

Subjects

Pathway Name	p-value
Phagosome Formation	1.41 x 10 ⁻⁸
iCOS-iCOSL Signaling in T Helper Cells	3.01 x 10 ⁻⁸
CD28 Signaling in T Helper Cells	5.98 x 10 ⁻⁸
Role of Macrophages, Fibroblasts and Endothelial Cell in RA	6.31 x 10 ⁻⁸
Role of NFAT in Regulation of Immune Response	1.40 x 10 ⁻⁷

Fig. S4. Heatmap and pathway analysis of genes (rows) differentially expressed at the second post-surgery time point compared to baseline in the placebo group. Columns represent individual subjects. The table shows pathway enrichment (RA: Rheumatoid Arthritis)