

Additional File 4: Figure S4

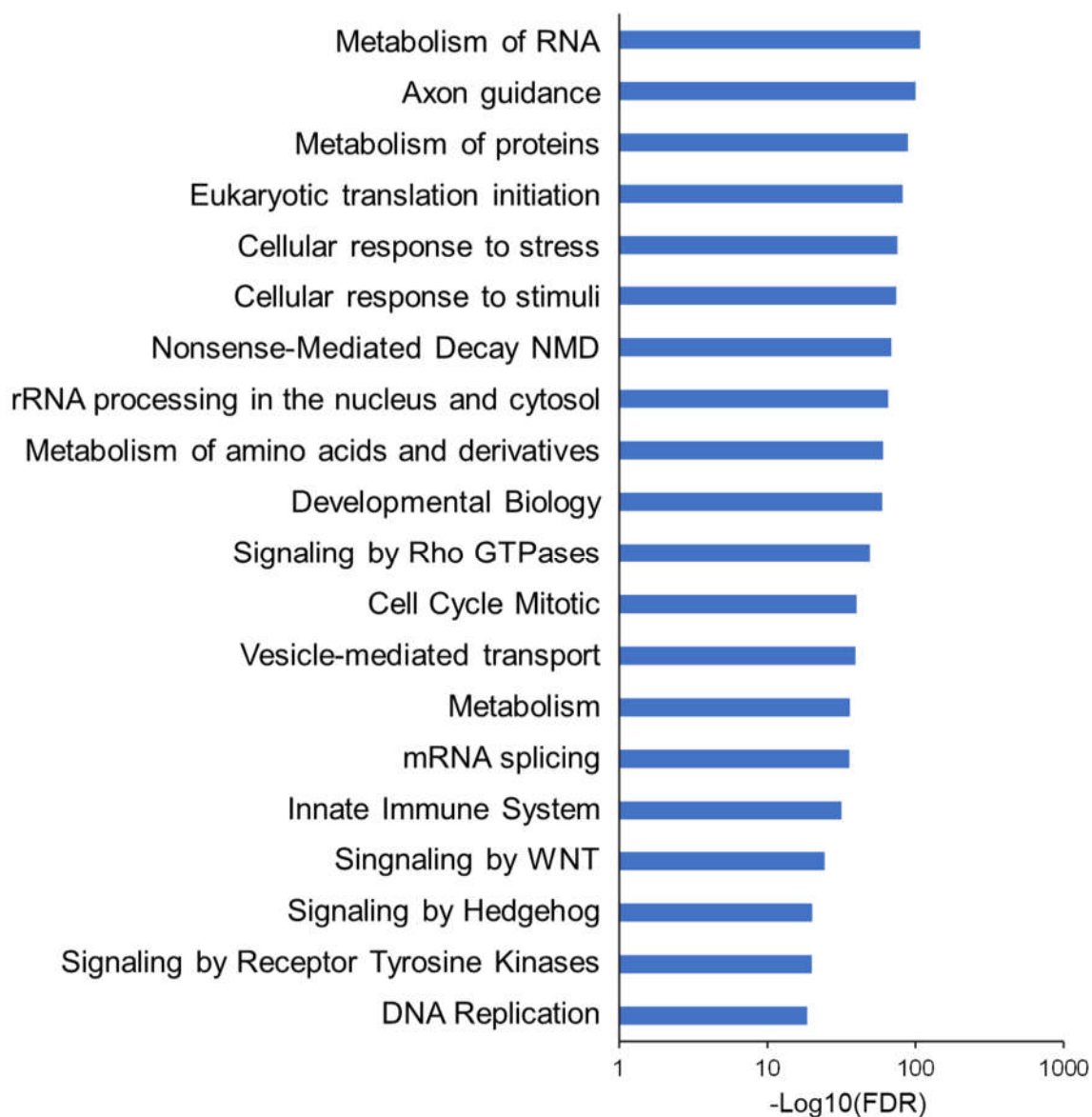


Figure S4. Pathways enrichment analysis of 1290 proteins identified as common in hiPS-EVs derived from different oxygen conditions: normoxia, hypoxia 5% O₂ and hypoxia 3% O₂ (detected in at least 6 out of 9 samples). The analysis was performed in the STRING web tool.