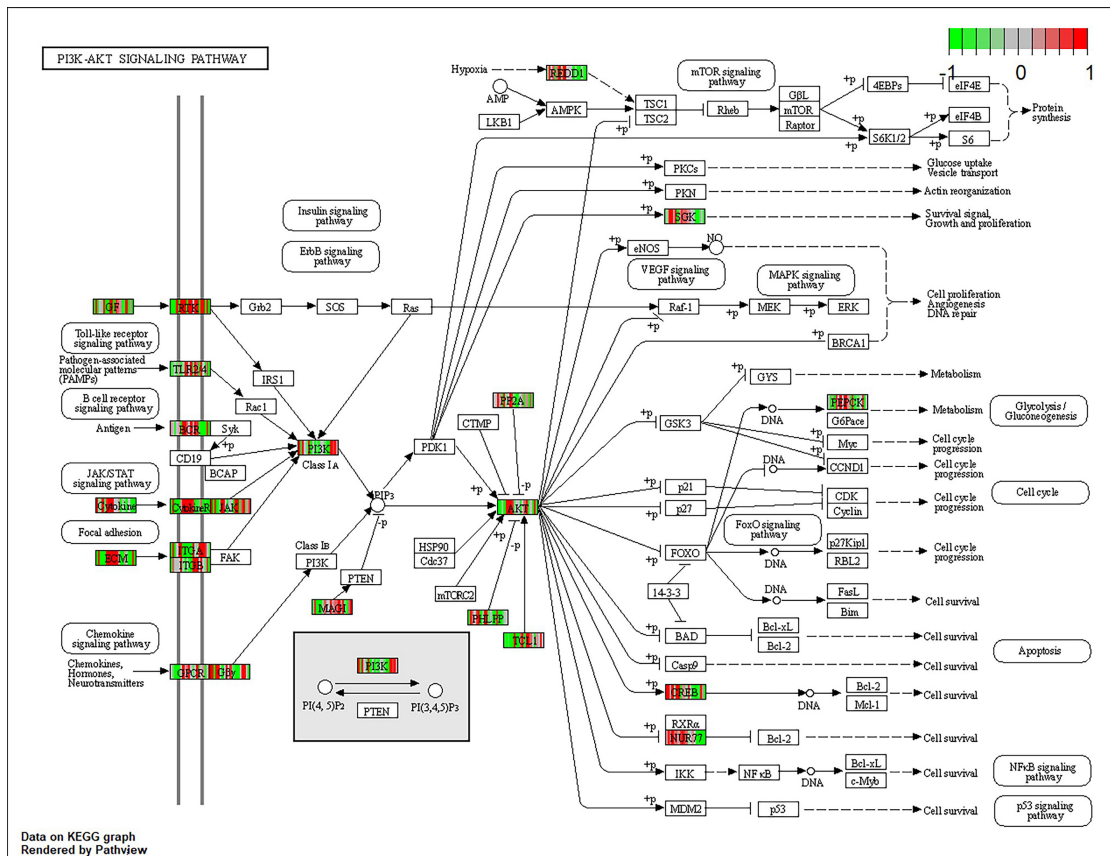
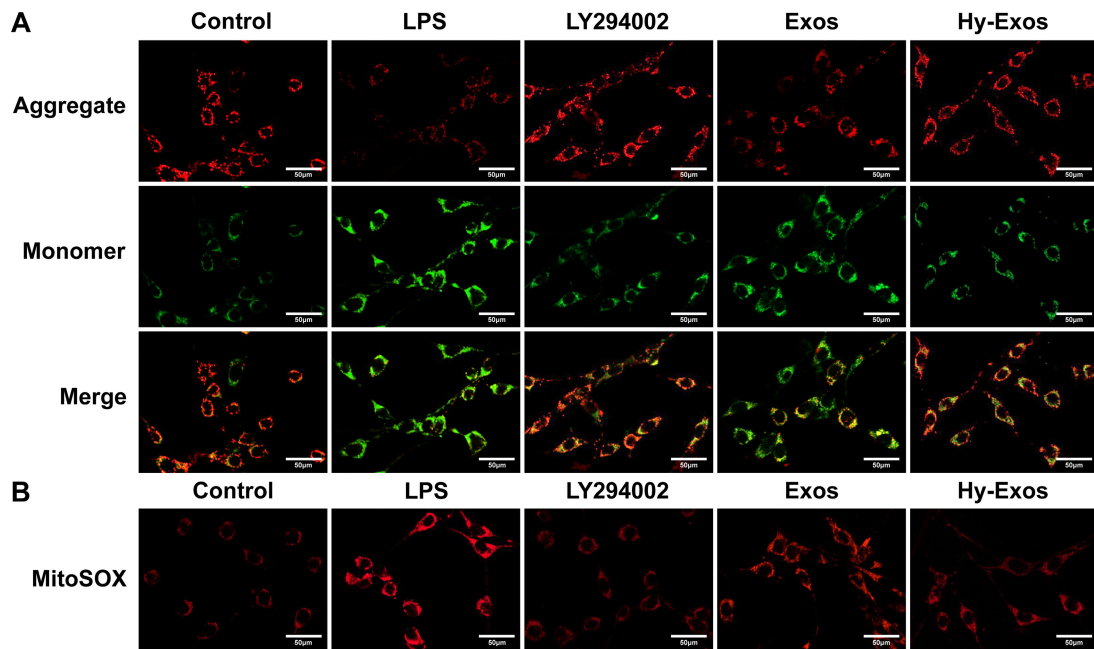


**Figure S1. Characterization of Hy-HF-MSCs.** A. Cell morphology of Hy-HF-MSCs. Scale bar = 100 μm. B. Adipogenic differentiation of Hy-HF-MSCs. C. Osteogenic differentiation of Hy-HF-MSCs. D. Flow cytometry was used to detect Hy-HF-MSC-specific antigenic markers.



**Figure S2. Enrichment results of DEGs in the PI3K-AKT signaling pathway.**



**Figure S3. The PI3K inhibitor LY294002 alleviated mitochondrial dysfunction in MODE-K cells.** A. Mitochondrial membrane potential in MODE-K cells as measured by a JC-1 staining kit. Scale bar = 50  $\mu\text{m}$ . B. Mitochondrial ROS was detected in MODE-K cells. Scale bar = 50  $\mu\text{m}$ .

Table S1. Primer sequences for quantitative real-time PCR.

Genes	Forward primer (5'-3')	Reverse primer (5'-3')
<i>IL-1<math>\beta</math></i>	GCAACTGTTCCCTGAACTCAACT	ATCTTTTGGGGTCCGTCAACT
<i>TNF-<math>\alpha</math></i>	CCTCTCTCTAATCAGCCCTCTG	GAGGACCTGGGAGTAGATGAG
<i>IL-4</i>	GCCATATCCACGGATGCGACAA	GGTGTTCCTTCGTTGCTGTGAGGA
<i>IL-10</i>	GCTCTTACTGACTGGCATGAG	CGCAGCTCTAGGAGCATGTG
<i><math>\beta</math>-actin</i>	GGCTGTATTCCCCTCCATCG	CCAGTTGGTAACAATGCCATGT
<i>miR-92b-3p</i>	GTCCGCTATTGCACTCGTCCCGGCCTCC	GTGCGTGTTCGTGGAGTC
<i>miR-484</i>	GCGTCAGGCTCAGTCCCCT	AGTGCAGGGTCCGAGGTATT
<i>miR-214-3p</i>	ACGAGAACACAGCAGGCACAG	ATCCAGTGCAGGGTCCGAGG
<i>miR-30a-5p</i>	AACGAGACGACGACAGAC	TGTAAACATCCTCGACTGGAAG
<i>miR-205-5p</i>	CGTCCTTCATTCCACCGG	AGTGCAGGGTCCGAGGTATT
<i>U6</i>	CCGTATGACCTCCTTCCACAGA	TCTGTCCACCTCTGAAACCAGG