



Supplementary Material

Multiparametric evaluation of post-MI small animal models using metabolic ($[^{18}\text{F}]\text{FDG}$) and perfusion-based ($[^{18}\text{F}]\text{Fkardio}$) heart viability markers

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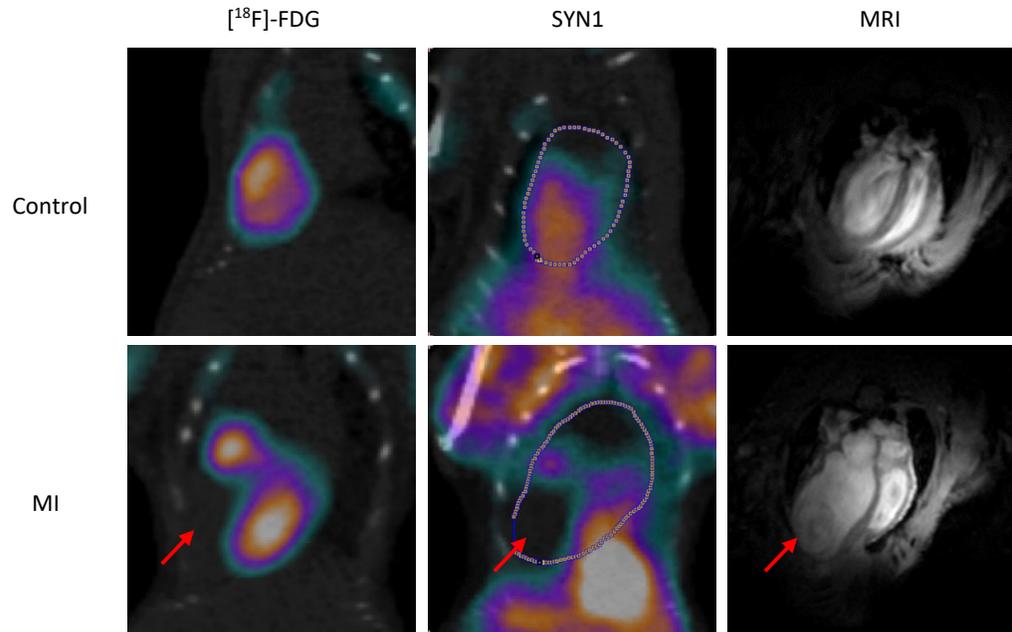


Figure S1. Comparison of molecular imaging of control and MI mice.

Control mouse (top row) and post MI mouse (bottom row). Note the difference in the loss of [¹⁸F]-FDG and SYN1 uptake in the apex region. MRI imaging confirms the damage of the heart muscle left ventricle, which is also visible in the images. The size of post MI heart is bigger due to the organ remodeling.

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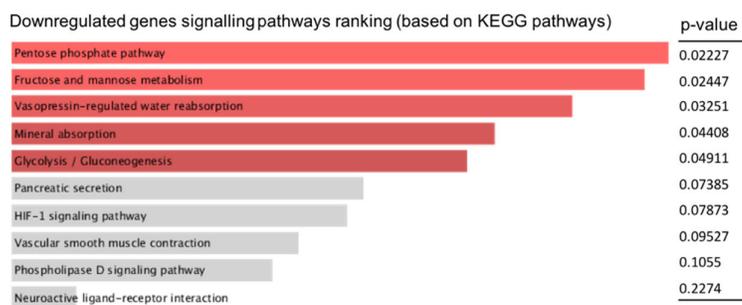
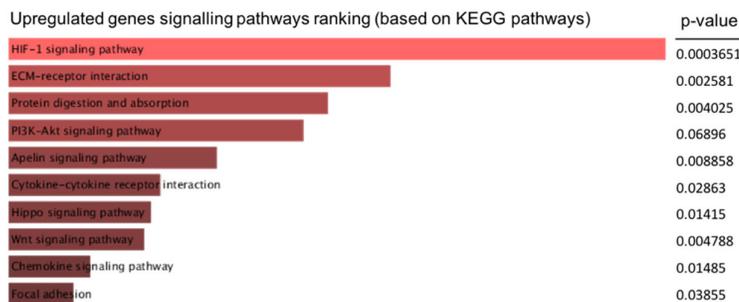
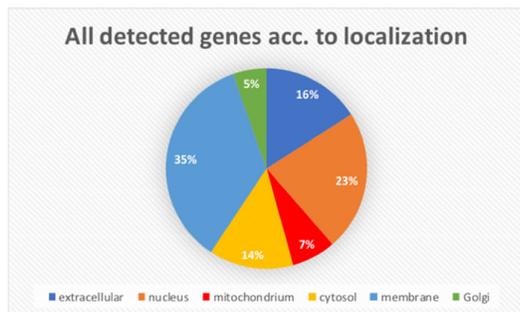


Figure S2. Summary of the expression data between MI and control groups

- A. Localization of the differentially expressed genes (DEG) shows that the major changes occur in the extracellular and membrane compartments, accompanied with the nuclei expression changes.
- B. Major up- and downregulated signaling pathways in post MI group. Signaling pathway ranking is based on reversed p-values. P-values are presented next to the chart.