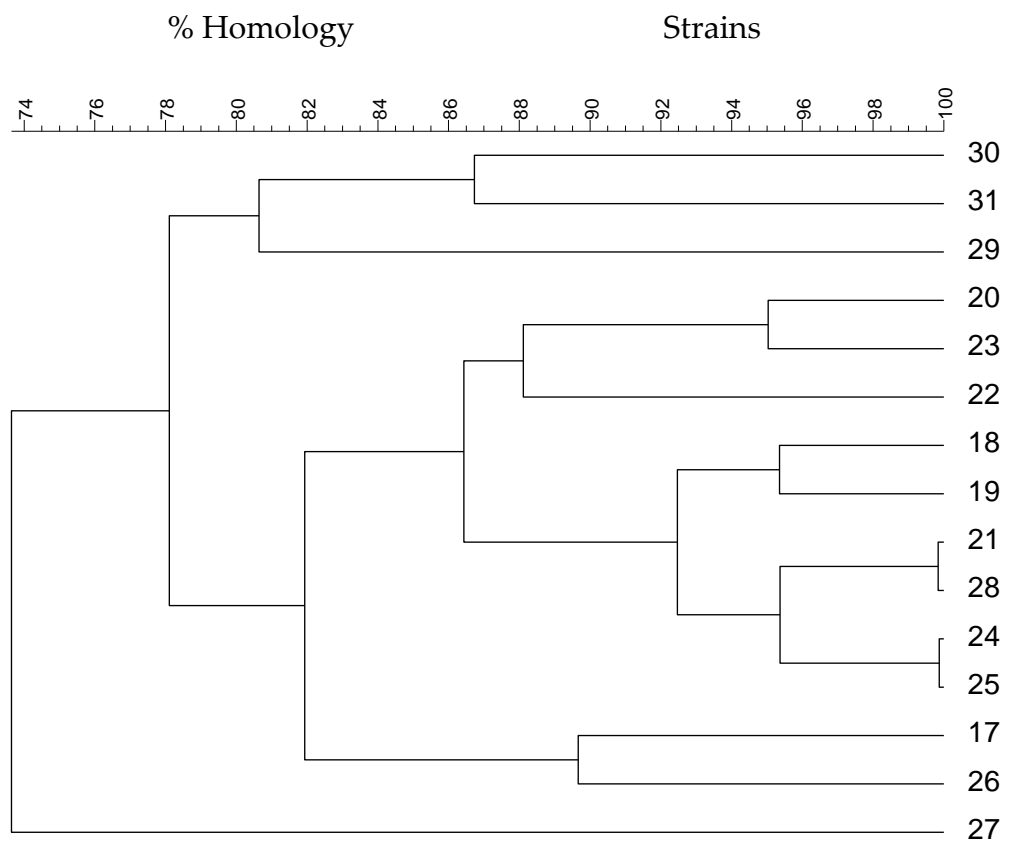
Valentina Monistero, Hans Ulrich Graber, Claudia Pollera, Paola Cremonesi, Bianca Castiglioni, Enriqueta Bottini, Alejandro Ceballos-Marquez, Laura Lasso-Rojas, Volker Kroemker, Nicole Wentz, Inge-Marie Petzer, Carlos Santisteban, Jeff Runyan, Marcos Veiga dos Santos, Bruna Gomes Alves, Renata Piccinini, Valerio Bronzo, Mohamed Salah Abbassi, Meriam Ben Said and Paolo Moroni

Unweighted pair-group method with arithmetic averages (UPGMA)-based dendrogram derived from the combined RS-PCR profiles and the virulence factors of the *S. aureus* strains considered in this study.

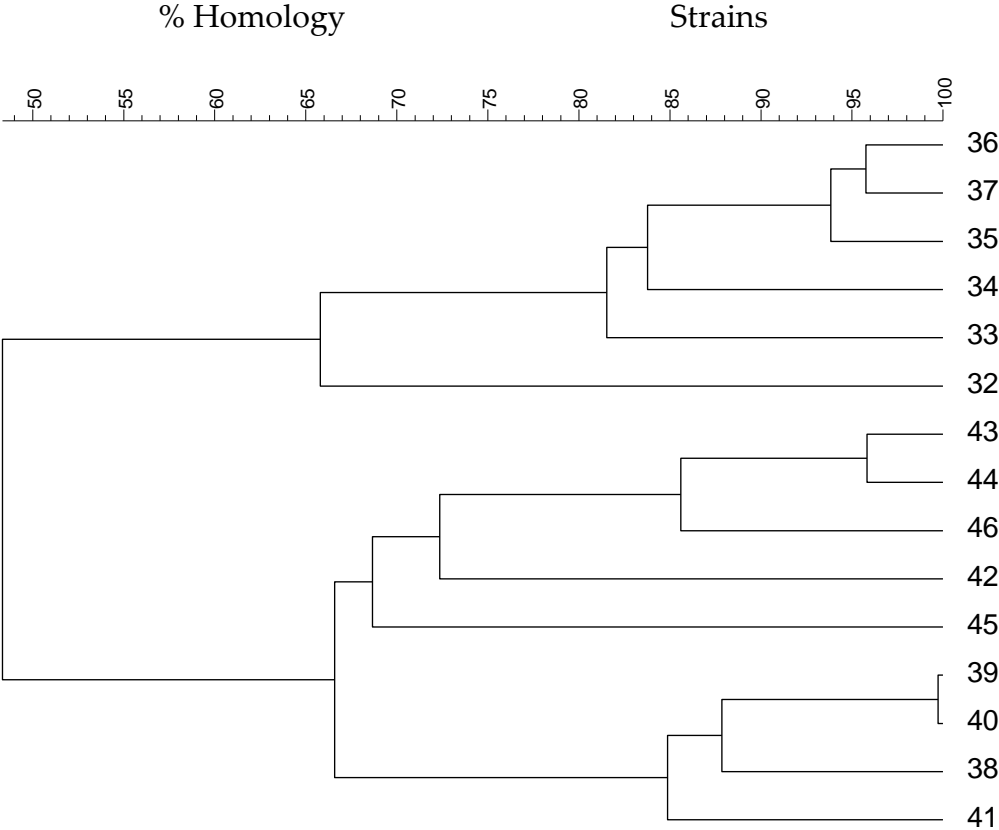
## Argentina



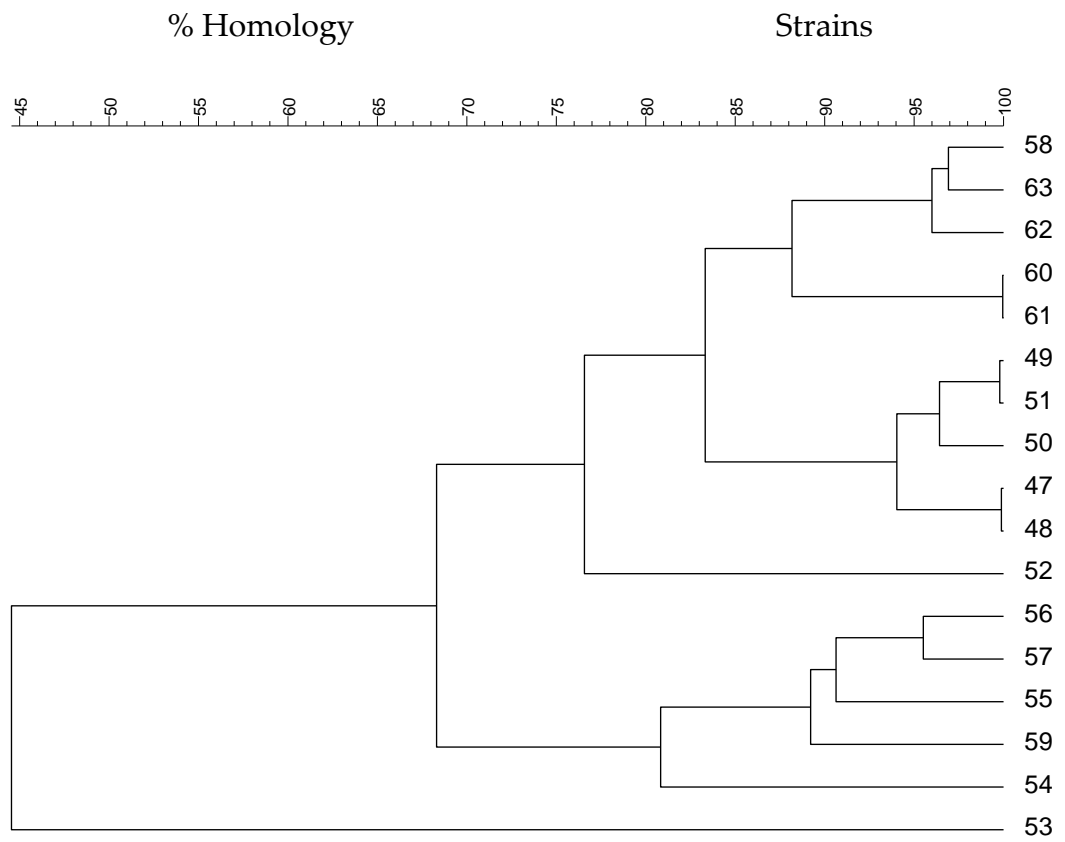
**Brazil**



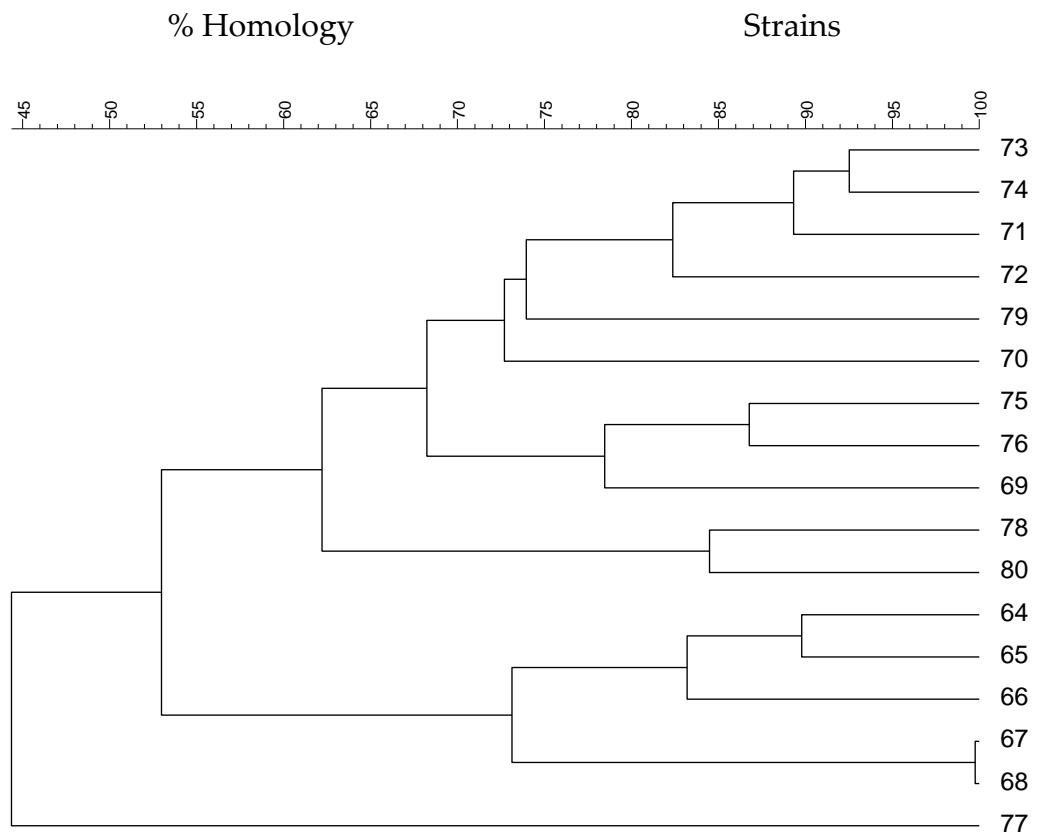
Colombia



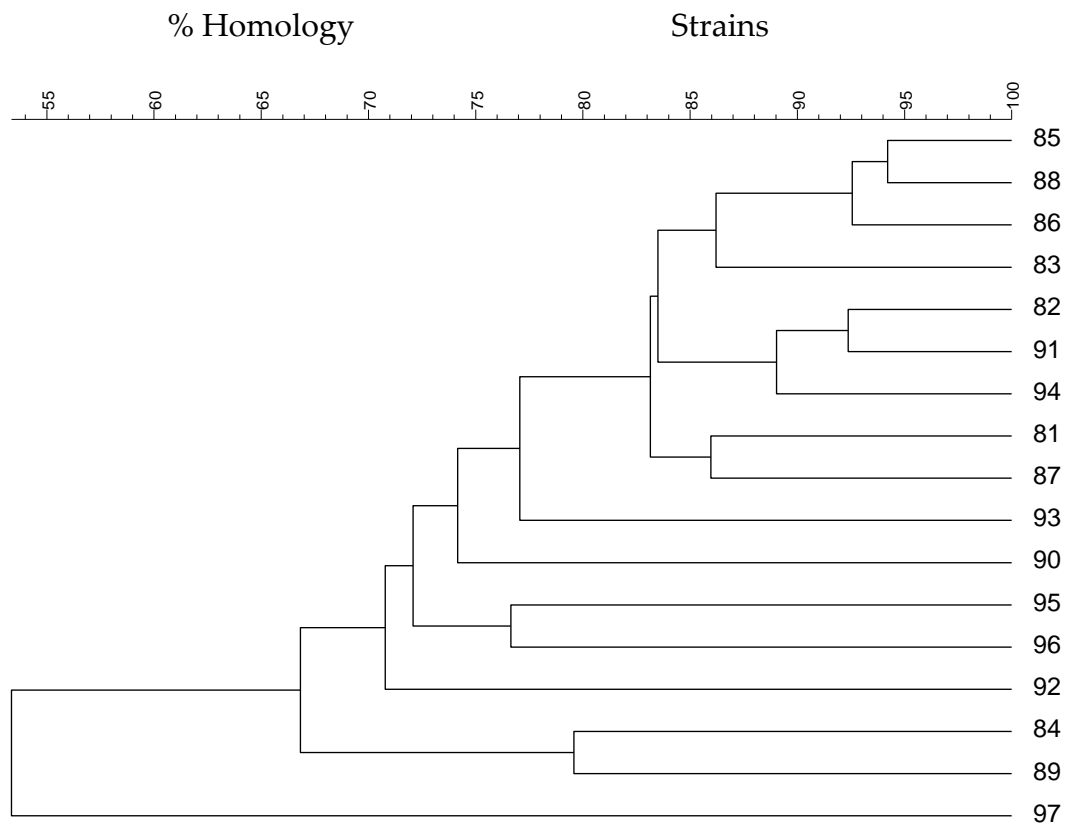
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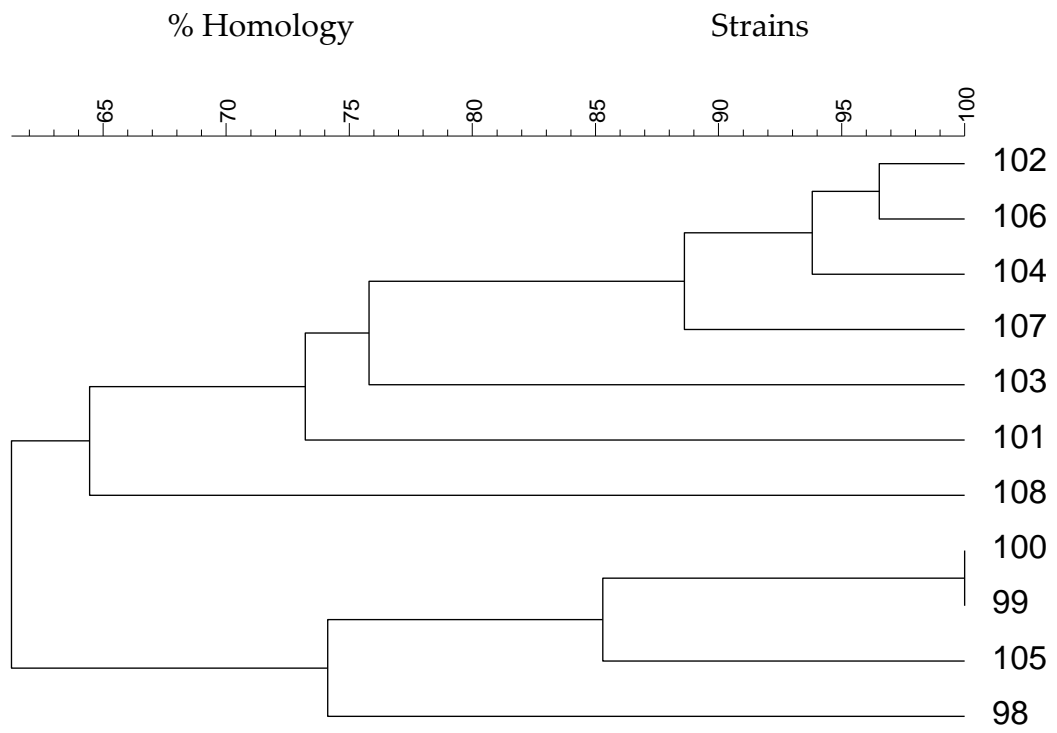
Italy



**New York State**



**South Africa**



Tunisia

