Supplemental Figure S1

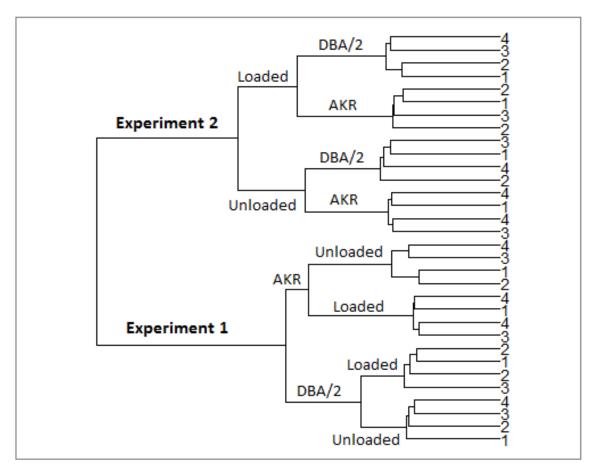


Figure S1: Hierarchical clustering analysis of 32 samples included in the study. Four replicates each for control and cholesterol loaded macrophages from AKR and DBA/2 strain, both independent experiments were included (loaded; unloaded samples).