

WWP2 Loss-of-Function in macrophages ameliorates pathological cardiac fibrosis

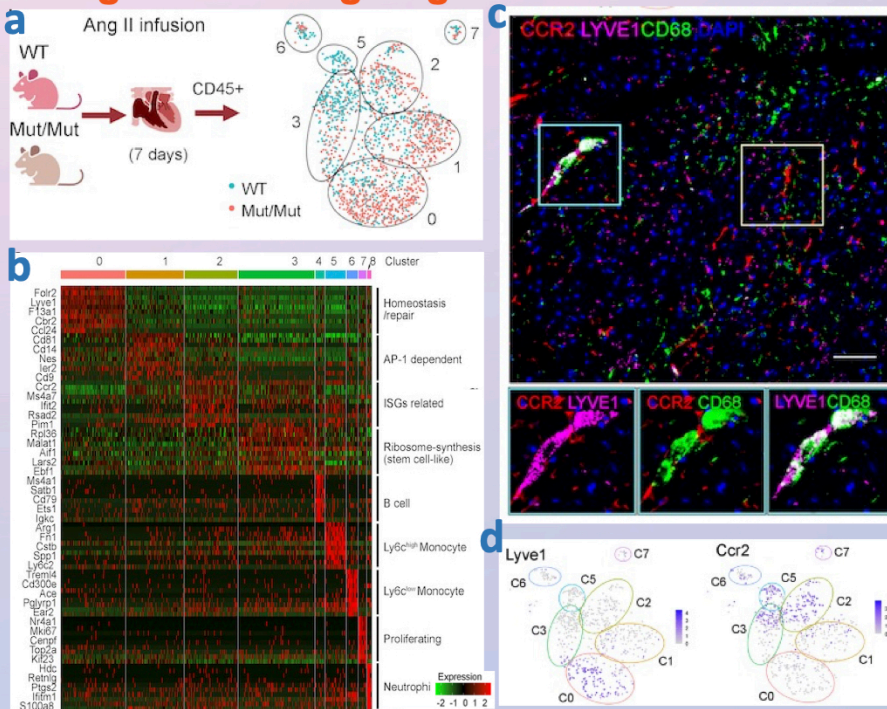
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Background

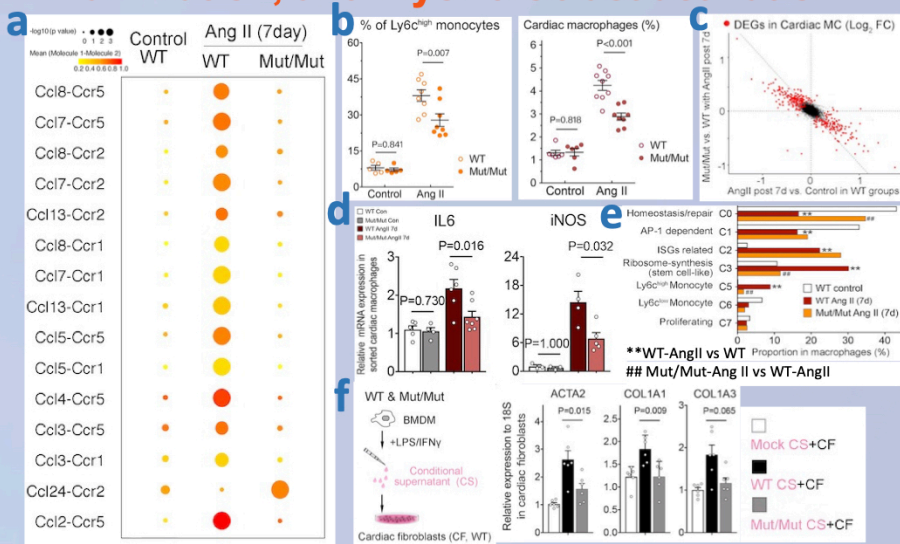
Cardiac fibrosis is the end-stage pathology of many cardiac conditions. We previously showed that the E3 ubiquitin ligase, WWP2, regulates a pro-fibrotic network conserved across different cardiac diseases¹. However, the role of WWP2 in macrophages is unknown. Utilizing single-cell RNA sequencing of a murine model of hypertensive heart disease with or without WWP2-KO, we investigated the role of WWP2 in regulating macrophages during cardiac fibrosis. Transcriptomic analysis showed a downregulation of inflammation-related genes in the macrophages of WWP2-KO mice. Crucially, the therapeutic effect of WWP2 LOF was maintained in macrophage-specific-WWP2-KO mice. Lastly, IRF7 was identified as a mechanistic target for WWP2 in macrophages.

1) Profiling cardiac macrophages during fibrogenesis using Ang-II infusion model



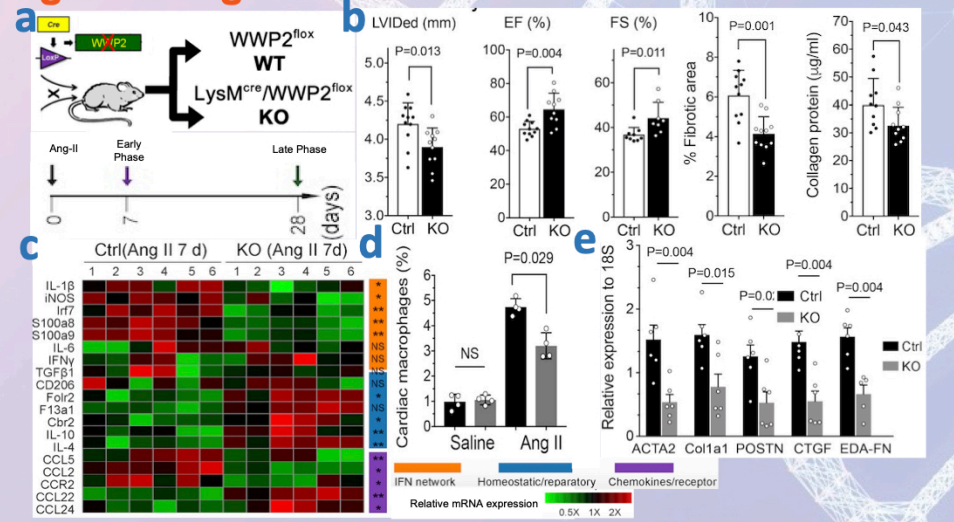
a) scRNA-sequencing of CD45⁺ macrophages from murine left ventricles (LV), b) Heatmap of top marker genes for macrophage clusters, c) Immunofluorescence of macrophage markers in murine LV, d) UMAP visualization of key marker expression

2) WWP2 LOF reduces chemokine signalling, inflammation, and myofibroblast activation



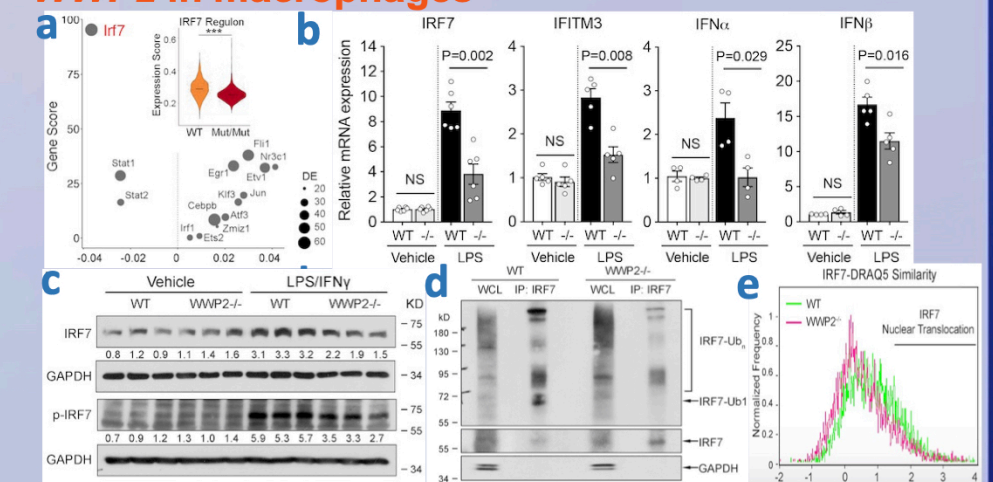
WWP2 LOF downregulates a) chemokine signaling (scRNA ligand-receptor analysis), b) % of Ly6c^{hi} and cardiac macrophages (flow cytometry, 7 day post Ang-II infusion), c) Ang-II induced signature (scRNA DE analysis), d) key inflammatory genes (qPCR), e) proportion of inflammatory Ly6c^{hi} monocytes (while upregulating proportion of homeostatic macrophages), f) myofibroblast activation

3) Macrophage-specific WWP2 LOF is protective against Ang-II induced cardiac fibrosis in vivo



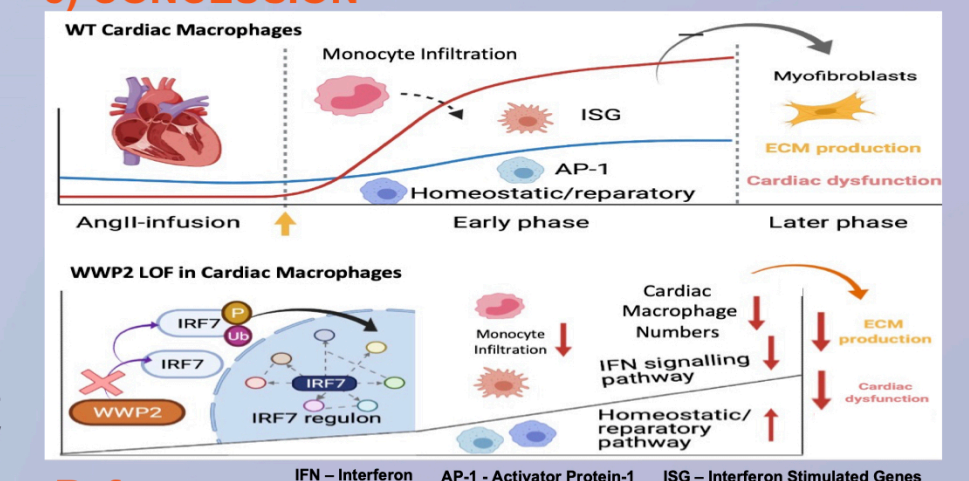
a) Cre/LoxP generation of LysM^{cre}/WWP2^{lox} mice and experimental schematic, b) Macrophage-specific WWP2 KO improves ejection fraction (EF), fractional shortening (FS), and reduces left ventricular internal dimension (LVID) end diastole, % fibrotic area, and collagen (28 days), c) Heatmap of qPCR of key inflammatory, homeostatic, and chemokine genes (7 days), d) % Cardiac macrophages (7 days), e) qPCR of key fibrogenic genes in murine LVs (28 days)

4) IRF7 is a potential mechanistic target of WWP2 in macrophages



a) Gene score vs differential expression (KOVsWT) of identified regulons, b) qPCR of IRF7 related genes, c) Western blot of IRF7/p-IRF7 in WT/WWP2^{-/-} BMDMs +/- LPS/IFN γ , d) Ubiquitination analysis of IRF7 in BMDMs, e) Imaging flow cytometry of non-nuclear/nuclear localization of WT/WWP2^{-/-} BMDMs

5) CONCLUSION



References

1. Chen, H., Moreno-Moral, A., Pesce, F., Devapragash, N., Mancini, M., Heng, E.L., Rotival, M., Srivastava, P.K., Hamston, N., Shkura, K. and Rackham, O.J., 2019. WWP2 regulates pathological cardiac fibrosis by modulating SMAD2 signaling. *Nature communications*, 10(1), pp.1-19.