



Supplementary Figure 1. Plots of principal component 1 (PC1) against principal component 2 (PC2) based on normalized read counts from each subject of normal control group (C1-5 from left to right, top row), CGD patients (D1-4 from left to right, middle row) and of IFNGRD patients (N1-6 from left to right, bottom row) analyzed as different RNA seq datasets. There is distinct separation of IFN- γ treated versus untreated samples of all normal control and CGD patients however the majority of IFNGRD patients do not form distinct groups between IFN- γ treated and untreated samples.