

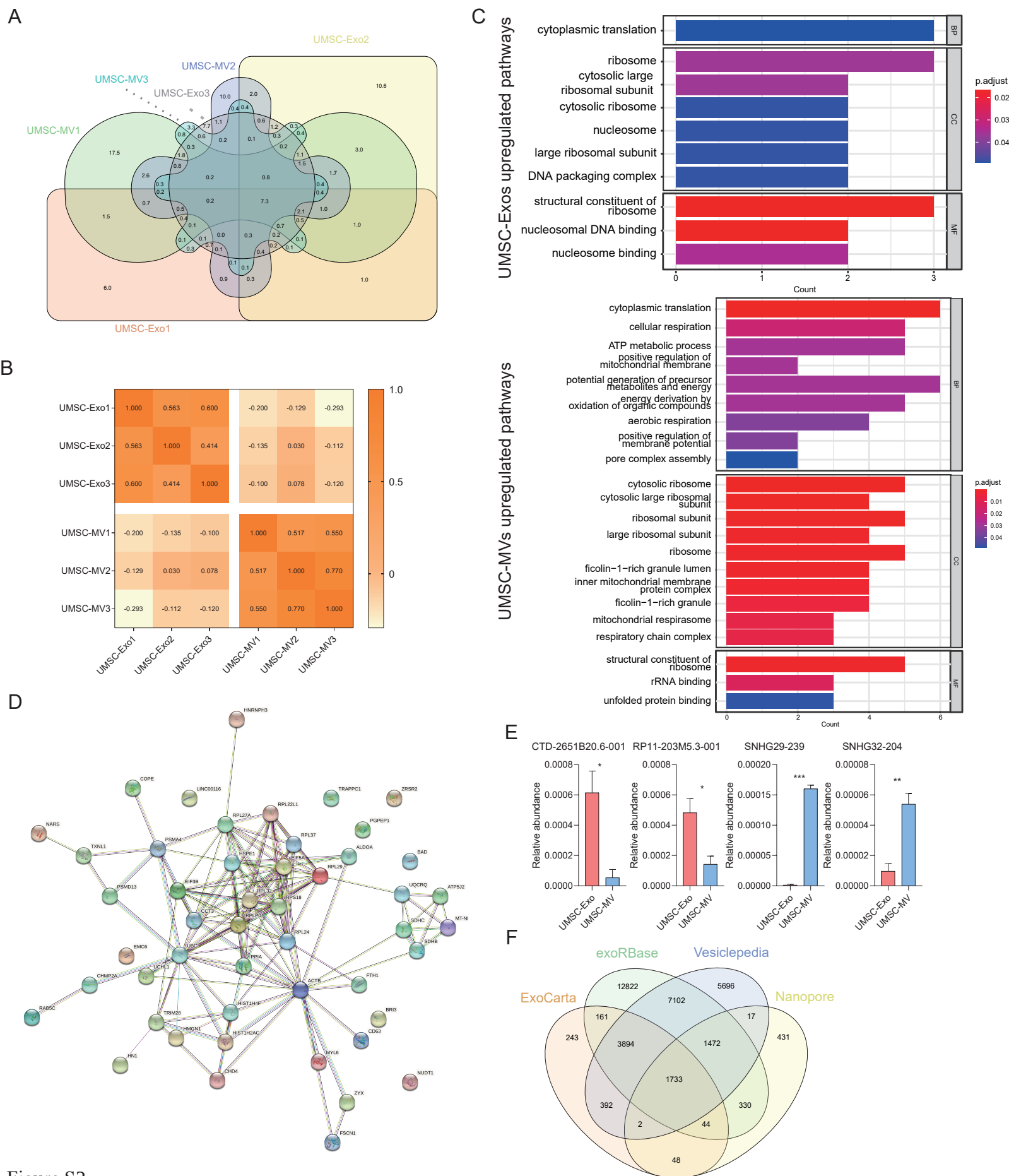


Figure S2

(A) The transcript distribution between exosomes and micro-vesicles. (B) Heatmap of Pearson correlation coefficients of different expressed transcripts between exosomes and micro-vesicles. (C) Diagram of pathways differentially expressed gene. (D) Establishment of PPI network of the differentially expressed genes using STRING database. (E) The expression of four differentially expressed lncRNAs. (F) The gene distribution between public databases and nanopore sequencing.