

Additional File 3: Figure S3

A

Sample	#proteins	AVERAGE/SD	# non-redundant proteins in group
EVs_1_N	2020	2081	2615
EVs_2_N	2165	75	
EVs_3_N	2057		

Sample	#proteins	AVERAGE/SD	# non-redundant proteins in group
EVs_1_H5	1904	1952	2408
EVs_2_H5	2069	102	
EVs_3_H5	1883		

Sample	#proteins	AVERAGE/SD	# non-redundant proteins in group
EVs_1_H3	1968	2042	2566
EVs_2_H3	2171	112	
EVs_3_H3	1986		

B

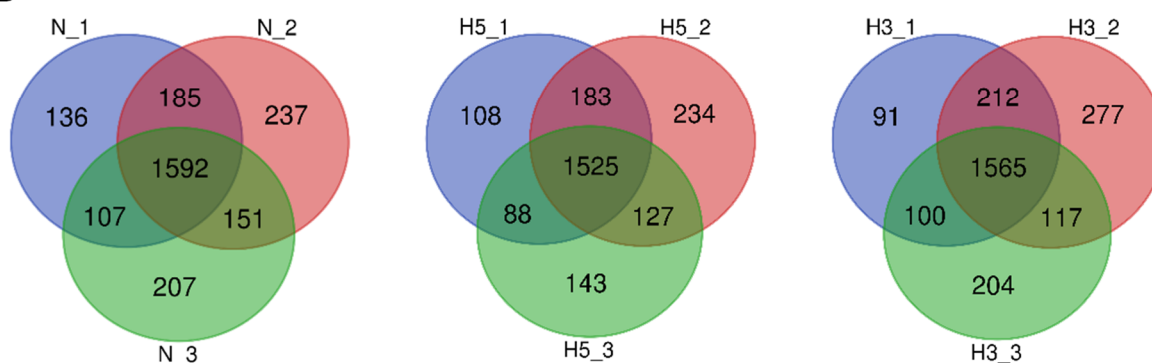


Figure S3. Proteomic analysis of hiPSC-EVs. **A.** Number of proteins identified in each EV sample, based on at least two peptides. **B.** Venn diagrams showing common and distinct proteins in three EV samples from each group. Abbreviations: N – normoxia (21% O₂); H5 – hypoxia at 5% O₂; H3 – hypoxia at 3% O₂.