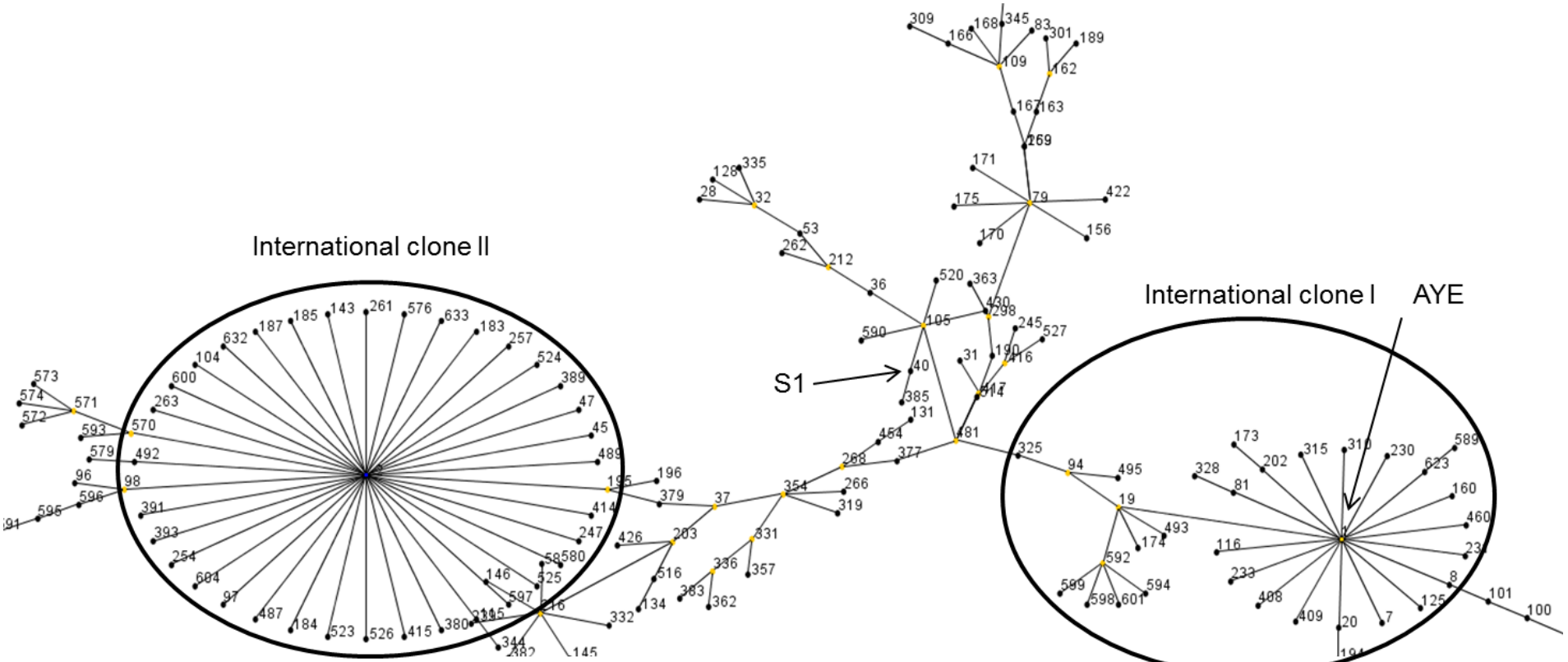Fig S1 Relationships between 615 *A. baumannii* isolates based on MLST data (Pasteur scheme) as calculated by the BURST algorithm. Circles indicate major international groups responsible for the majority of disease. Arrows indicate strains used in this study. ATCC 19606 represents a singleton ST (ST 52) and is not on this network.