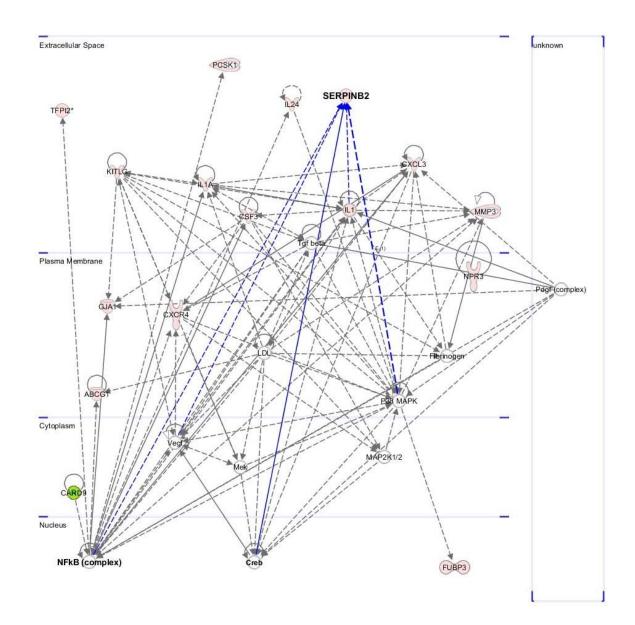
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*Cysteinyl leukotriene signaling through perinuclear CysLT*₁ *receptors on vascular smooth muscle cells transduces nuclear calcium signaling and alterations of gene expression*



Supplementary Fig 1. Functional gene network associated with upregulation by LTC_4 stimulation (derived from Ingenuity Pathway analysis). Network shown, suggested to be implicated in cellular movement and hematopoietic system function, was found to be markedly upregulated (Fishers' exact test score 1×10^{-35}) by LTC_4 stimulation based on analysis of significantly differentially expressed genes (Table 1). Red shading indicates significantly upregulated genes. Intensity of the shading is proportional to the fold change. Interactions involving PAI-2 are highlighted in blue.