## SUPPLEMENTAL DATA

Functional fibroblast growth factor receptor-1 is expressed in murine βTC3 cells. FGFR1 expression has been observed previously in adult mouse islets (1, 2). To elucidate endogenous islet  $\beta$ -cell FGFR1 expression and activity, we examined the murine insulin-secreting βTC3 cell By RT-PCR, we confirmed line (Fig. S1). expression of the FGFR1IIIc mRNA transcript that was not detectable in the  $\alpha TC6$  cell line (Fig. S1A). FGFR1IIIc plasmid DNA was used as a positive control. Full-length endogenous receptor was also identified by Western immunoblotting as a protein band with a molecular mass of ~150 kDa (Fig. S1B, top). This band exhibited slight retardation in protein mobility with FGF-1 stimulation, consistent with tyrosine phosphorylation of activated receptor. Phosphorylation of the FRS2 docking protein (~90 kDa molecular mass) was also enhanced with FGF stimulation confirming activation of receptor (Fig. S1B; middle). Taken together, these data are consistent with endogenous FGFR1 expression and activity in the βTC3 cell line, providing a translational model system for islet studies.

Generation of functional fluorescent proteintagged FGFR1 isoforms. To measure FGFR1dependent ERK1/2 activity in islet  $\beta$ -cells and to generate tools for examining the molecular mechanisms of FGFR activation, we created Venus fluorescent protein-tagged constructs of full-length  $(R1\beta v)$ and kinase-deficient (KDR1\betav) FGFR1c. Construct expression was driven by the rat insulin promoter (3). By using lentiviral infection (MOI ~5), we achieved 80-90% expression efficiency in βTC3 cells (Fig. S2A). Western immunoblotting of transfected cell lysates was performed using antibodies directed against Venus fluorescent protein ( $\alpha$ GFP) and the C-terminus of FGFR1 (Flg (C-15)) to confirm and characterize construct expression (Fig. S2B). The full length receptor isoform R1<sub>b</sub>v was identified as a protein band at the expected molecular mass of approximately 150 kDa by both antibodies (Fig. S2B; solid arrow). KDR1v was detected as a protein band at the expected molecular mass of 80-90 kDa by

only the  $\alpha$ GFP antibody since this isoform is deficient for the C-terminal tail recognized by the Flg receptor antibody (Fig. S2B; asterisk). Protein bands of approximately 28-30 kDa in lysates of all transfected cells reflect expression of Venus indicating some enzymatic cleavage of the fluorescent protein in samples expressing receptor constructs (Fig. S2B; arrowhead). To determine membrane association of the receptor we subsequently constructs. measured fluorescent protein diffusion using live cell fluorescence recovery after photobleaching (FRAP) (Fig. S2C). Significantly slower rates of recovery were observed for the FGFR1 constructs ( $t_{1/2} \sim 12$  s for R1 $\beta$ v and KDR1 $\beta$ v) than the Venus construct ( $t_{1/2} \sim 0.7$  s), consistent with association at the plasma membrane. These data indicate that the fluorescent protein-tagged receptor isoforms are expressed appropriately in βTC3 cells and translocate to the plasma membrane.

Rat insulin promoter driven expression of R1<sub>βv</sub>, KDR1<sub>βv</sub> and Venus. Although islets are comprised of a majority of  $\beta$ -cells (~85 %), a number of other endocrine cells are present including  $\alpha$ -,  $\delta$ -, pancreatic polypeptide-, and ghrelin-cells. To confirm that the constructs driven by the rat insulin promoter were expressed exclusively in  $\beta$ -cells, we examined fluorescence protein expression in islets immunostained for insulin (Fig. S3A) and glucagon/somatostatin (non-ß cell; Fig. S3B). co-localization of Venus examining By fluorescent protein and insulin, it was evident that the constructs were expressed in insulin positive  $\beta$ -cells only (Fig. S3A). However, while all R1<sub>βv</sub>-, KDR1<sub>βv</sub>- and Venus-infected cells were positive for insulin, not all insulinpositive cells were positive for Venus suggesting <100% fluorescent protein transfection efficiency. By examining Venus fluorescence in association with glucagon- and somatostatin-associated immunofluorescence. we confirmed that the Venus-tagged constructs were not expressed in non- $\beta$ -type cells. These data demonstrate that the rat insulin promoter confines FGFR construct expression exclusively in  $\beta$ -cells in intact islets.

## SUPPLEMENTAL FIGURE CAPTIONS

**Figure S1. Murine \betaTC3 cells express functional FGFR1.** (A) RT-PCR of total RNA using primers designed to amplify FGFR1IIIc (*top*) and GAPDH (*bottom*) in  $\beta$ TC3 cells ( $\beta$ ),  $\alpha$ TC6 cells ( $\alpha$ ), water negative control (-), and R1 $\beta$ v plasmid DNA positive control (+). Sense (5'-CAA CCT GCC TTA TGT CCA GA-3') and antisense (5'-GCT TCC AGA ACG GTC AAC CA-3') primers were used [35 cycles of 94 °C for 1 min, 55 °C for 1 min, and 72 °C for 1 min]. The housekeeping transcript glyceraldehyde-3-phosphate dehydrogenase (GAPDH) was a positive control of RNA expression and was amplified over 35 cycles using sense (5'-ATC GAG CTC ATC CCA TCA CCA TCT TCC AGG-3') and antisense (5'-ACA TCT AGA GCC ATC ACG CCA CAG TTT CCC-3') primers. (B) Western immunoblotting of whole cell lysate (10 µg/lane) from cultures stimulated with FGF-1 (10 ng/ml; times as indicated) using FGFR1 antibodies recognizing the N-terminus (*top*), phospho-FRS2 (*middle*), and  $\beta$ -actin (loading control, *bottom*).

Figure S2. Expression and membrane association of Venus-tagged full-length (R1 $\beta$ v) and kinasedeficient (KDR1 $\beta$ v) FGFR1c in  $\beta$ TC3 cells. (A) Live cell images of Venus-, R1 $\beta$ v-, and KDR1 $\beta$ vassociated fluorescence in  $\beta$ TC3 cells. Receptor isoforms are excluded from the nucleus compared to Venus control protein. (B) Western immunoblotting of whole cell lysates (10 µg/lane) from cells expressing Venus, R1 $\beta$ v, KDR1 $\beta$ v or wild type  $\beta$ TC3 (WT) using antibodies recognizing either Venus (left panel) or the C-terminus of FGFR1 (Flg, C-15) (C) Fluorescence recovery after photobleaching (FRAP) curves revealing protein diffusion rates in cells expressing Venus (- $\bullet$ -), R1 $\beta$ v (- $\circ$ -), and KDR1 $\beta$ v (- $\blacktriangle$ -). Data are expressed as the average fluorescence recovery ( $\Delta$ F/F<sub>0</sub>)  $\pm$  standard error of the mean for three independent experiments.

Figure S3. Insulin-positive islet  $\beta$ -cells express fluorescent protein constructs driven by the rat insulin promoter. (A) Islets were infected with a low MOI of Venus, full-length R1 (R1 $\beta$ v), and kinase-deficient R1 (KDR1 $\beta$ v) and immunostained with anti-insulin antibody. Fluorescence images for Venus fluorescence protein (FP) and insulin were acquired and overlaid (FP/insulin). (B) Non-insulin producing cells were identified in a control islet expressing Venus protein by co-labeling for non- $\beta$ -cells (glucagon and somatostatin). The overlay image (Venus/Non- $\beta$ -cell) demonstrates that Venus-expressing cells do not express glucagon or somatostatin.

## REFERENCES

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