S5 Fig. Principal Component Analyses (PCA) of Gc and PMN transcriptomes including 0-hour samples. A) PCA of Gc transcriptomes based on 1582 core genes. PC1 separates the 130 and H041 strains, and PC2 separates the different infection conditions and timepoints. B) PCA of PMN transcriptomes. PC1 separates the two timepoints (0h vs 1h), and PC2 separates the different PMN donors. C) 3-dimensional PCA based on the top 1% of genes contributing to PMN principal components (PC2, PC3 & PC5, genes listed in S2 Dataset).