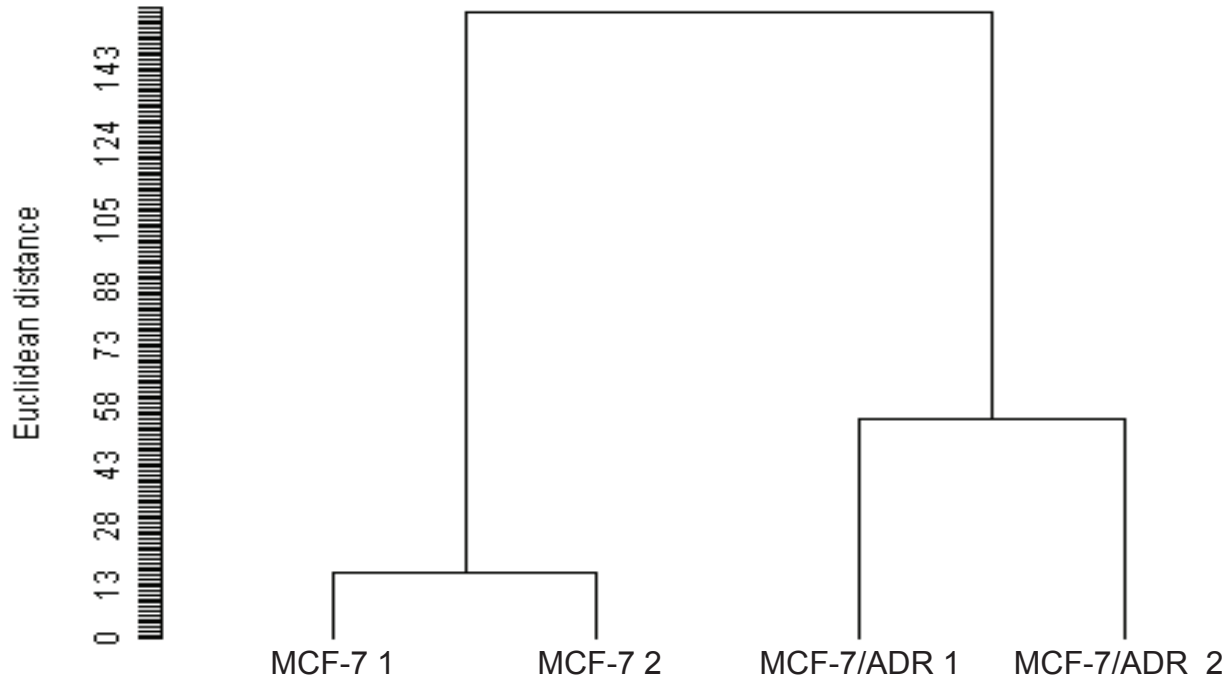


Calcagno et al.
Supplementary Figure 1



Supplementary Figure 1. Dendrogram from clustering experiments. Hierarchical clustering analysis used the Euclidean distance and a complete linkage algorithm. Hierarchical clustering of the 4976 genes remaining after filtering by exclusion parameters was used to determine the similarities of replicate gene expression profiles from MCF-7 and MCF-7/ADR cells. MCF-7 1= replicate 1 of MCF-7; MCF-7 2= replicate 2 of MCF-7; MCF-7/ADR 1= replicate 1 of MCF-7/ADR; MCF-7/ADR 2= replicate 2 of MCF-7/ADR.