

Supplementary Fig 1 . *C-jun, creb1* and *jnk1* expression during cardiac regeneration by ISH analyses



Supplementary Fig 2. Wnt2bb expression during cardiac regeneration by ISH analyses.



Supplementary Fig 3. Expression of *jun, jnk, creb* genes involved in noncanonical Wnt pathway analysis during cardiac regeneration.



Supplementary Fig 4. Identification of transgenic fish and gene expression efficiency



Supplementary Fig 5. *pka* activated in noncardiomyocyte cells during heart regeneration.



Supplement Fig 6. Wnt2bb dominant hearts have no affection on endocardium, epicardium, and fibroblasts.