Supplementary Material

Supplemental Figure S1

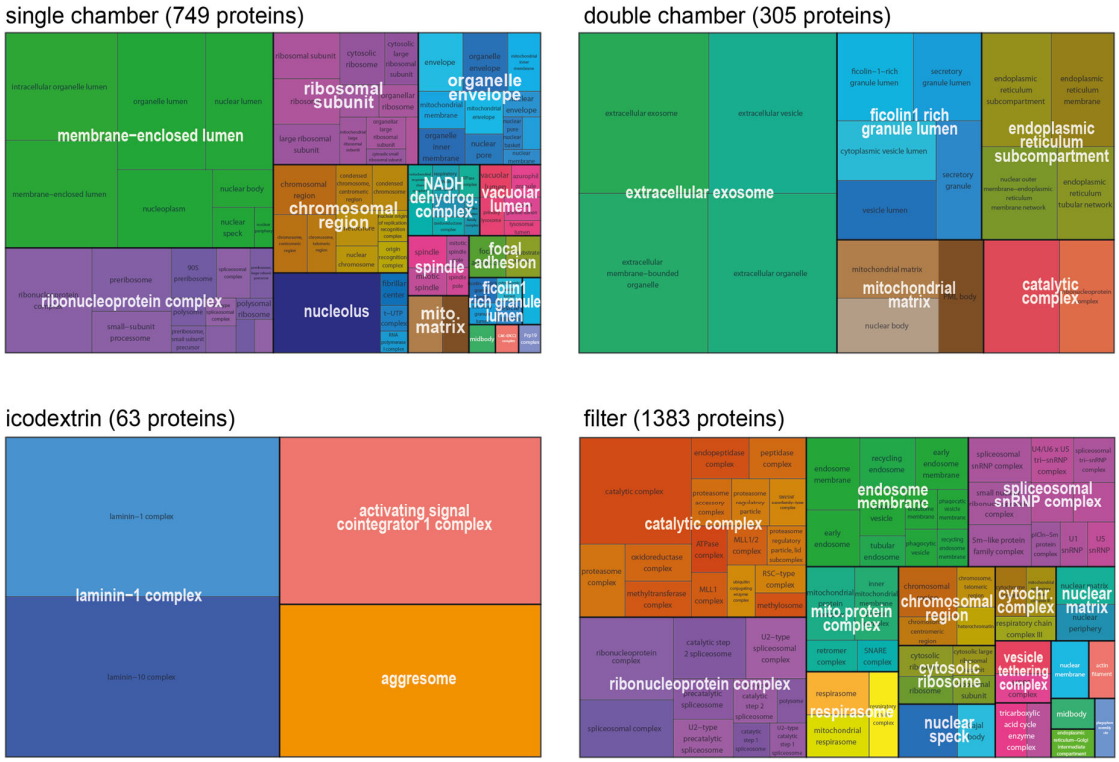

Figure S1. Gene ontology (GO) cellular component analysis of the proteins identified to be significantly regulated by individual PD fluids. Only GO categories with fold enrichment >2 were further analyzed with the Revigo tool to summarize terms in a parental term, the results are visualized as tree maps.