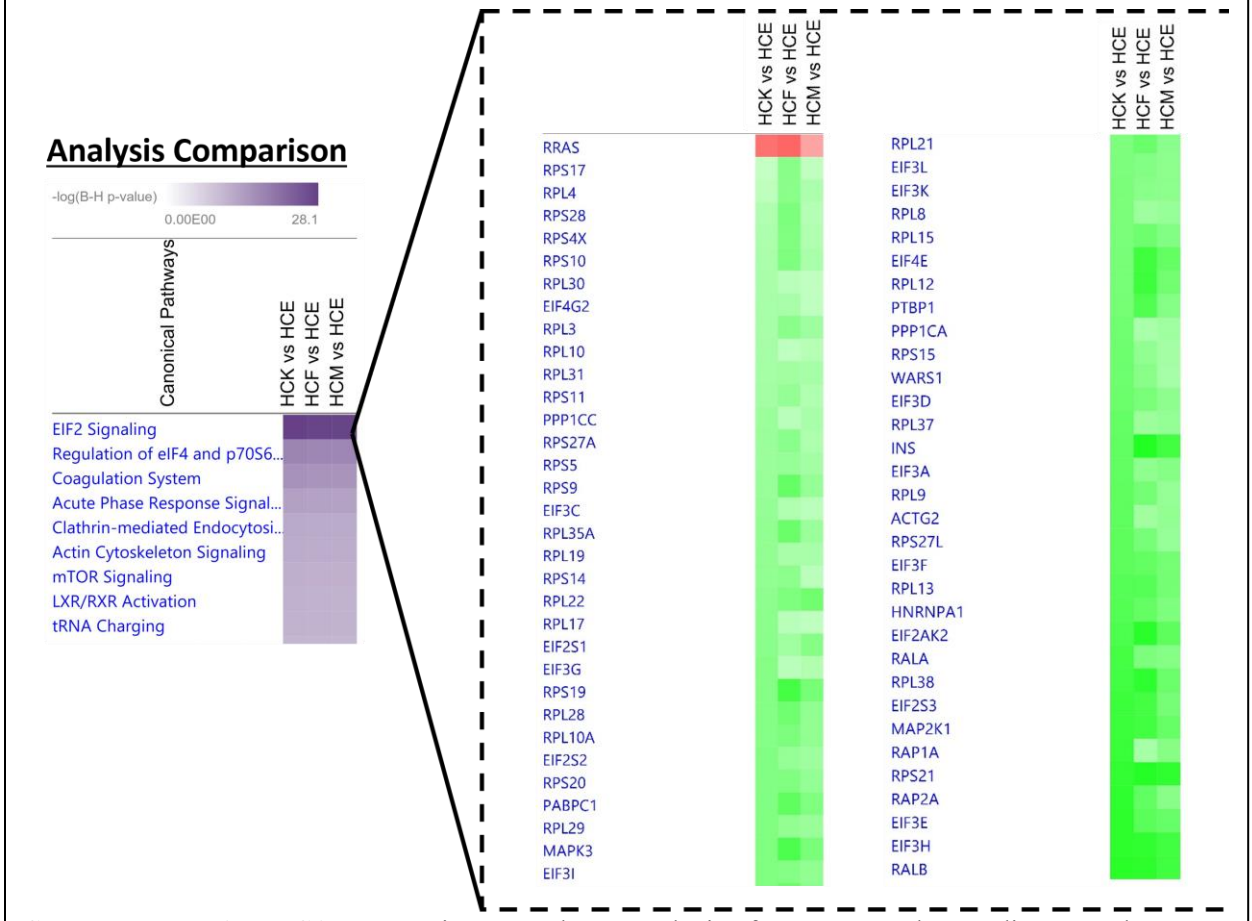

Supplemental Figure S1. Proteomic IPA pathway analysis of HCE-EVs. The top disease and biological functions were identified from IPA by using the dataset of 1209 protein with the EIF2 signaling pathway highlighted. The top scoring proteins were identified in HCE-EVs; green indicating elevated expressed and red indicates lower expression. Fisher exact test was used for determination of enrichment and p-value less than 0.05 was used as significant.