

Additional File 5: Figure S5

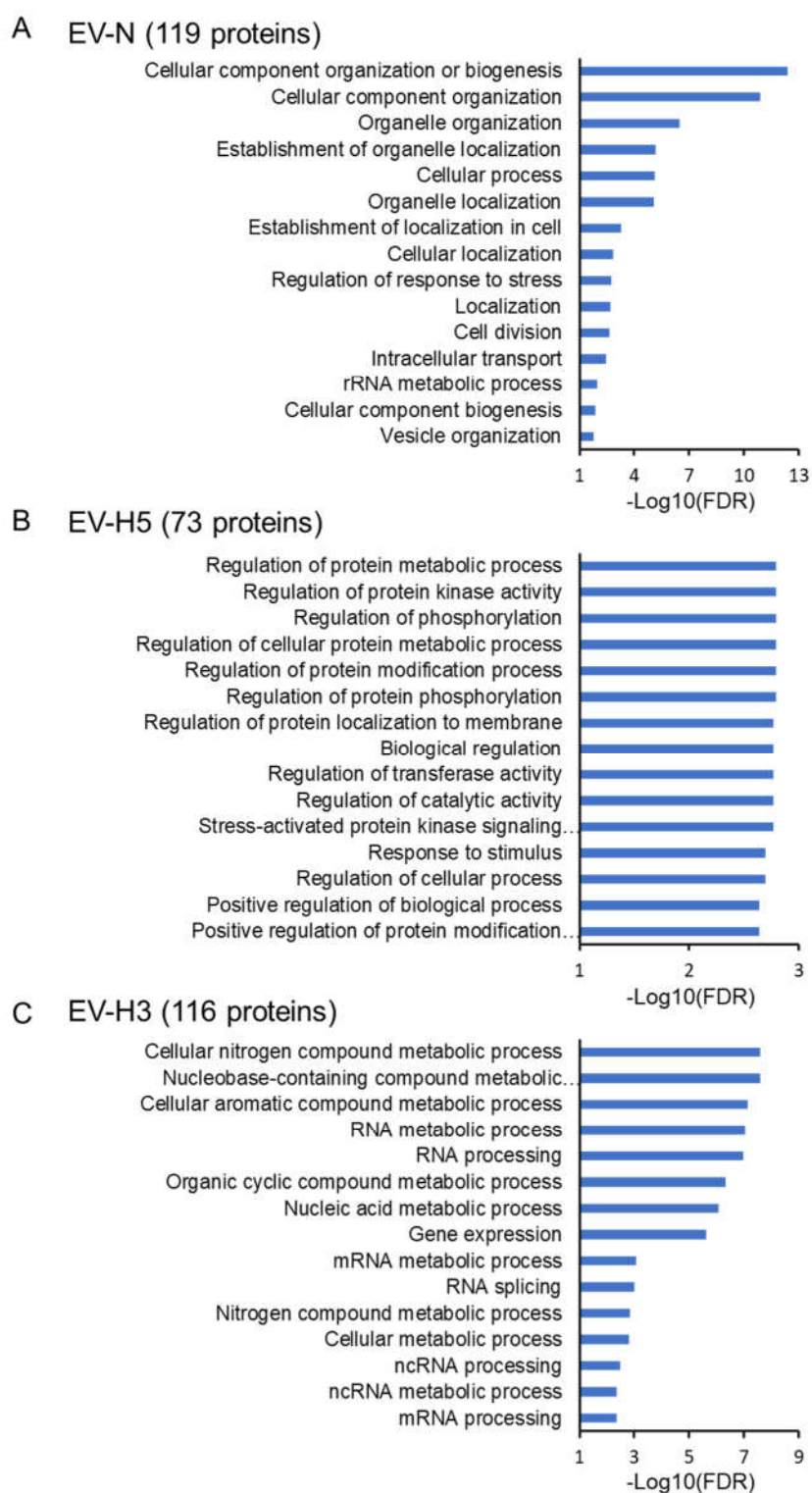


Figure S5. Pathways enrichment analysis of distinct proteins identified in hiPS-EVs derived from different oxygen conditions: (A) normoxia (EV-N), (B) hypoxia 5% O₂ (EV-H5) and (C) hypoxia 3% O₂ (EV-H3), performed in the STRING web tool. The top 15 pathways with the most significant FDR value are shown for each EV type. The number of proteins included in the analysis is indicated in the subheading of each panel.