



Figure S1. Phylogenetic tree of all 100 *L. monocytogenes* isolated tested based on the cgMLST analyses and prevalence of the selected genes (green – gene presence, red – gene absence). Column designations: 1. Isolate ID number, 2. Isolate CCs, 3. Isolate CTs, 4. Isolate source (voivodeship), 5. Year of isolation, 6. Isolate origin, 7. *bcrABC* genes, 8. SSI-1 genes, 9. SSI-2 genes, 10. *ermC* (*Tn6118_qac*) gene.