

Supplementary Figure 1: Heatmaps illustrating expression of genes distinguishing control and SpA peripheral blood in Set 1 and Set 2. All genes met the following criteria: >1.5 fold change difference in mean gene expression levels between SpA and control subjects and a significance analysis of microarrays (SAM) q value of < 5%. Each column represents gene expression in one subject and each row represents the signal from one probe set. Yellow indicates genes expressed at higher levels in SpA subjects; blue describes lower levels. The numeric data are provided in Supplementary Table 1.

