



Supplementary Figure S1: Neighbor Joining Tree of 28 *Acinetobacter baumannii* isolates based on 2.390 genes included in the cgMLST scheme of Ridom SeqSphere+. The tree is based on a comparison of publicly available *A. baumannii* genomes representing international clones IC1 (AYE, GenBank No. NC_010410.1; AB0057, NC011586.2), IC2 (1656-2, NC_017162.1; MDR-TJ, NC_017847.1), IC3 (NIPH 1669, APOQ01.1; Naval-81, AFDB02.1), IC4 (NIPH 1734, APOX01.1, Ab825, NTFR01.1), IC5 (AB030, NZ_CP009257.1; AF-401, NZ_CP018254.1), IC6 (Ab-3909, AEOZ01.1; AB4332, RJLV01.1), IC7 (161/07, JZCA01.1; LUH_6220, JZBW01.1), IC8 (XH693, LYIY01.1; LAC-4, NZ_CP007712) and IC9 (MRSN15574, VHGP01.1; AC-20, JABESZ01.1), and CP positive *A. baumannii* isolates from this study that could not be assigned to a known IC by PCR.