



Additional File 5. Relationship dendrogram of the 38 strains based on the genomic fluidity analysis proposed by Kislyuk *et al.* [55]. *A. baumannii* SDF is again outside the ACB complex, the two *A. nosocomialis* strains fall within the *A. baumannii* cluster and PHEA-2 is closest to DR1 and the two *A. calcoaceticus* strains. These differences may be explained by the fact that genomic fluidity is strongly affected by putative horizontal gene transfer events.