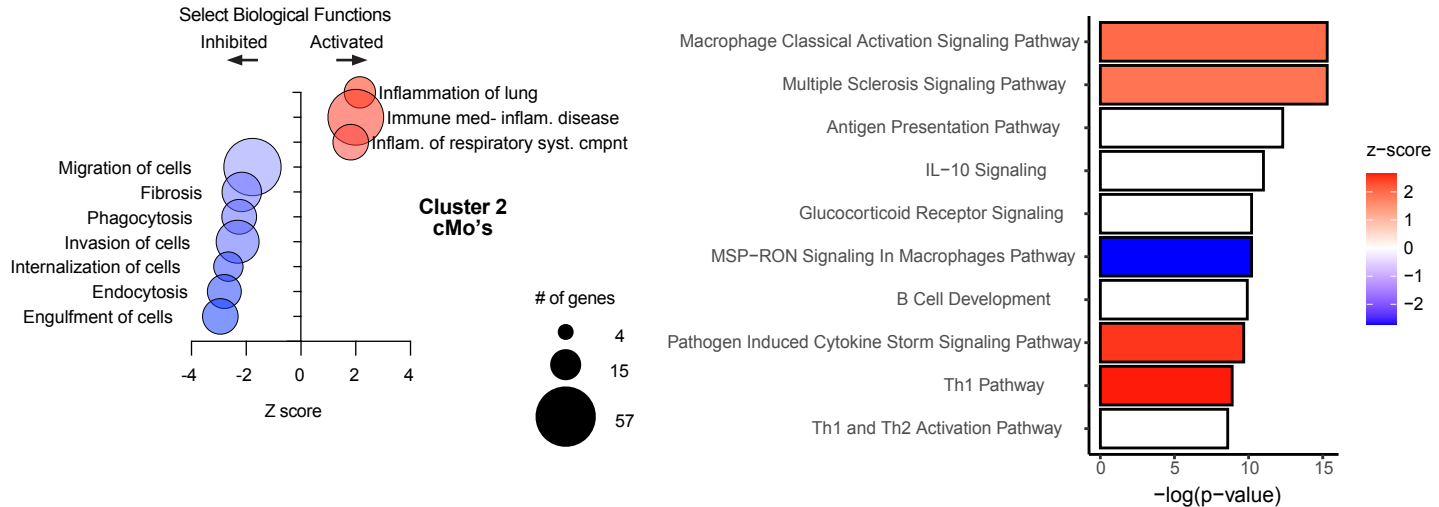


Classical Monocytes Cluster 2 (COPD vs ctl)



Supplementary Fig. 1. Top predicted downstream biological functions (A) and canonical pathways (B) using Ingenuity Pathway Analysis of differentially expressed genes in COPD vs control cells from mixed classical monocyte cluster 2 (n = 160, Padj<0.05, |log₂FC|>0.5).