

Figure S3: Target analysis of the top 50 highly expressed miRNAs in ADSC-EVs. (A) Heatmap of the top 50 highly expressed miRNAs in ADSC-EVs. (B) Cytoscape network of the interaction of the top five miRNAs abundant in ADSC-EVs and their predicted target genes (blue). (C) KEGG pathway map of the predicted target genes of the top five miRNAs enriched in ADSC-EVs. (D) GO

terms of top five miRNAs of ADSC-EVs in biological process, cellular component, and molecular functions.

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