

Supplementary Material 2. cgMLST analysis; Minimum spanning tree (created within Ridom Seqsphere+; setting: pairwise ignore missing values) of carbapenem-resistant *Acinetobacter baumannii* isolates (n=74). Clustering distance: 9. Isolates that are assigned to the same circle have no allelic differences. cgMLST clusters are highlighted in color. The numbers next to the lines indicate the number of allelic differences (the lengths of the lines are not proportional to the allelic differences for reasons of space).

