

## **Lineage Tracing Suggests Possibility for Adulthood Activation of Neural Crest Genetic Signature and Common Origin for Different Cardiac Progenitor Cell Types in the Atrial Appendages**

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### **Background:**

We recently identified three distinct cardiac progenitor cell (CPC) populations from the left atrial appendages (LAAs) of adult murine hearts. These populations have distinct phenotype and characteristics. Type A CPCs are c-kit+/Sca-1+, Type B CPCs are c-kit+/CD45+ and more mature Type C CPCs are spontaneously derived from Type B in vitro. The purpose of this study was to further analyze the connection between different progenitor cell populations using gene profile data, histology analysis and lineage tracing.

### **Methods:**

To analyze the transcriptomes of CPC populations, we performed whole-genome expression array experiments to the cell sorted Type A, Type B and Type C CPC populations. The LAA analyzed with immunostaining and tissue culture derived cells were analyzed using immunostaining and FACS. Neural crest lineage tracing was performed with Pax3-cre eYFP reporter mice.

### **Results**

Transcriptome comparison demonstrated significant up-regulation of neural crest differentiation –pathway genes in Type A CPCs and also in Type C CPCs, which are derived from Type B CPCs. Type B CPCs expressed a gene profile, which matches that of non-hematopoietic, resident macrophage –like cells found in other tissues. Tissue analysis after Pax3 lineage tracing demonstrated traced cells (eYFP) in the subepicardial zone of the LAA, co-stained with F4/80 macrophage marker and c-kit. Closely attached were epicardially lined F4/80+ cells with no eYFP signal. In tissue culture, Pax3-cre-eYFP signal was strongest in the Type B CPC population and lower in the Type A CPC population. Fibroblasts in the culture were eYFP negative.

### **Conclusion**

Our results suggest that rather than having distinct embryonic origins, the different types of CPCs in the LAA originate from the resident, non-hematopoietic macrophage progenitor population. The neural crest differentiation pathway can be activated in these cells during adulthood, possibly in cardiac stress situations, enabling differentiation to cardiac lineage.