Protein kinase A activates the Hippo pathway to modulate cell proliferation and differentiation

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The Hippo tumor suppressor pathway plays an important role in tissue homeostasis that ensures development of functional organs at proper size. The YAP transcription coactivator is a major effector of the Hippo pathway and is phosphorylated and inactivated by the Hippo pathway kinases Lats1/2. It has recently been shown that YAP activity is regulated by G-protein-coupled receptor signaling. Here we demonstrate that cyclic adenosine monophosphate (cAMP), a second messenger downstream from Gαs-coupled receptors, acts through protein kinase A (PKA) and Rho GTPTases to stimulate Lats kinases and YAP phosphorylation. We also show that inactivation of YAP is crucial for PKA-induced adipogenesis. In addition, PKA activation in Drosophila inhibits the expression of Yorki (Yki, a YAP ortholog) target genes involved in cell proliferation and death. Taken together, our study demonstrates that Hippo–YAP is a key signaling branch of cAMP and PKA and reveals new insight into mechanisms of PKA in regulating a broad range of cellular functions.

[Keywords: Hippo, PKA, YAP, adipogenesis; proliferation]

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the Hippo pathway through G-protein-coupled receptor (GPCR) signaling (Mo et al. 2012, Yu et al. 2012a,b). GPCR is the largest family of cell surface receptors encoded in the human genome and has been implicated in almost every aspect of physiological regulation. We observed that hormonal factors like LPA, S1P, and Thrombin can activate Gα12/13 to stimulate YAP/TAZ, which mediate the effect of these signals on gene expression, cell proliferation, and migration (Mo et al. 2012; Yu et al. 2012b). Similar observations were also reported by Wu and colleagues (Miller et al. 2012). In contrast, ligands of Gαs-coupled receptors, such as epinephrine and glucagon, stimulate Lats1/2 and result in inhibition of YAP/TAZ (Yu et al. 2012b). These findings suggest that the activity of YAP/TAZ can be positively or negatively modulated by a wide range of extracellular signals via GPCRs in a manner dependent on which intracellular signaling pathway is stimulated.

Activation of Gαs-coupled receptors usually results in accumulation of cyclic adenosine monophosphate (cAMP), an important second messenger with diverse physiological functions, including cell proliferation and differentiation (Cho-Chung 1990). Despite extensive studies, the precise molecular mechanisms of how cAMP regulates cell proliferation and differentiation is not fully understood (Stork and Schmitt 2002). In this study, we demonstrate that cAMP acts through protein kinase A (PKA, cAMP-dependent protein kinase) and Rho GTPases to stimulate Lats kinase activity and inhibit YAP/TAZ. Inhibition of YAP/TAZ is critical for cAMP and PKA to promote adipogenesis and suppress growth, establishing Hippo-YAP as a signaling branch downstream from cAMP and PKA.

**Results**

**cAMP signaling stimulates YAP phosphorylation**

Activation of Gαs-coupled receptors can stimulate adenyl cyclase (AC) and result in an increase of cAMP production (Sassone-Corsi 2012). We treated MDA-MB-231 breast cancer cells with epinephrine, a ligand for β2 adrenergic receptor that increases cAMP (Supplemental Fig. S1). As anticipated, we observed a transient induction of phosphorylation of the cAMP response element-binding protein (CREB), a direct target of PKA and an indicator of cAMP levels (Supplemental Fig. S1). Many pharmaceutical drugs are direct PDE inhibitors that can be used to increase cellular cAMP levels (Supplemental Fig. S1; Sassone-Corsi 2012). Several nonselective PDE inhibitors (theophylline, IBMX, and ibudilast) and PDE4 selective inhibitors (rolipram) all induced YAP phosphorylation (Fig. 1D), further supporting the role of cAMP in stimulating YAP phosphorylation. In addition, these data also suggest that PDE inhibitors might be useful tools for restricting YAP activity.

Figure 1. cAMP signaling induces YAP phosphorylation and inactivation. (A, B) MDA-MB-231 cells were treated with 10 μM epinephrine (Ep) or forskolin (FSK) for the indicated durations, and cell lysates were subjected to immunoblotting using the indicated antibodies. (C) Time course of YAP and CREB phosphorylation in response to epinephrine or forskolin (the value for time 0 was arbitrarily set). (D) MDA-MB-231 cells were treated with different PDE inhibitors—ibudilast (100 μM), IBMX (100 μM), rolipram (50 μM), or theophylline (1 mM)—for 1 h, and the phosphorylation status of YAP was determined by phos-tag gels. (E) HEK293A, MCF10A, U2OS, or MEF cells were treated with or without 10 μM forskolin for 1 h. YAP phosphorylation was assessed using phos-tag gels, and the same lysates were also used to blot for TAZ protein levels. (F) MCF10A cells were serum-starved overnight and treated with 10 μM forskolin for 1 or 4 h, mRNA was extracted, and the expression level of CTGF was determined using real-time RT–PCR.
We tested the effect of forskolin on YAP phosphorylation in multiple cell lines, including U2OS, MCF10A, HEK293A, and mouse embryonic fibroblasts (MEFs). In all cases, YAP phosphorylation was increased by forskolin treatment (Fig. 1E). TAZ is a YAP homolog similarly regulated by the Hippo pathway [Lei et al. 2008]. As expected, TAZ phosphorylation was increased, as indicated by the decreased electrophoretic mobility (Fig. 1E). Moreover, TAZ protein levels were modestly reduced in forskolin-treated cells because TAZ is destabilized by phosphorylation [Liu et al. 2010]. These results indicate that the cross-talk between cAMP signaling and the Hippo pathway is a conserved phenomenon in different cell types, and both Hippo pathway effectors, YAP and TAZ, are inactivated by cAMP.

YAP/TAZ are transcriptional coactivators. To determine the functional significance of intracellular cAMP on YAP activity, we determined expression of YAP/TAZ target genes. Indeed, the expression of CTGF, which is a direct YAP/TAZ target gene, was inhibited by forskolin in MCF10A cells (Fig. 1F), further supporting the idea that cAMP inhibits YAP and TAZ activity.

cAMP signals through PKA to stimulate YAP phosphorylation

Exchange protein activated by cAMP (Epac) and PKA are two downstream effectors mediating most physiological functions of cAMP [Supplemental Fig. S1]. Epac proteins and the regulatory (R) subunits of PKA contain cAMP-binding domains that function as cAMP sensors [Gloerich and Bos 2010; Taylor et al. 2012]. We investigated whether PKA or Epac signaling mediated the effect of cAMP on YAP phosphorylation. Overexpression of the catalytic (C) subunit α (PRKACA) induced YAP phosphorylation, whereas overexpression of the PKA kinase-dead mutant decreased YAP phosphorylation (Fig. 2A). In contrast, overexpression of wild-type or constitutively active Rap1b, an effector of Epac [Gloerich and Bos 2010], did not show a significant effect on YAP phosphorylation (Fig. 2B). These results indicate that PKA rather than Epac mediates the effect of cAMP on YAP inhibition.

PKA C subunits form a complex with R subunits under basal state, and the kinase activity is restricted; cAMP binding to the R subunits induces a conformational change and releases the C subunits and therefore results in PKA kinase activation [Supplemental Fig. S1; Taylor et al. 2012]. To study the involvement of PKA on YAP inactivation, we employed mutant PKA R subunits that interact with PKA C subunits in a manner unresponsive to cAMP. When mutant PKA R subunits (Rκα and Rκα) were overexpressed, forskolin-induced YAP phosphorylation was completely blocked (Fig. 2C). We also used shRNA to knock down the PKA C subunits, and when PKA Cα expression was down-regulated, the induction of YAP phosphorylation by forskolin or epinephrine was strongly compromised (Fig. 2D). Moreover, when cells were treated with a PKA inhibitor, KT5720 [Supplemental Fig. S1], YAP phosphorylation was decreased, and the effect of forskolin and epinephrine was largely blocked by KT5720 (Fig. 2E). We also examined primary hepatocytes isolated from mice. Stimulation with glucagon, a ligand known to activate PKA, increased YAP phosphorylation (Fig. 2F). Similar to that of epinephrine, the PKA inhibitor KT5720 abolished the effect of glucagon on YAP phosphorylation (Fig. 2F). Collectively, these data establish that PKA is the key mediator of cAMP in stimulating YAP phosphorylation.
cAMP stimulates Lats kinases to induce YAP phosphorylation

YAP is phosphorylated by the Lats1/2 kinases on five serine residues within the HXRXXS motifs, including S127, and can be phosphorylated on additional sites by other kinases [Zhao et al. 2010]. To determine whether cAMP regulates the Lats phosphorylation sites in YAP, a 5SA mutant YAP (with all five Lats targeting sites mutated to alanine) was transected into cells and then treated with or without forskolin. Forskolin failed to induce a significant change in phosphorylation of the YAP-5SA mutant, as assessed by phos-tag gel [Fig. 3A], suggesting that PKA likely acts through the Hippo pathway kinases to stimulate YAP phosphorylation. To test the function of MST kinases in cAMP-induced YAP phosphorylation, MST1/2 expression was down-regulated by siRNAs, and the phosphorylation status of YAP was relatively normal in response to forskolin treatment [Fig. 3B]. Overexpression of MST2 K/R [lysine mutated to arginine], a kinase-dead mutant, resulted in a lower basal YAP phosphorylation. However, under this condition, forskolin was still capable of inducing YAP phosphorylation [Fig. 3C], suggesting that MST may not be involved in cAMP response. In contrast, when Lats1/2 expression was down-regulated by siRNAs, the basal YAP phosphorylation was lower, and, notably, forskolin-induced YAP phosphorylation was significantly impaired [Fig. 3B]. When Lats2 K/R, a kinase-dead mutant, was overexpressed, the basal YAP phosphorylation was reduced, and, importantly, the effect of forskolin on YAP phosphorylation was abolished [Fig. 3D]. These data indicate that Lats is involved in YAP phosphorylation in response to cAMP.

We next determined whether cAMP could increase Lats kinase activity. Lats1 kinase was immunoprecipitated from cells treated with or without forskolin, and in vitro Lats kinase activity was measured using purified GST-YAP as a substrate. Our results indicate that Lats kinase activity is indeed induced by forskolin [Fig. 3E]. Collectively, the above results reveal that Lats kinases are required for cAMP and PKA to induce YAP phosphorylation.

Rho GTPases are required for PKA to modulate YAP phosphorylation

The response of YAP phosphorylation to cAMP is slower than that of CREB phosphorylation [Fig. 1C], suggesting that PKA may not directly phosphorylate a core component of the Hippo pathway. Consistently, we could not activate Lats1 in vitro using purified PKA [data not shown]. Recently, it has been reported that Rho GTPases can regulate the Hippo pathway and plays a major role from Gα12/13-coupled receptors to YAP phosphorylation [Dupont et al. 2011; Miller et al. 2012; Mo et al. 2012; Yu et al. 2012b; Zhao et al. 2012]. Interestingly, PKA has been shown to modulate the actin cytoskeleton by inhibition of RhoA, which is achieved by phosphorylation of RhoA, Rho GDP dissociation inhibitor [RhoGDI], or Rho guanine nucleotide exchange factors [RhoGEFs] (Qiao et al. 2008; Meiri et al. 2009; Tkachenko et al. 2011). We hypothesized that PKA might induce Lats1/2 activity by repressing RhoA. Indeed, the phosphorylation of myosin light chain 2 [MLC2], a target of Rho-associated protein kinase [ROCK], was reduced when cells were treated with forskolin [Fig. 4A], indicating a decreased RhoA activity when PKA is activated. When cells were transfected with wild-type or constitutively active RhoA, forskolin was unable to induce YAP phosphorylation [Fig. 4B]. Complementarily, when RhoGDI, an inhibitor of Rho GTPases, was overexpressed, the PKA inhibitor KT5720 was unable to induce YAP/TAZ dephosphorylation [Fig. 4C]. Taken together, these observations suggest that RhoA is a major mediator for cAMP or PKA to regulate YAP phosphorylation, and we propose that PKA increases Lats1/2 activity and YAP phosphorylation by inhibiting Rho GTPases.
Phosphorylation of Flag-YAP and endogenous TAZ was determined. For 1 h. Phosphorylation of Flag-YAP and endogenous TAZ was determined with 10 μM forskolin for 1 h, and cell lysates were subjected to immunoblotting. Phosphorylation of MLCl2, CREB, and YAP was determined. (B) Flag-YAP was cotransfected into HEK293A cells with wild-type or constitutively active RhoA, and, after 16 h, cells were stimulated with 10 μM forskolin for 1 h before Western blotting. (C) Flag-YAP was cotransfected into HEK293A cells with or without GFP-tagged RhoGDI, and, after 16 h of incubation in serum-free medium, cells were treated with or without KT5720 for 1 h. Phosphorylation of Flag-YAP and endogenous TAZ was determined.

Figure 4. Rho GTPases mediate the effect of PKA on YAP phosphorylation. (A) MDA-MB-231 cells were treated with 10 μM forskolin for 1 h, and cell lysates were subjected to immunoblotting. Phosphorylation of MLCl2, CREB, and YAP was determined. (B) Flag-YAP was cotransfected into HEK293A cells with wild-type or constitutively active RhoA, and, after 16 h, cells were stimulated with 10 μM forskolin for 1 h before Western blotting. (C) Flag-YAP was cotransfected into HEK293A cells with or without GFP-tagged RhoGDI, and, after 16 h of incubation in serum-free medium, cells were treated with or without KT5720 for 1 h. Phosphorylation of Flag-YAP and endogenous TAZ was determined.

Hippo pathway activation is required for cAMP- or PKA-induced adipogenesis

PKA and cAMP play important roles in cell lineage specification during metazoan development [Lane and Kalderon 1993]. For instance, PKA has been shown to promote adipogenesis [Rosen and MacDougall 2006], although molecular mechanisms underlying PKA-regulated cell differentiation are not fully understood. Notably, TAZ displays activity opposite to PKA and can inhibit adipogenesis [Hong et al. 2005]. As shown above, YAP and TAZ are negatively regulated by PKA; therefore, the Hippo pathway might function downstream from PKA in regulating cell differentiation.

To test the above hypothesis, the effect of PKA or the Hippo pathway on adipocyte differentiation of murine fibroblast 3T3-L1 cells was examined. We found that YAP phosphorylation was repressed by KT5720 and induced by forskolin or IBMX in 3T3-L1 cells [Fig. 5A]. TAZ protein level was increased by KT5720 and decreased by forskolin or IBMX [Fig. 5A], consistent with TAZ degradation upon phosphorylation [Liu et al. 2010]. Adipogenesis was initiated by addition of insulin, dexamethasone, and troglitazone (Tro) with or without PKA activator or inhibitor. The formation of lipid droplets was visualized by oil red staining. As expected, IBMX induced, whereas KT5720 inhibited, adipogenesis, indicating a stimulatory role of PKA in adipocyte differentiation [Fig. 5B]. Next, we examined the effect of YAP and TAZ on adipogenesis and found that knockdown of YAP and TAZ promoted adipogenesis, an effect similar to IBMX treatment [Fig. 5C,D]. Interestingly, knockdown of YAP and TAZ also largely blocked the effect of the PKA inhibitor KT5720 [Fig. 5E]. On the other hand, YAP overexpression strongly inhibited the ability of IBMX or forskolin to induce adipogenesis [Fig. 5F]. Furthermore, the expression of multiple adipogenesis markers was abolished by YAP overexpression even in the presence of IBMX or forskolin [Fig. 5G]. Therefore, YAP/TAZ activation is required for the PKA inhibitor to suppress adipocyte differentiation, whereas YAP/TAZ inhibition is crucial for IBMX to induce the differentiation program. Taken together, our data support a model in which modulation of YAP and TAZ activity is required for PKA signaling to regulate adipogenesis.

Inactivation of Yorki (Yki) by PKA in Drosophila

We next investigated whether the Drosophila PKA ortholog (Yki) is similarly regulated by PKA. In Drosophila S2R+ cells, when the Drosophila PKA ortholog (PKA-C1) was knocked down by dsRNA [RNAi], the Yki transcription activity was significantly increased, as assessed by a luciferase assay [Fig. 6A]. Therefore, the cross-talk between PKA and the Hippo pathway might be conserved in Drosophila. During Drosophila imaginal disc development, PKA-C1 has been shown to be a potent growth inhibitor, and loss of PKA function leads to ectopic limb formation [such as wing] [Jiang and Struhl 1995; Lepage et al. 1995; Li et al. 1995; Pan and Rubin 1995]. To test the possibility that PKA-C1 might regulate Hippo signaling for growth control, expression of several Yki target genes, including expanded (ex), Cyclin E (CycE), and Diap1, was determined at the transcript level. In larval wing discs, loss of PKA-C1 activity caused an increase of expression of ex, CycE, and Diap1 [Fig. 6B]. On the contrary, overexpression of PKA-C1 resulted in a moderate yet significant reduction of expression of these genes [Fig. 6B]. Moreover, PKA-C1 overexpression induced Yki phosphorylation [Fig. 6C] and was sufficient to reduce the level of Diap1 protein and promote programmed cell death, as revealed by increased caspase 3 staining [Fig. 6D–G]. These results suggest that PKA inhibits Yki activity in developing tissues to restrict proliferation and promote apoptosis.

Discussion

The signaling relay from cAMP to the Hippo–YAP pathway

In this study, we show that cAMP acts through PKA to stimulate Lats kinase activity and YAP phosphorylation, and the Rho GTPases likely mediate the effect of PKA to Hippo–YAP regulation [Fig. 7]. Although the Hippo homolog MST1/2 may not be involved in YAP regulation in response to cAMP, we would still prefer to retain the name of Hippo given the fact that YAP/TAZ are the only known major functional output of the Hippo pathway. Our data establish Hippo–YAP as a physiologically relevant signaling branch downstream from PKA. The precise
molecular mechanism connecting Rho to Lats kinase requires further investigation.

The kinase activity of endogenous Lats1 (Fig. 3E) and overexpressed Lats2 (data not shown) is increased upon PKA activation, and the effect of PKA activation on YAP phosphorylation is blocked by a kinase-dead mutant Lats2, suggesting that the effect of PKA on YAP phosphorylation is mediated by Lats kinases. In contrast, MST1/2 are not required for PKA-induced YAP phosphorylation because the effect of forskolin and epinephrine on YAP phosphorylation is intact when both MST1 and MST2 are down-regulated by siRNA (Fig. 3B), and expression of kinase-dead mutant MST2 does not block the effect of forskolin on YAP phosphorylation (Fig. 3C). Consistently, MST1 kinase activity and phosphorylation at the activation loop are not modulated by forskolin treatment (Yu et al. 2012b). However, phosphorylation of the hydrophobic motif of Lats1 is induced by cAMP signaling (Yu et al. 2012b), indicating that a kinase other than MST may phosphorylate the hydrophobic motif of Lats kinases upon PKA activation. Although unlikely, we cannot exclude the possibility that a residual amount of MST kinase activity is sufficient to activate Lats in response to cAMP. Moreover, it is also possible that PKA may promote Lats phosphorylation by inhibiting a phosphatase.

PKA phosphorylates proteins containing the RRXS/T consensus sequence, and several components of the Hippo pathway with the RRXS/T motif might be direct targets of PKA. Neurofibromin 2 [NF2, also known as merlin], a tumor suppressor and an upstream component of the Hippo pathway [McCartney et al. 2000; Hamaratoglu et al. 2006; Benhamouche et al. 2010; Zhang et al. 2010], has been shown as a direct target of PKA [Alflthan et al. 2004]. Based on our data, NF2 is not critical for PKA to induce YAP phosphorylation because the MDA-MB-231 cells have a homozygous NF2 mutation (Dupont et al. 2011), while YAP phosphorylation is properly regulated by cAMP. PKA can also phosphorylate mouse Lats2 at S171 and S362 following forskolin treatment, with the S171 site conserved in Lats1 and warts (Drosophila Lats ortholog). However, mutation of S171 or S362 of mouse Lats2 cannot block forskolin-induced Lats2 activation (FX Yu and KL Guan, unpubl.), indicating that Lats1/2 are unlikely to be direct targets of PKA responsible for cAMP-induced YAP phosphorylation. Our data are consistent with a model in which Rho functions between PKA and Lats1/2 kinases (Fig. 7).

RhoA regulates the Hippo pathway by modulating the actin cytoskeleton. Formation of actin filaments or generation of cellular tension results in YAP dephosphorylation, nuclear localization, and activation [Dupont et al. 2011; Fernandez et al. 2011; Sansores-Garcia et al. 2011; Wada et al. 2011; Mo et al. 2012; Yu et al. 2012b; Zhao et al. 2012]. In addition to RhoA, other Rho family members, such as Rac and Cdc42, can also regulate the Hippo pathway kinases (Zhao et al. 2012). Therefore, the effect of PKA on the Hippo pathway may not be solely mediated by RhoA. Other Rho GTPases or their effectors may participate in the signaling pathway from PKA to Lats [Fig. 7]. In support of this, PKA has been shown to phosphorylate PAK [Howe and Juliano 2000], which in principle can lead to rearrangements of the actin cytoskeleton.

Figure 5. YAP/TAZ mediate the effect of cAMP in adipogenesis. (A) 3T3-L1 cells were treated with 10 μM forskolin or 100 μM IBMX for 1 h, or serum-starved 3T3-L1 cells were treated with 5 μM KT5720 for 1 h, and YAP phosphorylation and TAZ protein levels were determined. (B) 3T3-L1 cells were incubated under adipocyte differentiation conditions with IBMX or KT5720. (C) 3T3-L1 cells were transfected with control or YAP and TAZ siRNAs (siYT), and the knockdown efficiency was determined by immunoblotting [shown in C]. (D) These cells are also subjected to adipogenesis. (E) 3T3-L1 cells were transfected with control or YAP and TAZ siRNAs. Cells were treated with Tro and IBMX in the presence of vehicle (DMSO) or KT5720 as indicated. Adipocyte differentiation was measured by oil red staining. (F) Overexpression of YAP abolished IBMX- or forskolin-induced adipogenesis. (G) Following differentiation (as in F), cells were lysed, and the expression of adipogenesis marker genes was determined by real-time RT–PCR; the mRNA level was normalized to that of cells incubated in growth medium.
Ga12/13- and Gaq/11-mediated signaling activates Rho GTPases and YAP (Yu et al. 2012b). The importance of Rho GTPases in PKA-mediated YAP inactivation suggests that Ga-mediated signals antagonize with Ga12/13- and Gaq/11-mediated signals on the activity of Rho GTPases, which in turn results in induction or repression of YAP phosphorylation. Therefore, differential regulations of Rho GTPases by numerous extracellular molecules will fine-tune the activity of the Hippo pathway and determine cellular responses such as cell proliferation, apoptosis, and differentiation (Fig. 7). This also provides a mechanism of signal integration when cells have to respond to a wide range of extracellular signals.

YAP/TAZ inhibition mediates cellular functions of PKA

PKA is the first protein kinase purified and is involved in a wide range of physiological regulations [Taylor et al. 2012]. This study indicates that inhibition of YAP/TAZ contributes to the physiological function of cAMP or PKA. For example, cAMP can promote adipocyte differentiation, and this process is dependent on inhibition of YAP and TAZ [Fig. 5]. Besides adipocyte differentiation, PKA has been shown to induce neuronal differentiation and inhibit osteoblast differentiation (Ravni et al. 2006; Yang et al. 2008). One may speculate that YAP/TAZ play a role in neurogenesis or osteogenesis in response to cAMP signals. Consistent with this notion, YAP or TAZ has been shown to promote osteogenesis and inhibit neuronal differentiation [Hong et al. 2005; Cao et al. 2008; Zhang et al. 2012]. Interestingly, RhoA has functions similar to those of YAP/TAZ during various cell differentiation processes [McBeath et al. 2004]. Therefore, inhibition of Rho GTPases and YAP/TAZ may serve as a common mechanism in PKA-regulated cell differentiation.

PKA exerts a growth inhibitory effect on most cell and tissue types. YAP and TAZ are putative oncoproteins, and their activation stimulates cell proliferation and inhibits apoptosis. Therefore, PKA may inhibit cell growth by inactivating YAP/TAZ. This notion is supported by the functional analyses in Drosophila in which PKA inhibits the expression of cyclin E and Diap1. Based on the data presented in this study, we propose that inhibition of YAP/TAZ plays a key role in mediating the growth inhibitory effect of PKA. YAP/TAZ activation by either increased protein expression or reduced phosphorylation is associated with a large number of human cancers [Chan et al. 2008; Steinhardt et al. 2008]. Many pharmaceutical drugs directly target cellular cAMP levels. We speculate that elevation of cAMP by either PDE inhibitors or adenylyl cyclase activators may suppress tumor growth, particularly for those with high activity of YAP or TAZ.

Figure 6. PKA inhibits Yki in Drosophila. (A) In Drosophila S2R+ cells, knockdown of PKA-C1 by RNAi increased Yki/Sd reporter activity. (B) Relative transcript levels of ex, CycE, and Diap1 genes in wild-type (blue), C5-Gal4/UAS-PKA-C1 RNAi (red), and C5-Gal4/UAS-PKA-C1 (green) larval wing discs. (C) Yki phosphorylation was increased in C5-Gal4/UAS-PKA-C1 larval wing discs. (D-F) en-Gal4/UAS-PKA-C1 UAS-GFP larval wing discs exhibiting expression of GFP marker (D, green), Diap1 protein (E, red), and Caspase3 (F, white). (G) Merge of D–F.

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Figure 7. Regulation of the Hippo pathway by cAMP-PKA signaling. Upon stimulation of Gαs-coupled GPCR, activation of PKA by cAMP leads to inhibition of Rho GTPases, which indirectly inhibit Lats kinase activity. Stimulation of Gα12/13- or Gαq/11-coupled receptors antagonize the effect of cAMP or PKA on YAP phosphorylation by inducing Rho GTPases. Inhibition of YAP and TAZ mediates functions of cAMP and PKA on adipogenesis, cell proliferation, and apoptosis.
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Materials and methods

Cell culture

MDA-MB-231 cells were cultured in DMEM/F12 medium [Invitrogen]. HEK293A, HEK293T, U2OS, and MEFs were cultured in DMEM medium [Hyclone]. Primary hepatocytes were isolated from 12-wk-old male mice using a standard protocol and incubated in DMEM medium. All of the above cells were supplemented with 10% fetal bovine serum [FBS] [Omega Scientific] and 50 μg/mL penicillin/streptomycin [P/S]. MCF10A cells were cultured in DMEM/F12 supplemented with 5% horse serum [Invitrogen], 20 ng/mL EGF, 0.5 μg/mL hydrocortisone, 10 μg/mL insulin, 100 ng/mL chola toxin, and 50 μg/mL P/S. For serum starvation, cells were incubated in DMEM or DMEM/F12 without supplements. All cell lines were maintained at 37°C with 5% CO2.

Chemicals

Epinephrine, glucagon, dexamethasone, troglitazone, and rolipram were purchased from Sigma-Aldrich. IBMX, forskolin, KT5720, ibudilast, and theophylline were purchased from Tocris.

Transfection

Cells were transfected with plasmid DNA using PolyJet DNA in vitro transfection reagent (Signagen Laboratories) according to the manufacturer’s instructions. Dr. Mark Ginsberg [University of California at San Diego] generously provided the GFP-GDI plasmid. The pCMV-Flag-YAP, pCDNA3-MST2 K/R plasmids have been described elsewhere (Zhao et al. 2010, 2012). RhoA, the PKA catalytic subunit, and pCDNA3-Lats2 K/R plasmids were delivered into cells using RNAiMAX (Invitrogen) according to the manufacturer’s instructions. Dr. Mark Ginsberg (University of California at San Diego) generously provided the GFP-GDI plasmid. The pCMV-Flag-YAP, pCDNA3-MST2 K/R, and pCDNA3-Lats2 K/R plasmids have been described elsewhere (Zhao et al. 2010, 2012). RhoA, the PKA catalytic subunit, and Rap1b were in a pCDNA3 vector. The PKA regulatory subunit mutants [llα and llα] were in a pEGFP-C1 vector.

RNAi

Smartpool siRNAs were purchased from Dharmacon, and siRNAs were delivered into cells using RNAiMAX [Invitrogen] according to the manufacturer’s instructions. Lentiviral shRNAs in the pLKO.1 vector were purchased from Sigma-Aldrich, and virus was made in HEK293T cells using pMD2.g and PsPAX2 as packaging plasmids. Virus was filtered and used to infect targeting cells. The The RNAi Consortium (TRC) IDs for shRNAs used for the PKA catalytic subunit [PRKACA] are TRCN0000001372 and TRCN0000001373.

Immunoprecipitation and Lats kinase assay

Cells were lysed using mild lysis buffer (50 mM HEPES at pH 7.5, 150 mM NaCl, 1 mM EDTA, 1% NP-40, 10 mM pyrophosphate, 10 mM glycerophosphate, 50 mM NaF, 1.5 mM Na3VO4, protease inhibitor cocktail [Roche], 1 mM PMSF). Cell lysates were cleared and used for immunoprecipitation. The Lats1 antibody [Cell Signaling Technology] was mixed with cell lysates for 1 h at 4°C, and then protein A agarose beads were added in for 1 h. After four washes with lysis buffer, beads were washed once with wash buffer [40 mM HEPES, 200 mM NaCl] and once with kinase assay buffer [30 mM HEPES, 50 mM potassium acetate, 5 mM MgCl2]. The immunoprecipitated Lats1 was then subjected to a kinase assay in the presence of 500 μM cold ATP, 10 μCi [γ-32P]ATP, and 1 μg of GST-YAP expressed and purified from Escherichia coli as substrate. The reaction mixtures were incubated for 30 min at 30°C, terminated with SDS sample buffer, and subjected to SDS-PAGE and immunoblotting.

RNA extraction, reverse transcription, and real-time PCR

Following forskolin treatments or adipogenesis, cells were washed with cold phosphate-buffered saline [PBS] and subjected to RNA extraction using an RNasey Plus minikit [Qiagen]. RNA samples [1 μg] were reverse-transcribed to complementary DNA [cDNA] using iScript reverse transcriptase [Bio-Rad]. After dilution, cDNA levels were quantified by real-time PCR using KAPA SYBR FAST quantitative PCR master mix [Kapa Biosystems] and the 7300 real-time PCR system [Applied Biosystems]. The primer pairs [h and m indicate human and mouse, respectively] used in this study were β-actin [h], GGCAGACGAGGATGAGAAGGAG ATCA/AAGCTTGGCTGAGCAGTGGGA; CTGF [h], CAA TGACCAAGCTTCTCG/TGGTGACCAGAAAGTCCT; HPRT [m], GCAATACAGCCCCAATATGG/ACAAAGTGCCGGCTG TATCCAA/C/EBPb [m], GCAAAGCGCAAGAGCTGCTGGA/ CCTTCGTCTGGCAGCTCGACGT; PPARγ [m], CTGGCGGTGTT CAGAAAGTCCT/C/CCCAAACCTGATTGCGTGAAGACA; Adiposin [m], TCGCCCGCTGAACTCCTTACAATAATGTCGAC TACCCCGTCG; FABP4 [m], CGATGAAAATCCCGCGAC GAG/AGTACCGCCTTTTCATAACACA; and Adiponectin [m], AGATGGCACCCTCGTGGAGAGAA/ACATAAGCCTGGTCT CAGCGCT.

Adipogenesis

Murine 3T3-L1 cells were maintained in DMEM medium containing 10% calf serum [Hyclone]. To initiate adipocyte differentiation, confluent 3T3-L1 cells were switched into FBS-containing DMEM medium. In addition, insulin, dexamethasone, and troglitazone [Tro] were added. In selected samples, IBMX [250 μM] or forskolin [100 μM] was used to increase cAMP and PKA activity. For experiments using KT5720, 3T3-L1 cells were pretreated with KT5720 [5 μM] overnight in advance, and fresh KT5720 was added when adipogenesis was initiated. Two days later, the medium was changed into DMEM containing 10% FBS and insulin. After another 2 d, cells were incubated in DMEM with FBS. Cells were typically harvested on day 6 depending on the formation and maturation of lipid droplets. Cells were then subjected to RNA extraction or oil red [Sigma-Aldrich] staining according to the manufacturer’s instructions.

Luciferase assay

S2R+ cells were cultured in 24-well plates at standard conditions. Pka-C1 was down-regulated by dsRNAs. All the samples were cotransfected with 10 ng of the copia-Renilla luciferase reporter.
as a normalization control and 200 ng of 3xSd luc (gift from Dr. Jin Jiang) firefly luciferase reporter using Cellfectin II (Invitrogen). Fifty nanograms each of pUAST-Yki, pUAST-HA-sd, and pAc-Gal4 was used in each well to promote the firefly luciferase expression. Luciferase activity was measured after 48 h of incubation using the Dual-Glo luciferase assay kit (Promega) according to the manufacturer’s protocol.

**Drosophila stocks, quantitative RT–PCR, and immunocytochemistry**

All fruit flies were maintained under standard conditions. Fly stocks UAS-PKA-C1 [ID #55554] and UAS-PKA-C1-RNAi [ID #31277] were obtained from the Bloomington Drosophila Stock Center. Total mRNA from *Drosophila* late third-instar larval wing discs was isolated using the Qiagen RNAeasy kit (Qiagen), and mRNA was reverse-transcribed using Quanta qScriptDNAsuperMix (Qiagen). Real-time PCR was performed using PerfeCTa SYBR Green FastMix (Qiagen), and data were collected via the Applied Biosystem StepOnePlus real-time PCR system (Life Technologies). The relative amount of specific mRNAs under each condition was calculated after normalization to the *Histone 3* transcripts. A standard procedure for immunocytochemistry was followed in this study. Wing discs from late third-instar larvae were dissected in cold PBS and fixed in 4% PFA for 30 min at room temperature. Tissues were incubated in primary antibody overnight at 4°C, followed by 2 h of secondary antibody incubation at room temperature. Primary antibodies anti-Diap1 (1:200; Cell Signaling Technology) were used in this study. Images were collected with an Olympus Fluoview 1000 confocal laser-scanning microscope.

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