Inhibition of TBK1/IKKε Promotes Regeneration of Pancreatic β-cells

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β-cell proliferation induction is a promising therapeutic strategy to restore β-cell mass. By screening small molecules in a transgenic zebrafish model of type 1 diabetes, we identified inhibitors of non-canonical IκB kinases (IKKs), TANK-binding kinase 1 (TBK1) and IκB kinase ε (IKKε), as enhancers of β-cell regeneration. The most potent β-cell regeneration enhancer was a cinnamic acid derivative (E)-3-(3-phenylbenzo[c]isoxazol-5-yl)acrylic acid (PIAA, which, acting through the cAMP-dependent protein kinase A (PKA), stimulated β-cell-specific proliferation by increasing cyclic AMP (cAMP) levels and mechanistic target of rapamycin (mTOR) activity. A combination of PIAA and cilomostat, an inhibitor of β-cell-enriched cAMP hydrolyzing enzyme phosphodiesterase (PDE) 3, enhanced β-cell proliferation, whereas overexpression of PDE3 blunted the mitogenic effect of PIAA in zebrafish. PIAA augmented proliferation of INS-1β-cells and β-cells in mammalian islets including human islets with elevation in cAMP levels and insulin secretion. PIAA improved glycemic control in streptozotocin (STZ)-induced diabetic mice with increases in β-cell proliferation, β-cell area, and insulin content in the pancreas. Collectively, these data reveal an evolutionarily conserved and critical role of TBK1/IKKε suppression in expanding functional β-cell mass.

Inflammation to islets has emerged as a key contributor to the loss of functional β-cell mass in both type 1 diabetes (T1DM) and type 2 diabetes (T2DM)1,2. In T1DM, β-cells are the target of an autoimmune assault. Chronic low-grade inflammation and activation of the immune system are major factors in obesity-induced insulin resistance and T2DM. Therefore, immunotherapies designed to block β-cell apoptosis may stand as a unifying target for diabetes treatment. Despite this rationale, the slow rate of β-cell regeneration in adult humans3,4 limits the efficacy of immune-intervention trials. Accordingly, among multiple small mitogenic molecules identified5–18, several of them have either not shown or shown minor functional effects in human β-cells5–7. Moreover, some of them displayed off-target effects16,17,18. Thus, identifying β-cell regenerating agents that specifically increase residual functional β-cells and coupling them with immunomodulators represent an auspicious treatment for T1DM and T2DM9–9.

Non-canonical IκB kinases (IKKs), TANK-binding kinase 1 (TBK1) and IKKε, have high sequence homology with comparable phosphorylation profiling of substrate(s)18. These kinases regulate inflammatory reactions primarily through their action on the interferon regulatory factor (IRF) pathway21,22. Independent of their role in acute immune responses, TBK1 and IKKε were shown to be induced in response to obesity-dependent inflammation and directly phosphorylate phosphodiesterase (PDE) 3B23, a major cyclic AMP (cAMP) hydrolyzing PDE isoform in adipocytes24. Consequently, pharmacological inhibition of TBK1/IKKε with amlexanox, a small molecule inhibitor of these kinases, increased cAMP levels in adipocytes23. This led to the secretion of interleukin-6 (IL-6) and the activation of the hepatic Signal Transducer and Activator of Transcription 3 (STAT3)25, resulting in weight loss and reduced hepatic gluconeogenesis in obese mice26. In addition, IKKε was shown to be among putative targets of diarylamine WS6, a small molecule that promoted human β-cell proliferation in vitro27,28. Despite

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Intracellular cAMP levels modulated by their rate of synthesis via adenyl cyclase and their rate of degradation via PDEs are essential for β-cell replication, survival, and insulin secretion. G-protein coupled receptors (GPCRs) adenosine receptor 2a (A2a) and 2b (A2b) activate Gs and stimulate production of cAMP to increase in β-cell regeneration (Fig. 1D, I). Although TBK1/IKKε has demonstrated to blunt Ucp1 expression in response to βAR agonists in 3T3-L1 adipocytes, the signaling regulatory networks that link TBK1/IKKε, cAMP levels, and mTOR activity to proliferation and functional restoration of β-cells remain elusive. In this study, through chemical screens using the zebrafish model of type 1 diabetes, we identified TBK1/IKKε inhibitors (TBK1/IKKε-Is) as enhancers of β-cell regeneration. Pharmacological and genetic functional analyses in zebrafish using the most promising hit-compound (E)-3-(3-phenylbenzo[c]isoxazol-5-yl)acrylic acid (PIAA) indicated that suppression of TBK1/IKKε augments β-cell-specific proliferation by increasing cAMP levels and mTOR activity via PDE3. PIAA improved function and replication of mammalian β-cells including primary human β-cells. Furthermore, PIAA improved glucomycin control and induced β-cell proliferation with increase in insulin content in the pancreas in streptozocin (STZ)-induced diabetic mice.

Results

Chemical screens identify TBK1/IKKε inhibitors as enhancers of β-cell regeneration in zebrafish. To identify bioactive compounds that facilitate pancreatic β-cell regeneration, we screened a library of 75 small molecules with well-characterized biological and pharmaceutical activity in a transgenic zebrafish model of type 1 diabetes. We used the Tg(ins:CFP-NTR)468 line, in which β-cells are eradicated by nitroreductase (NTR), an enzyme that converts the chemical metronidazole (MTZ) to a DNA interstrand cross-linking agent. To easily follow the ablation and regeneration of β-cells, we used an additional transgenic line, Tg(ins:Kaede)486, which expresses the brightly green fluorescent protein Kaede in β-cells. Tg(ins:CFP-NTR)468; Tg(ins:Kaede)486 larvae were treated with MTZ at 3 days post-fertilization (dpf) for 24 hours to induce β-cell apoptosis, followed by washing out of MTZ (at 4 dpf, defined as 0 hours post-ablation (hpa)) and subsequent recovery in the presence or absence of chemical compounds for 48 hours (4–6 dpf, corresponding to 0–48 hpa). Using this system, we identified that the compound BX795 can approximately double the number of regenerated β-cells at 48 hpa (Fig. 1B, I). BX795 is a small molecule inhibitor that represses both 3-phosphoinositide-dependent kinase 1 (PDPK1) and eukaryotic translation initiation factor 4E (eIF4E)-binding protein 1 (4E-BP1) activity and increases in β-cell regeneration via PDE3. We screened a library of small molecules with well-characterized biological and pharmaceutical activity in a transgenic zebrafish model of type 1 diabetes. We used the Tg(ins:CFP-NTR)468; Tg(ins:Kaede)486 line, in which β-cells are eradicated by nitroreductase (NTR), an enzyme that converts the chemical metronidazole (MTZ) to a DNA interstrand cross-linking agent. To easily follow the ablation and regeneration of β-cells, we used an additional transgenic line, Tg(ins:Kaede)486, which expresses the brightly green fluorescent protein Kaede in β-cells. Tg(ins:CFP-NTR)468; Tg(ins:Kaede)486 larvae were treated with MTZ at 3 days post-fertilization (dpf) for 24 hours to induce β-cell apoptosis, followed by washing out of MTZ (at 4 dpf, defined as 0 hours post-ablation (hpa)) and subsequent recovery in the presence or absence of chemical compounds for 48 hours (4–6 dpf, corresponding to 0–48 hpa). Using this system, we identified that the compound BX795 can approximately double the number of regenerated β-cells at 48 hpa (Fig. 1B, I). BX795 is a small molecule inhibitor that represses both 3-phosphoinositide-dependent kinase 1 (PDPK1) and non-canonical IKKs, TBK1 and IKKε (Fig. 1C). To determine which pathway’s suppression is primarily responsible for augmenting β-cell regeneration, we tested the potent PDPK1 inhibitor BX912. BX912 caused minimal increase in β-cell regeneration (Fig. 1D, I). Hence, we further tested the following biologically known as well as uncharacterized TBK1/IKKε inhibitors (TBK1/IKKε-Is): amlexanox, azabenimidazol (AZ) derivatives 5c and 5e, and a cinnamic acid derivative (E)-3-(3-phenylbenzo[c]isoxazol-6-yl)acrylic acid (iso-PIAA; Figs 2A–C and S1A, B; see details in Supplemental Experimental Procedures). Each of these compounds augmented the number of regenerated β-cells, with PIAA showing the highest efficiency (Fig. 1E–I). PIAA blocked the activity of IKKε with a half maximal inhibitory concentration of (IC50) approximately 1.07 μM and that of TBK1 at 0.41 μM (Fig. 2A, B). It did not block IKKα or IKKβ at these concentrations (IC50 of PIAA for IKKα: 47.2 μM; IKKβ: 50.5 μM). In addition, PIAA downregulated polyinosine:polycytidylic acid (poly I:C)-stimulated phosphorylation of interferon responsive factor-3 (IRF3), a substrate of TBK1/IKKε, in INS-1 rat pancreatic β-cells (Figs 2F and S2). These results suggest that TBK1/IKKε inhibition, rather than PDPK1 repression, enhances β-cell regeneration.

To elucidate the interactions between TBK1/IKKε and inhibitors, we performed molecular docking simulations using the well-characterized TBK1 crystal structure in complex with BX795 (PDB entry 4EUT). We observed that amlexanox and PIAA adapt docking poses that closely mimic the crystallographically obtained structure of BX795 by forming multiple hydrogen bonds (H-bonds) with key residues within the kinase domain of TBK1 (Fig. 2E–G). The core aromatic moieties, the carboxylate of amlexanox and PIAA, and the thiophene amide of BX795 all bound within an unoccupied region of the kinase domain (Fig. 2E–G), while BX795, being a longer molecule, used its urea moiety to extend toward the outer rim of the kinase domain through interaction with the carbonyl group of Pro-90 (Fig. 2F, G). To corroborate the docking simulations, we further performed structure-activity relationship (SAR) studies of PIAA. We synthesized four analogs: PIAA-1 (an analog lacking the carboxylate group), PIAA-2 (a methyl ester analog of PIAA), PIAA-3 (an analog with an open isoxazolyl ring), and PIAA-4 (an amide analogue of PIAA) (Figs 2H and S1A). The carboxylate group and the intact isoxazolyl ring were both required in enhancing β-cell regeneration, since analogs lacking either group were inactive (Fig. 2I).
Thus, the SAR data and modeling of TBK1 and inhibitor interactions delineate the molecular basis of the selectivity of the TBK1/IKKe-I used in in vivo chemical screens. Taken together, these results indicate that suppression of TBK1/IKKe-I augments β-cell regeneration in the zebrafish model of type 1 diabetes.

Repression of TBK1/IKKe increases β-cell regeneration by primarily promoting their proliferation. To exclude a substantial contribution of pre-existing β-cells to regeneration of β-cells, we converted the fluorescence of the Kaede protein from green to red by exposing the [Tg(ins:CFP-NTR)] expression (green) of larvae at 48 hpa treated with DMSO (A), BX795 (B), BX912 (D), amlexanox (E), AZ-5c (F), AZ-5e (G), and PIAA (H), respectively. While TBK1/IKKe-I substantially expanded [Tg(ins:CFP-NTR)]-expressing cell population (white squares and insets) during regeneration (B and E–H) compared to DMSO (A), the PDPK1 inhibitor BX912 showed minimal effect (D). (C) BX795 is a dual inhibitor of PDPK1 and TBK1/IKKe. (I) Quantification of the number (mean ± SD) of total regenerated β-cells at 48 hpa (in A-B and D–H; 4.3 ± 1.3 (DMSO), 9.0 ± 2.2 (BX795), 4.7 ± 1.2 (BX912), 11.0 ± 3.4 (amlexanox), 7.4 ± 1.4 (AZ-5c), 7.0 ± 1.3 (AZ-5e), and 17.7 ± 2.4 (PIAA)). Cells in 20 planes of confocal images from 25 individual larvae were counted per condition. ***P < 0.001.
suppression of TBK1/IKKε-cell function.

β-cell ablation, showed that TBK1/IKKε-Is did not lead to an overshoot in β-cell number (Fig. S4E–J and data not shown). These results suggest that suppression of TBK1/IKKε enhances β-cell proliferation during the most dynamic period of β-cell regeneration without inducing a general increase in proliferation of other cell types.

Next, we examined the ability of TBK1/IKKε-Is to restore normoglycemia. Free glucose levels were elevated after β-cell ablation but declined from 24–72 hpa (corresponding to 5–7 dpf) in DMSO- and TBK1/IKKε-Is-treated larvae (Fig. 5I). Importantly, normal levels of free glucose were recovered significantly faster in TBK1/IKKε-Is-treated, especially PIAA-treated, larvae than in DMSO-treated larvae (Fig. 5I). Altogether, these data suggest that inhibition of TBK1/IKKε induces selective increase in β-cell number that accelerates restoration of sufficient overall β-cell function.

Repression of TBK1/IKKε enhances β-cell replication via cAMP-PKA-mTOR activation. To explore the mechanisms of how TBK1/IKKε suppression stimulates β-cell replication, we measured the regeneration efficiency of β-cells simultaneously treated with several β-cell replication-pathway inhibitors in the presence

Figure 2. Kinase profiling, molecular docking, and structure-activity relationship analyses reveal selectivity of the TBK1/IKKε inhibitors. (A, B) Chemical structures and kinase profiling of PIAA and iso-PIAA. Dose responses of PIAA and iso-PIAA were generated to determine the potency of the inhibitors (IC50). (C) Ball and stick model of PIAA (grey) and iso-PIAA (red) docked into the binding pocket of TBK1. PIAA adapts a docked pose that has its isoxazole nitrogen and carboxylate moieties engaged in stronger interactions, relative to the same moieties on iso-PIAA, with THR-156 and ASP-157 at the active site of TBK1. (D) Chemical structures of amlexanox and BX795. (E–G) Molecular docking simulations showing interactions of TBK1/IKKε-Is and TBK1. Ball and stick model of PIAA (grey), amlexanox (blue), and BX795 (purple) docked into the binding pocket of TBK1 (E and F). Space filling model of PIAA (grey) and BX795 (purple) docked into the binding pocket of TBK1 (G). The core moieties of PIAA, amlexanox, and BX795 all bound within an unvaried region of the kinase domain of TBK1, while the urea moiety of BX795 extends toward the outer rim of the TBK1 kinase domain and interacts with the carbonyl group of Pro-90. Specifically, the carboxylate moieties of PIAA and amlexanox are placed next to the carboxylate side chain of ASP-157 buried in the TBK1 active site, forming a strong low-barrier H-bonding between these two carboxylic acid groups. (H) Chemical structures of four PIAA analogs. The moieties that were replaced and different from the original PIAA structure are marked in red (PIAA-1, PIAA-2, and PIAA-4) or blue (PIAA-3). (I) Quantification of the number (mean ± SD) of total regenerated β-cells at 48 hpa treated with DMSO, PIAA-1, PIAA-2, PIAA-3, PIAA-4, and PIAA, respectively (4.8 ± 0.8 (DMSO), 4.2 ± 1.3 (PIAA-1), 6.2 ± 1.3 (PIAA-2), 4.0 ± 1.2 (PIAA-3), 6.0 ± 2.2 (PIAA-4), and 18.6 ± 3.4 (PIAA)). Cells in 20 planes of confocal images from 10 individual larvae were counted per condition. **P < 0.001. (J) Representative Western blot showing a PIAA dose-dependent decrease of pIRF3 levels in rat INS-1 cells.
of TBK1/IKKE–Is, specifically PIAA. Whereas the phosphoinositide 3-kinase (PI3K) inhibitor LY294002 and Akt/Protein Kinase B (PKB) inhibitor MK2206 caused statistically significant increases in β-cell regeneration when combined with PIAA (Fig. S5K, L and S5N), the mechanistic target of rapamycin (mTOR) inhibitor rapamycin and the cAMP-dependent protein kinase A (PKA) inhibitor PKI-(6–22)-amide both suppressed PIAA-mediated β-cell regeneration (Fig. S5I, J and S5N). We observed the same effect when each of LY294002 and MK2206 was tested alone (Fig. S5E, F and S5M). These results implicate that cAMP-PKA and mTOR signaling mediate the β-cell regeneration response to TBK1/IKKE–Is (Fig. S5O) and that PI3K-Akt signaling, cooperatively with TBK1/IKKE signaling, may play a repressive role in β-cell regeneration (Fig. S5O).

Given the previous studies showing PKA directly phosphorylating mTOR and RAPTOR in white adipose browning via βAR-cAMP-PKA-mTORC1 signaling pathway42 and TBK1 and IKKe direct phosphorylating and activating PDE3B in vitro35, and our replication-pathway inhibitor data (Fig. S5J and S5N), we hypothesized that TBK1/IKKe–Is induce the activity of the cAMP-PKA-mTOR for β-cell regeneration via suppression of the TBK1/IKKe–PDE3 signaling axis (Fig. 6A). To test this hypothesis, we first measured the cellular cAMP levels and determined the phosphorylation status of the S6 kinase 1 (S6K1), a downstream target of the mTOR signaling cascade43, in the PIAA-treated regenerating larvae. Treatment with PIAA led to pronounced increases in both cAMP levels and S6K1 protein phosphorylation (Figs 6B, C, S6A and S7). Furthermore, a combination of PIAA and PDE3 inhibitor cilostamide enhanced β-cell regeneration with increases in cAMP levels, number of β-cells that were EdU incorporated, and immunoreactive for phosphorylated ribosomal protein S6 (pRPS6), another downstream target of the mTOR signaling cascade43, compared to individual compound-treated larvae (Figs 6E–6G, S6B, and S8A–E). These effects were suppressed by rapamycin treatment (Figs 6F–6I, 6J, K, and data not shown). In a converse experiment, we assessed the effects of ectopic expression of pde3a on mitogenic potential of TBK1/IKKe–Is using a heat-inducible transgene Tg(hsp:pde3a; hsp:GFP)64. Pde3a is the only PDE3 isoform in zebrafish. When pde3a expression was induced during recovery period in the presence of PIAA, the proportion of new β-cells, which were EdU incorporated and pRPS6-positive, was decreased compared to PIAA-only-treated larvae (Fig. S8F–K). These data suggest that suppression of TBK1/IKKe bestows an increase in β-cell number by regulating cAMP and mTOR activity through PDE3 in the zebrafish model of type 1 diabetes (Fig. S8L).

**TBK1/IKKe inhibition augments β-cell function and proliferation in mammalian systems.** To determine whether the effects of TBK1/IKKe suppression on β-cells are conserved across species, we first performed glucose-stimulated insulin secretion (GSIS) assay in primary rat and human islets as elevation of cAMP levels has shown to lead to enhanced β-cell replication, survival, and insulin secretion10,28,29. PIAA treatment significantly increased glucose stimulation indices in rat and human islets (Fig. 7I, M). Next, we investigated the mitogenic effects of TBK1/IKKe inhibition by analyzing the ability of PIAA to increase β-cell proliferation in INS-1 cells66. Treating INS-1 cells with PIAA resulted in increased percentage of proliferating Insulin-positive cells (co-expressed Ki-67 (Fig. 7A–C and Supplementary Table 1) and levels of cAMP (Fig. 7D). Furthermore, PIAA...
augmented phosphorylation of PKA substrate and S6K1, RPS6, and Grb10 proteins, which are established targets of mTOR, as well as ERK1/2, whereas it triggered minimal phosphorylation of Akt (Figs 7E, S9A, B, and S10) in INS-1 and INS-1-derived 832/13 cells. These data suggest that suppression of TBK1/IKKε promotes replication of β-cells via activation of the cAMP-mTOR signaling axis in mammalian systems. PIAA also increased β-cell proliferation in both dispersed and whole rat islets in a dose-dependent manner (Fig. 7F–H, Supplementary Table 2, and data not shown), consistent with its mitogenic effect on β-cell formation even in the absence of injury in zebrafish (Fig. S11A–D). Importantly, treatment of PIAA on primary human β-cells using islets obtained from 3 cadaveric organ donors caused a notable, dose-dependent induction of β-cell proliferation (Fig. 7J–L and Supplementary Table 3).

We further investigated whether PIAA could increase β-cell regeneration in the streptozotocin (STZ)-induced mouse model of type 1 diabetes. PIAA administration started causing a substantial reduction of non-fasting blood glucose levels after 4–5 days of intraperitoneal injection (Fig. 8A). Significant improvement in glucose and insulin tolerance was observed compared with vehicle treatment (Fig. 8B, C). Morphometric analysis of pancreas sections showed that the β-cells, not α-cells, in PIAA-treated mice were more likely to be Ki67+, indicating that they were proliferating at a higher rate (Fig. S8D–F and Supplementary Table 4). Moreover, β-cell area and insulin content were increased in PIAA-treated compared with vehicle-treated diabetic mice (Fig. 8G, H). There was no difference in the weight of the mice based on treatment, neither at the start nor at the end of the experiments, indicating that the mice were not generally affected by PIAA treatment (data not shown). Taken together, these

Figure 4. TBK1/IKKε inhibitors have modest effects on α-to-β-cell transdifferentiation but strongly enhance β-cell proliferation. (A–C) Confocal images of [Tg(ins CFNTR)692; Tg(ins:Kaede)692] larvae at 24 hpa, concurrently treated with EdU and DMSO (A), amlexanox (B), or PIAA (C), respectively, from 0–24 hpa, stained for Somatostatin (blue). (D) Quantification of the number (mean ± SD) of Insulin and Somatostatin-double positive cells at 24 hpa in A–C: 0.7 ± 0.6 (DMSO), 1.7 ± 0.8 (amlexanox), and 1.8 ± 0.5 (PIAA). (E–G) Confocal images of [Tg(ins CFNTR)692; Tg(ins:Kaede)692] larvae at 24 hpa, concurrently treated with EdU and DMSO (E), amlexanox (F), or PIAA (G), respectively, from 0–24 hpa, stained for Glucagon (blue). Note that the number of Insulin and Glucagon-double positive cells (blue arrows) was increased in TBK1/IKKε-I-treated recovering larvae (F, G) compared to DMSO-treated larvae (E). PIAA-treated larvae also showed an EdU-incorporated β-cell (white arrow) (G). (H) Quantification of the number (mean ± SD) of Insulin and Glucagon-double positive cells at 24 hpa in E–G: 1.5 ± 1.1 (DMSO), 3.4 ± 1.3 (amlexanox), and 4.4 ± 1.9 (PIAA). (I–K) Confocal images of [Tg(ins CFNTR)692; Tg(ins:Kaede)692] larvae at 48 hpa, concurrently treated with EdU and DMSO (I), amlexanox (J), or PIAA (K), respectively, from 24–48 hpa. The number of β-cells that incorporated EdU (white arrows) was significantly increased in TBK1/IKKε-I-treated recovering larvae (J, K) compared to DMSO-treated larvae (I). (L) The percentage (mean ± SD) of regenerated β-cells that incorporated EdU at 48 hpa: 4.0 ± 7.0% (DMSO), 32.0 ± 6.0% (amlexanox), and 39.0 ± 6.0% (PIAA). Cells in 20 planes of confocal images from 10 individual larvae were counted per condition. **P < 0.01.
data suggest that inhibition of TBK1/IKKe leads to improvement of β-cell function and induction of β-cell replication across multiple species including primary human β-cells and diabetic mice.

**Discussion**

In this study, we identified TBK1/IKKe-I s as selective enhancers of β-cell regeneration in a transgenic zebrafish model of type 1 diabetes. We further demonstrated that inhibition of TBK1/IKKe promotes amplification of β-cells in mammalian systems including primary rat and human islets as well as STZ-induced diabetic mice. The proliferative effects of TBK1/IKKe-I s are likely to be mediated by the cAMP-PKA-mTOR signaling axis via PDE3, indicating that TBK1/IKKe play a previously unappreciated role in modulating β-cell mass.

Utilizing small molecule inducers of β-cell proliferation is one of the most tangible methods to restore functional β-cell mass. We have pinpointed that inhibition of TBK1/IKKe promotes β-cell proliferation in multiple
species including primary human β-cells. Of essential importance is that suppression of TBK1/IKKε using amlexanox and PIAA, which exhibited the highest potency among tested, can increase β-cell proliferation selectively without inducing a general increase in proliferation of other cell types/tissues. Furthermore, a longer treatment of β-cell ablated zebrafish with these small molecules did not lead to over-proliferation of β-cells once normoglycemia was approached. Since oncogenicity can arise as a result of modulating mitogenic or regenerative pathways⁸, specific TBK1/IKKε-Is' selectivity to β-cells and ability to increase β-cell proliferation primarily during the most active period of β-cell regeneration present valid strategies for expanding β-cell mass. Furthermore, PIAA exhibited substantial efficiency in β-cell regeneration with minimum toxicity. Biologically less characterized azabenzimidazole (AZ) derivatives 5c and 5e have IC50 of 0.032 μM of PIAA and cilostamide, 6.0 ± 2.0, of which 1.0 ± 1.0 (PIAA and rapamycin), and 5.2 ± 1.1, of which 0.8 ± 0.8 (cilostamide and rapamycin) incorporated EdU with pRPS6 immunoreactivity (E-E'). (K) The percentage (mean ± SD) of regenerated β-cells that incorporated EdU with pRPS6 immunoreactivity at 48 hpa (in D-I'; 6.1 ± 9.5% (DMSO), 47.3 ± 7.5% (PIAA), 50.4 ± 8.8% (cilostamide), 54.2 ± 4.2% (PIAA and cilostamide), 13.9 ± 12.7% (PIAA and rapamycin), and 13.7 ± 13.0% (cilostamide and rapamycin)). Cells in 20 planes of confocal images from 10 individual larvae were counted per condition. ***P < 0.001.

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Figure 6. Suppression of the TBK1/IKKε-PDE3 signaling axis promotes β-cell proliferation by increasing cAMP levels and mTOR activity. (A) Schematic of the TBK1/IKKε-PDE3 signaling that modulates cAMP-PKA-mTOR pathway. The sites of inhibition by PIAA and cilostamide are shown in red. (B) Quantification of cAMP levels (mean ± SD) at 48 hpa (0.4 ± 0.1 pmol/larva (DMSO) and 0.0 ± 0.0 pmol/larva (PIAA)). (C) Representative Western blot showing increased pS6K1 levels in PIAA-treated recovering larvae. (D-I) Confocal images of Tg(ins:CFP-NTR) larvae at 48 hpa, concurrently treated with EdU and DMSO (D-I'), PIAA (E-E'), cilostamide (F-F'), a combination of PIAA and cilostamide (G-G'), a combination of PIAA and rapamycin (H-H'), or a combination of cilostamide and rapamycin (I-I'), respectively, from 0–48 hpa, stained for pRPS6 (blue). The number of EdU-incorporated (white arrows) and pRPS6-positive (white arrowheads) β-cells was increased in recovering larvae treated with both PIAA and cilostamide (G-G') compared to individual compound-treated larvae (E-E' and F-F'). Rapamycin substantially suppressed the PIAA- and cilostamide-dependent increases in the number of EdU-incorporated and pRPS6-positive β-cells (H-H'). (J) Quantification of the number (mean ± SD) of total regenerated β-cells (green bars) and regenerated β-cells that incorporated EdU with pRPS6 immureactivity (white bars) at 48 hpa (in D-I'; 5.0 ± 1.3 total regenerated β-cells, of which 0.3 ± 0.5 (DMSO), 17.8 ± 2.8, of which 8.4 ± 1.7 (PIAA), 12.7 ± 1.9, of which 5.0 ± 1.5 (cilostamide), 24.7 ± 1.2, of which 13.3 ± 0.6 (PIAA and cilostamide), 6.0 ± 2.0, of which 1.0 ± 1.0 (PIAA and rapamycin), and 5.2 ± 1.1, of which 0.8 ± 0.8 (cilostamide and rapamycin) incorporated EdU with pRPS6 immunoreactivity). (K) The percentage (mean ± SD) of regenerated β-cells that incorporated EdU with pRPS6 immunoreactivity at 48 hpa (in D-I'; 6.1 ± 9.5% (DMSO), 47.3 ± 7.5% (PIAA), 50.4 ± 8.8% (cilostamide), 54.2 ± 4.2% (PIAA and cilostamide), 13.9 ± 12.7% (PIAA and rapamycin), and 13.7 ± 13.0% (cilostamide and rapamycin)). Cells in 20 planes of confocal images from 10 individual larvae were counted per condition. ***P < 0.001.

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Previous studies have suggested that modulation of cAMP levels via GPCR in β-cells is essential for β-cell replication, survival, and insulin secretion. Our results provide compelling evidence that inhibition of TBK1/IKKε enhances selective β-cell proliferation by increasing cAMP levels via PDE3. Time-course analysis in zebrafish treating PDE3 inhibitor cilostamide, PIAA, and a combination of PIAA and cilostamide demonstrated the correlation between the levels of cAMP and expansion of β-cell mass: the most dynamic period of...
Figure 7. PIAA induces proliferation of rat and human β-cells. (A–B') Confocal images of rat INS-1 β-cells treated with DMSO (A–A') and PIAA (B–B’), respectively, stained for Ki67 (red) and Insulin (green). (C) The percentage (mean ± SD) of Ki67 and Insulin-double positive cells (in A–B'; 43.1 ± 5.2% (DMSO) and 87.3 ± 10.4% (PIAA)). (D) Quantification of cAMP levels (mean ± SD) (5.8 ± 0.2 pmol/well (DMSO) and 8.7 ± 0.4 pmol/well (PIAA)). (E) Representative Western blot showing increased phosphorylation of PKA substrate and ERK1/2 in PIAA-treated INS-1-derived 832/13 β-cells. Note that PIAA-treatment augmented phosphorylation of mTOR targets RPS6 and Grb10 but did not trigger phosphorylation of AKT^{T308} and AKT^{S473}. (F–G') Confocal single-plane images of whole rat islets treated with DMSO (F–F') and PIAA (G–G'), respectively, stained for Ki67 (red, white arrows) and Insulin (green). (H) The percentage (mean ± SD) of Ki67 and Insulin-double positive cells in whole rat islets increased in a dose-dependent manner with treatment of PIAA (1.1 ± 0.3% (DMSO), 2.7 ± 0.9% (20 µM), 3.7 ± 1.5% (40 µM), and 5.5 ± 0.6% (80 µM)). n = 5 replicates per condition. (I) Glucose stimulation indices of rat islets treated with DMSO or PIAA (300 islet equivalents per column, triplicate). (J–K') Confocal single-plane images of human islets treated with DMSO (J–J') and PIAA (K–K'), respectively, stained for Ki67 (red, white arrows), Topro (blue), and INSULIN (green). (L) The percentage (mean ± SD) of Ki67 and Insulin-double
β-cell regeneration corresponds to high levels of cAMP, whereas the period reaching to normoglycemia without over-proliferation of β-cells links to low levels of cAMP. Moreover, PIAA treatment of INS-1 rat β-cells resulted in increased percentage of proliferating Insulin-positive cells (co-expressed Ki-67) with concurrent augmentation of cAMP levels. It is likely that PKA stimulation is responsible for linking PIAA-enhanced cAMP levels and subsequent β-cell proliferation. In INS-1 and INS-1-derived 832/13 cells, PIAA increased phosphorylation of PKA substrate and ERK1/2, which was shown to be driven β-cell proliferation via cAMP-PKA signaling cascade upon GIP/GIPR stimulation. Phosphorylation of mTOR targets including S6K1, RPS6, and Grb10, was also greatly augmented with PIAA treatment. Both mammalian and zebrafish mTOR proteins contain 3 conserved PKA target RR/KXS motifs and RAPTOR proteins have 1 RRX motif (IX and CHS, unpublished observation) that were directly phosphorylated by PKA upon treatment with βAR agonists in vitro. PIAA did not trigger phosphorylation of Akt1 and Akt2, suggesting that the effect of TBK1/IKKe repression is at least in part through a cAMP-PKA-mTOR signaling, not via a well-established Akt-mTOR signaling. The suggested suppressive effect of PI3K-Akt signaling in β-cell regeneration in our studies is consistent with a report showing the inhibitory role of PI3K signaling in β-cell formation in zebrafish and may act through PDE3. Akt has been demonstrated to phosphorylate PDE3B in vitro and P3K has shown to directly activate Akt1 and stimulate Akt2 in a PDK1-dependent manner. Consistently, treatment of ciliostamide with PI3K and Akt inhibitors led to increase in β-cell regeneration in zebrafish (data not shown). In this regard, PDK1 triggering activation of both Akt and S6K1 may be the reason for minimal increase in β-cell number with PDK1 inhibition. While oxo-AR agonists described above stimulate cAMP-PKA-mTOR, and several PDE inhibitors including a PDE3 inhibitor cilostamide have displayed their potency to stimulate β-cell replication in a cAMP-dependent manner, enhanced insulin secretion was observed in Pde3b knockout (KO) mice. However, Pde3b KO mice fail to suppress hepatic glucose production and display insulin resistance with a number of cAMP-signal transduction components being altered in Pde3b-deficient livers. Contrarily, genetic deletion of IKKe and pharmacological inhibition of TBK1/IKKe led to improved insulin sensitivity through the inhibition of hepatic glucose production with decrease in PDE3B activity and increase in cAMP levels in adipocytes, not in livers, in obese mice. Thus, despite the ameliorated insulin tolerance in PIAA-treated STZ diabetic mice may be a secondary effect of hyperglycemic reversal by augmenting β-cell mass, our findings of modulation of PDE3 activity and cAMP levels via suppression of TBK1/IKKe is likely to enable us to pinpoint bona fide therapeutic approaches that increase the number of functionally adequate β-cells with direct or indirect improvement of insulin sensitivity.

In progression of T1 and T2DM, decreasing β-cell mass by cytokine- and/or glucolipotoxicity-induced apoptosis is a common feature. Thus, prevention of β-cell loss can be an alternative approach for increasing β-cell mass in diabetes. Considering our studies demonstrating that repression of TBK1/IKKe increases cAMP levels and the previous studies showing that anti-apoptotic gene bcl2 is induced by the cAMP-PKA-cAMP response element binding protein (CREB) signaling axis, it is plausible to speculate that suppression of TBK1/IKKe can preserve β-cell mass. Intriguingly, adipose-specific genetic ablation of TBK1 attenuates diet-induced obesity with exaggeration in glucose intolerance/insulin resistance, while genetic deletion of IKKe increases energy expenditure with improvement in insulin sensitivity on a high fat diet. Thus, a careful dissection and elucidation of TBK1- and/or IKKe-controlled signaling networks will shed light on modulating β-cell survival with concomitant increase in functional β-cell mass, opening up new avenues of therapies for mitigating diabetes.

Experimental Procedures

Zebrafish strains. Adult fish and embryos/larvae were raised and maintained under standard laboratory conditions. We used the following published transgenic lines: Tg(ins:CFP-NTR) and Tg(ins:Kaede). Zebrafish studies conducted and protocols used were approved by the Institutional Animal Care and Use Committees of Mayo Clinic and Georgia Institute of Technology, and were in accordance with National Institutes of Health guidelines.

β-cell ablation, chemical treatment, and photoconversion. To ablate β-cells, Tg(ins:CFP-NTR) or Tg(ins:Kaede) larvae were treated with freshly prepared 5 mM metronidazole (MTZ) (Sigma) from 3 dpf to 4 dpf in the dark, followed by washing out the MTZ, and subsequent 48 hours recovery in the presence of DMEM, individual, or combination of the chemical compounds. The names and the concentrations of the chemical compounds are described in the Supplemental Experimental Procedures. To demonstrate that all β-cells are ablated by MTZ treatment, Tg(ins:Kaede), expressing β-cells were converted from green to red by exposing them to UV light at 3 dpf immediately after MTZ treatment.

Chemical screening. To perform a chemical screen for compounds enhancing β-cell regeneration, we tested 75 compounds from the Stem Cell Signaling Compound Library (Selleckchem). Further details are described in the Supplemental Experimental Procedures.

Immunohistochemistry. Immunohistochemistry on whole-mount zebrafish larvae and 5-ethyl-2′-deoxyuridine (EdU) analysis were performed as previously described. For mammalian islet culture, after chemical treatment, islets were washed with PBS and fixed in 4% paraformaldehyde (PFA) for 1 hour.
For mouse studies, 8 μm thick sections were obtained by using a cryostat microtome (CryoStar NX70 Cryostat). See Supplemental Experimental Procedures for additional information.

**Glucose and cAMP measurements.** Glucose measurements were performed 3 times on 10 zebrafish larvae per condition using a fluorescence-based enzymatic detection kit (Biovision Inc.). cAMP content of whole-zebrafish larvae and rat INS-1 cells was analyzed using commercial ELISA kits (Enzo Life Sciences, Inc.). Further details are described in the Supplemental Experimental Procedures.

**Mammalian ex vivo islet culture and mice experiments.** Male Lewis rat pancreatic donors were purchased from Charles River (Wilmington, MA).

Human islets from healthy donors were purchased from Prodo Laboratories (Irvine, CA). 6–8-wk old C57BL/6 male mice (Jackson Laboratory) were used. Studies conducted and protocols used were approved by the Institutional Animal Care and Use Committee of Mayo Clinic and Georgia Institute of Technology and were in accordance with National Institutes of Health guidelines. See Supplemental Experimental Procedures for additional information.

**Statistical analysis.** All statistical analyses were performed using GraphPad Prism (version 7). P-values less than 0.05 were considered statistically significant.

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Author Contributions

C.H.S. conceived and designed the experiments. J.X., J.-Y.F., J.D.W., D.J.P., J.N., and C.H.S. performed the experiments. S.T., I.O.R., and A.K.O. provided reagents/materials and performed molecular docking simulations. J.X., J.-Y.F., J.D.W., A.J.G., D.-S.C., A.V.M., A.K.O., and C.H.S. analyzed the data. A.J.G., D.-S.C., A.V.M., A.K.O., and C.H.S. wrote the paper.

Additional Information

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