

The level of fibrotic protein in atrial fibroblasts. (A, B) Western blot analysis of protein levels in EXO-Pacing and untreated atrial fibroblasts (*P<0.05 and **P<0.01). (C, D) Western blot analysis of protein levels in atrial fibroblasts and atrial myocytes. (*P<0.05, ***P<0.001). The levels of fibrotic proteins in atrial fibroblasts.









The level of fibrotic protein in atrial fibroblasts. (A, B) Western blot analysis of protein levels in atrial fibroblasts treated with exosomes derived from atrial myocytes (*P<0.05, **P<0.01).

- EXO-NC
- EXO-Pacing
- EXO-Pacing+miR-200a inhibitor
- EXO-miR-200a inhibitor



dephosphorylation-

Bioinformatics analysis on the relationship between miR-210-3p and fibroblast activity. (A, B) GO analysis of the differentially expressed exosomal miR-210-3p derived from atrial myocytes showed enrichment in the regulation of proteins involved in fibroblast growth factor-activated receptor activity.

Classification

Biological Process Cellular Component Molecular Function



2.81 2.5 2.19 1.88 1.25 0.94 0.62 0.31 50 100 150 200 250 300 350 400 450

-log₁₀ (P value) No.of Genes



derived exosomes. (C, D) Expression levels of miR-210-3p in atrial fibroblasts and atrial fibroblast-derived exosomes. (E) Coculture of atrial fibroblasts with exosomes derived from atrial myocytes (***P<0.001 compared with the Ctr and NC groups). (F) qRT-PCR analysis of the transfection efficiency of atrial myocytes (*P<0.05 and ***P<0.001 compared with the Ctr group). (G-I) The mRNA levels of α-SMA, collagen I and TGF β 1 were evaluated using qRT–PCR (*P<0.05).







Exosomal miR-210-3p derived from atrial myocytes promotes atrial fibroblast proliferation and activation. (A-C) Western blot analysis of protein levels in atrial fibroblasts treated with atrial myocyte-derived exosomes (*P<0.05). (D) CCK-8 assay to detect the proliferation of atrial fibroblasts treated with atrial myocyte-derived exosomes (*P<0.05 compared with EXO-NC, and #P<0.05 and ##P<0.01 compared with EXO-Pacing). (E) Immunofluorescence staining showing the relative expression levels of α-SMA (red) and DAPI (blue) in atrial fibroblasts. Scale bar: 100µm.



Effects of miR-210-3p KO on atrial fibrosis in rats. (A) qRT–PCR analysis of the expression of miR-210-3p in various organs of rats (n=7; *P<0.05 and **P<0.01 compared with the WT group). (B, D) Western blot analysis of TGF β 1, α -SMA and Collagen I expression levels in atrial tissues (n=7; *P<0.05 and **P<0.01 compared with the KO+Ang II group). (C) qRT–PCR analysis of TGFβ1, α-SMA and Collagen I expression levels in atrial tissues (n=7; **P<0.01 and ***P<0.001 compared with the KO+Ang II group). (E, F) Immunohistochemical staining for α-SMA; scale bar: 50 μm (n=7; ***P<0.001 compared with the KO+Ang II group).







RNA sequencing analysis. (A, B) RNA-Seq analysis of differentially expressed mRNAs showing that GPD1L regulates atrial fibroblast proliferation and activation.



miR-210-3p promotes proliferation and collagen synthesis by targeting GPD1L/PI3K/AKT. The expression levels of proteins (A, B) and mRNAs (C) in the si-NC and si-GPD1L groups (*P<0.05 and **P<0.01). (D, E) Western blot analysis of PI3K, p-AKT and AKT levels in atrial fibroblasts (*P<0.05). (F) The ratio of p-AKT/AKT levels (*P<0.05). (G-I) qRT-PCR analysis of α -SMA, collagen I and TGF β 1 expression in atrial fibroblasts (*P<0.05, **P<0.01, and ***P<0.001).