Supplementary material to article by Y. Tan et al. "Nanostring Analysis of Skin Biopsies from Patients with Henoch-Schönlein Purpura Reveals Genes Associated with Pathology and Heterogeneity in the Disease Process"

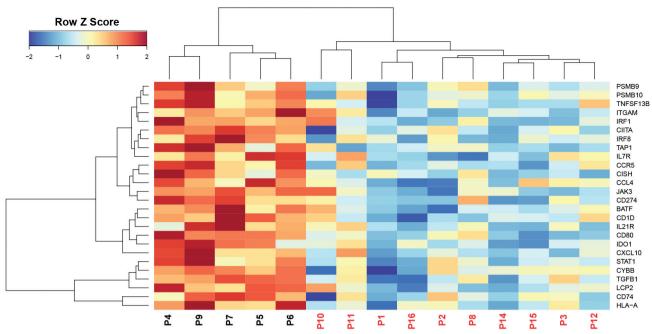


Fig. S2. Dendrogram of genes identified by Ingenuity Pathway Analysis in the 2 HSP patient clusters. HSP-normal samples are labelled in black and HSP-inflamed samples are labelled in red. The log10 normalized counts were transformed into row z-scores for each gene and depicted in a colour scale with blue as the minimum value and red as the maximum value. The samples and genes are clustered using hierarchical clustering of the Euclidean distance with average linkage.