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A Hippo "AKT" Regulates Cardiomyocyte Proliferation

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Heart failure is a costly and deadly disease affecting over 23 million patients worldwide¹. At the core of the pathophysiology of heart failure is the inability of the adult mammalian heart to regenerate following myocardial loss, which is in marked contrast to terelost fish^{2–4}, urodele amphibians^{5–8}, and mammalian neonates^{9, 10}. In mammals most of the cardiomyocytes are permanently withdrawn from cell cycle soon after birth, and despite extensive efforts to identify regulators of cardiomyocyte cell cycle in mammals^{11–15}, the signaling cascades that activate/repress cell cycle in mammalian cardiomyocytes remain unclear.

Hippo signaling, which is an evolutionary conserved pathway that regulates cell proliferation, survival differentiation, and organ size¹⁶, has received significant attention in the field of regenerative medicine recently. Upon activation of the Hippo pathway in mice, Ste20 family kinases MST1/2 interact with scaffolding protein WW45 to phosphorylate and activate LATS1/2 kinases which in turn forms a complex with its cofactor MOB1 to phosphorylate and inactivate YAP and TAZ, the downstream transcriptional regulators of the signaling pathway. In contrast, when the Hippo pathway is inactivated, YAP and TAZ accumulate in the nucleus to interact with TEA domain (TEAD) family of proteins, along with other proteins such as SMADs, OCT4 or AMOT, to promote gene expression for cellular proliferation and organ growth¹⁷.

Recently, several studies have uncovered the importance of Hippo signaling pathway in heart development and regeneration. Knock-down of an upstream effector of Hippo cascade, Salv (WW45)¹⁸, and also forced expression of a constitutively active form of YAP (S127A in human and S112A in mouse) in the fetal heart, promoted cadiomyocyte proliferation and thickening of myocardial wall^{19, 20}. In contrast, specific deletion of Yap in cardiomyocytes resulted in cardiac hypoplasia and lethality^{19, 20}, demonstrating the necessity of Hippo signaling pathway in cardiomyocyte proliferation during embryonic development. The significance of Hippo pathway in postnatal cardiac homeostasis and repair has also been demonstrated where cardiomyocyte specific knockout of Yap and/or Taz results in lethal cardiomyopathy¹³, whereas expression of YAPS112A in mouse heart stimulated postnatal re-activation of cardiomyocyte proliferation and enhanced cardiac function in mice after myocardial infarction (MI) injury ^{13, 21}. In addition, deletion of Salv or Lats1/2 in postnatal mice with postnatal day 7 apex resection or adult with MI promotes heart regeneration²².

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These results indicate that Hippo signaling is a potentially important target for promoting myocardial regeneration.

Despite these results, downstream mediators of Hippo signaling pathway that regulate cardiomyocyte proliferation are not fully understood. Previous studies using constitutively active YAP revealed that the Hippo-YAP pathway augments insulin-like growth factor (IGF) signaling, which in turn induces activation of the PI3K–AKT pathway¹⁹. Phosphorylated AKT inactivates GSK-3 β by increasing its phosphorylation, leading to the stabilization of β -catenin, which in turn is required for Yap-mediated proliferation. This pathway has been well studied in a variety of disease models such cancer and diabetes^{23–25} ^{26, 27}.

In this issue, Lin et al^{28} reported the identification of one of the direct transcriptional targets of YAP, an isoform of PI3K catalytic subunit PIK3CB, that regulates cardiomyocyte proliferation as a downstream mediator of Hippo-YAP signaling. ChIP-seq analysis combined with three different systems - (1) overexpression of YAP in cardiomyocyte-like HL cells, (2) overexpression of YAP in rat neonatal ventricular cardiomyocytes (NRVMs), and (3) cardiomyocyte specific Yap homozygous knockout, identified YAP-binding sites which are significantly enriched in genes related to heart development. Among candidate YAP target genes, the authors focused on *Pik3cb*, the function of which has not been understood thus far. A YAP-bound region containing conserved TEAD motif, was in the first-intron of *Pik3cb*, through which YAP-TEAD activates transcription of *Pik3cb*. Importantly, the authors showed that forced-expression of PIK3CB is sufficient to activate PI3K-AKT pathway, and hence to regulate cardiomyocyte proliferation. Adenoviral transfection of *Pik3cb* to NRVMs, as well as in vivo overexpression of YAP driven by cTNT promoter in neonatal cardiomyocyte using AAV9, both significantly activated AKT by triggering the phosphorylation of AKT, and induced cardiomyocyte proliferation assessed by BrdU uptake and immune-staining with phosphorylated histone H3 (pH3) antibody. Mice with cardiomyocyte-specific Yap deletion showed reduced phosphorylated AKT but not total AKT, which is consistent with the findings in cancer cell lines²⁹ and neonatal cardiomyocytes¹⁹.

Moreover, Lin et al showed that *Pik3cb* is necessary for Yap-mediated activation of AKT and cardiomyocyte proliferation. AAV9-mediated overexpression of YAP together with scrambled control or specific shRNA targeting *Pik3cb* showed that while YAP overexpression (with scrambled *Pik3cb* shRNA) promoted cardiomyocyte proliferation as previously described^{13, 19, 20}, addition of *Pik3cb* shRNA resulted in a diminished effect of YAP overexpression on AKT phosphorylation and cardiomyocyte proliferation. Although these are convincing results, utilizing a *Pik3cb* knockout model would eliminate the potential off-target effects of shRNA. Finally, the authors show that AAV9-mediated overexpression of PIK3CB in the cardiomyocyte-specific Yap knockout mice induced cardiomyocyte proliferation, improved contractile function, and attenuated cardiomyocyte hypertrophy to an extent, demonstrating that *Pik3cb* can partially rescue the Yap knockout phenotype in cardiomyocytes. In summary, Hippo-YAP mediated activation of PI3K/AKT pathway, along with cardiomyocyte growth, is at least in part mediated by direct transcriptional activation of *Pik3cb* by Hippo-signaling mediator YAP/TEAD complex.

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Despite these important findings, some questions remain. For example, much of the terminal effect on cell cycle is attributed to p27, however the mechanism of regulation of p27 or the effect on other CDK Inhibitors has not been fully examined. In addition, in the loss of function studies both decreased proliferation and survival are noted. However, the isolated effects on cell survival and proliferation are not clearly dissected, although admittedly this may be quite a difficult task. Finally, it would be important for future studies to examine upstream regulators of Hippo-YAP pathway, and how this pathway is regulated in the postnatal heart. Nevertheless, this report demonstrates that *Pik3cb* is an important link between Hippo-YAP and PI3K-AKT pathways, and brings us one step closer to an understanding of molecular mechanism regulating cardiomyocyte growth and proliferation.

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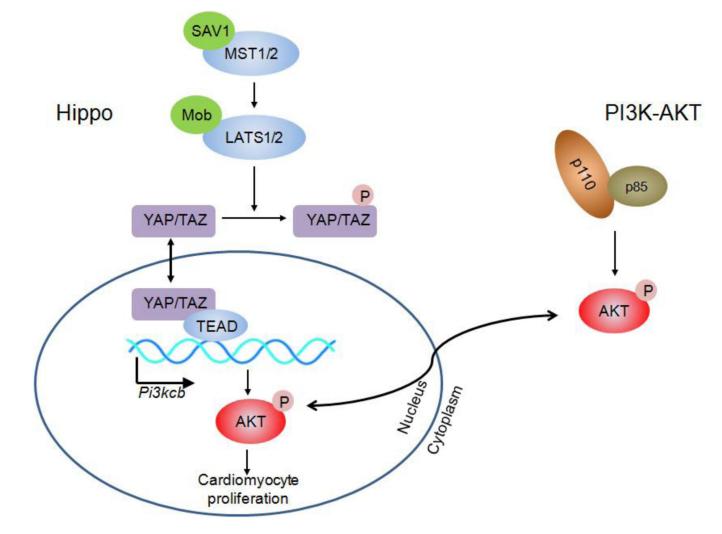


Figure 1. Hippo and PI3K-Akt signaling pathways stimulate the cardiomyocyte proliferation *Pik3cb* is a gene encoding for the catalytic subunit p110 of the Class IA PI3K. Lin et al found that *Pik3cb* is a direct target of YAP which links Hippo and PI3K-Akt signaling pathways to stimulate cardiomyocyte proliferation.