

**HIF2 α is an essential molecular brake for postprandial hepatic glucagon response
independent of insulin signaling**

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Supplement Table 1 (Related to Experimental procedure- Real-time PCR)

Primers	Sequence
G6pase F	GTGTCCAGGACCCACCAATA
G6pase R	ACTGTGGGCATCAATCTCCT
Pepck F	CTGGATGAAGTTTGATGCC
Pepck R	TGTCTTCACTGAGGTGCCAG
Epo F	CATCTGCGACAGTCGAGTTCTG
Epo R	CACAACCCATCGTGACATTTTC
Hif-2 α F	TGAGTTGGCTCATGAGTTGC
Hif-2 α R	TATGTGTCCGAAGGAAGCTG
Irs-1 F	GGAGGATTTGCTGAGGTCAT
Irs-1 R	CTATGCCAGCATCAGCTTCC
Irs-2 F	CACAATTCCAAGCGCCACAA
Irs-2 R	CATCACCTCCTCCCAGGGTA
Pgc-1 F	TGAGGACCGCTAGCAAGTTT
Pgc-1 R	TGTAGCGACCAATCGGAAAT
Pkar1 α F	CCCTCGAGTCAGTACGGATG
Pkar1 α R	GCATTCCTTCGGGAATACTTT
Pkar2 α F	CCTCCTCTTCTTCATCAGGG
Pkar2 α R	GAGTGACTCGGACTCGGAAG
Pkar2 β F	GATCATCGGTTTTGGGATGT
Pkar2 β R	ATAAACCGGTTCAAGGCG
Pkac α F	ATTCTGAGAAGGGGTCTCCC
Pkac α R	AAGAAGGGCAGCGAGCAG
Pkac β F	TCCTCAAGCCCAGCATTACT
Pkac β R	CAAGAAAGGCAGCGAAGTG
PDE1B F	TGCAGTAGAGTATGTGGGGC
PDE1B R	AGAAACCCAAGTTCCGAAGC
PDE2A F	CCCCATCACTCCTCATCATC
PDE2A R	AGAACCAGCCATTGCTCTTG
PDE3A F	CGAGGACCAAGGAAGAGATTC
PDE3A R	ATGAGCTGCTCCCTGGGTAT
PDE3B F	ATCGCCTCTTGGTCTGCCAG
PDE3B R	GCCTTCTGTCCATCTCAAATGTAGG
PDE4A F	CAGAACCTCAGCAAGCGCCA
PDE4A R	CCTGAGGACCTGGATACGG
PDE4C F	CGGACTGAGCGTCTAGGAGA
PDE4C R	GGGACACAGAGAAACCTTGG
PDE4D F	TGCTCAGGTCTTGGCTAGCCTGC
PDE4D R	TCCTCCAGGGTCTCGCTGGC
PDE7A F	GTGTTAATCAGCCGTTTCTTATT
PDE7A R	CAAAGCACCTATCTGAGCCT
PDE7B F	TCAATGAATGGGTAGGAGCC
PDE7B R	GAGCCCTGAACAGAGTGTCA
PDE10A F	AAATTATATGGTTTGACGGATG
PDE10A R	ACAGTCTCTGCACTAACAC

Supplement figure legends

Supplement Figure 1 *Progressive improvement in glucose and glucagon tolerance is independent of IRS-2 levels at 1-week following VHL disruption (Related to Figure 1).*

(A) Glucose tolerance test in mice at two week following disruption of VHL in liver (Vhl^{LivKO}). * $p < 0.05$ compared to $Vhl^{F/F}$. Five mice were assessed per group. (B) qPCR analysis of *Irs-1* and *Irs-2* in the livers of $Vhl^{F/F}$ and Vhl^{LivKO} at one and two week following disruption of VHL in liver (Vhl^{LivKO}). Expression was normalized to β -actin mRNA. Each bar represents the mean value \pm S.E.M. * $p < 0.05$ compared to $Vhl^{F/F}$. (C) Western blot analysis of IRS-2 in primary hepatocytes extracted at 1-week following VHL disruption or (D) in livers at 7-, 14- and 30-days following VHL disruption. Three mice assessed per group. (E) Western blot analysis of pFOXO1 and FOXO1 in cytosolic and nuclear fraction from Vhl^{LivKO} at 1-week following VHL disruption. Three mice assessed per group. (F) Glucagon tolerance test in mice at two week following disruption of VHL in liver (Vhl^{LivKO}). Four to five mice were assessed per group. * $p < 0.001$ compared to $Vhl^{F/F}$. Four to five mice were assessed per group.

Supplement Figure 2 *Decreased gluconeogenesis in Vhl^{LivKO} mice could not be restored by PGC-1 overexpression or STAT-3 inhibition (Related to Figure 2).* (A) Pyruvate tolerance test in mice at two weeks following disruption of VHL in liver (Vhl^{LivKO}). * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ compared to $Vhl^{F/F}$. Four to five mice were assessed per group. (B) Western blot analysis of insulin stimulated AKT phosphorylation in primary hepatocytes isolated from $Vhl^{F/F}$ and Vhl^{LivKO} mice that were pretreated for 1-hour with 50 nM LY294002 or 50 nM MK-2206. (C) qPCR analysis of PGC-1 in the primary

hepatocytes of *Vhl*^{F/F} and *Vhl*^{LivKO} mice (D) *G6pase* in the primary hepatocytes of *Vhl*^{F/F} and *Vhl*^{LivKO} mice infected with recombinant adenovirus expressing PGC-1 α (Ad-PGC-1 α). (E) Western blot analysis for pSTAT3 and STAT3 in primary hepatocytes of *Vhl*^{F/F} and *Vhl*^{LivKO} mice. (F) Western blot analysis for pSTAT3 and STAT3 in primary hepatocytes of *Vhl*^{LivKO} mice treated with the STAT3 inhibitor (STAT3i), 100 μ M S3i-201 (Apexbio, Houston, TX) for 2-hours. (G) qPCR analysis of *G6pase* and *Pepck* in the primary hepatocytes of *Vhl*^{F/F} and *Vhl*^{LivKO} treated with STAT3i for 2-hours. Gene expression studies are done in triplicates and expression was normalized to β -actin mRNA. Each bar represents the mean value \pm S.E.M. ***p < 0.001 compared to *Vhl*^{F/F}. ###p < 0.05 compared to untreated.

Supplement Figure 3 *Improved glucose homeostasis and glucagon resistance in *Vhl*^{LivKO} mice is HIF-2 α dependent (Related to Figure 3)*. (A) Fed and fasting glucose monitored in WT (*Vhl/Hif-1 α* ^{F/F}) and mice with compound disruption of VHL in liver (*Vhl/Hif-1 α* ^{LivKO}). (B) Glucose, (C) glucagon and (D) pyruvate tolerance test performed in WT and *Vhl/Hif-1 α* ^{LivKO} mice. **p < 0.01, ***p < 0.001 compared to *Vhl/Hif-1 α* ^{F/F}. Seven to eight mice were assessed per group.

Supplement Figure 4 *Decrease in gluconeogenic genes after refeeding is associated with decreased CREB phosphorylation (Related to Figure 4)*. (A) Serum insulin levels measured in mice fed, overnight fasted or overnight fasted and refed for 30 minutes to 2-hours. (B) Western blot analysis in whole cell lysates of mice that were either fasted overnight or refed for 30 minutes to 2-hours. (C) qPCR analysis of *G6pase* and *Pepck*

mRNA in mice that were either fed, overnight fasted or refed for 30 minutes to 2-hours. Expression was normalized to β -actin mRNA. Each bar represents the mean value \pm S.E.M. * $p < 0.05$, *** $p < 0.001$ compared to fasted mice. Five mice were assessed per group.

Supplement figure 5 *Abrogation of glucagon signaling in Vhl^{LivKO} mice is due to increased hydrolysis of cAMP (Related to Figure 5)*. (A) G6pase luciferase assays in primary hepatocytes of $Vhl^{F/F}$ and Vhl^{LivKO} mice that were infected with Ad-G6pase-luciferase virus and then pre-treated with 0.2 μ M Okadaic acid for 2-hours, followed by treatment with 50 nM glucagon. Luciferase assays were performed in triplicates and activities normalized to protein content. Each bar represents the mean value \pm S.E.M. # $p < 0.05$, ### $p < 0.001$ compared to untreated. *** $p < 0.001$ compared to $Vhl^{F/F}$. (B) Western blot analysis of pCREB in primary hepatocyte lysate incubated invitro with recombinant PKA (2500U/ml; NewEngland Biolabs) for 30 minutes at 30C. (C) qPCR analysis of PKA subunit mRNA in the primary hepatocytes of $Vhl^{F/F}$ and Vhl^{LivKO} mice. (D) Western blot analysis of PKA subunits in the primary hepatocyte lysates from $Vhl^{F/F}$ and Vhl^{LivKO} mice. (E) Immunostaining for PKA-C in the primary hepatocytes from $Vhl^{F/F}$ and Vhl^{LivKO} mice that were treated with 50 nM glucagon for 30 minutes. (F) Western blot analysis for PKA-C in the nuclear lysates of $Vhl^{F/F}$ and Vhl^{LivKO} mice. (G) Co-immunoprecipitation with PKA-C antibody in the primary hepatocytes from $Vhl^{F/F}$ and Vhl^{LivKO} mice that were treated with 50 nM glucagon for 30 minutes and then probed for PKA-C and CREB. (H) qPCR analysis of AKAP-12 and Syk in the primary hepatocytes of $Vhl^{F/F}$ and Vhl^{LivKO} mice. (I) Western blot analysis for AKAP-12 in the primary

hepatocytes from *Vhl*^{F/F} and *Vhl*^{LivKO} mice. (J) qPCR analysis to assess glucagon induced *G6pase* and *Pepck* in the primary hepatocytes of *Vhl*^{F/F} and *Vhl*^{LivKO} mice that were pre-treated with Piceatannol (Syk inhibitor) or PMA (AKAP inhibitor) for 2-hours. qPCR analysis for *Pepck* in the primary hepatocytes of *Vhl*^{F/F} and *Vhl*^{LivKO} mice that were treated with (K) forskolin or (L) 200 μ M 8-br-cAMP for 2-hours. (M) Western blot analysis for p-PKA substrate antibody in the primary hepatocytes from *Vhl*^{F/F} and *Vhl*^{LivKO} mice that were treated with 200 μ M 8-br-cAMP for 30 minutes. Gene expression studies are done in triplicated and expression was normalized to β -actin. Each bar represents the mean value \pm S.E.M. #p < 0.05, ###p < 0.001 compared to untreated. *p < 0.05, ***p < 0.001 compared to *Vhl*^{F/F}. (N) Glucagon stimulated *Pepck* mRNA in *Vhl*^{F/F} and *Vhl*^{LivKO} primary hepatocytes pre-treated with 1 mM IBMX for one hour. (O) qPCR analysis for glucagon induced *G6pase* expression in the primary hepatocytes of *Vhl*^{LivKO} mice that were pre-treated with 20 μ M Bay-60-7550 (PDE2), 20 μ M Milrinone (PDE3), and 250 μ M Papaverine (PDE10) for 2-hours. qPCR analysis of PDEs in the primary hepatocytes of (P) *Vhl*^{F/F} and *Vhl*^{LivKO} mice or (Q) *Vhl/Hif2 α* ^{F/F} and *Vhl/Hif2 α* ^{LivKO} mice. *p < 0.05 compared to *Vhl*^{F/F} or *Vhl/Hif2 α* ^{F/F}. Gene expression studies are done in triplicates and expression was normalized to β -actin mRNA. Each bar represents the mean value \pm S.E.M. *p < 0.05, ***p < 0.001 compared to *Vhl*^{F/F}. # p < 0.05, ###p < 0.001 compared to Vehicle, \$p < 0.05, \$\$\$p < 0.001 are IBMX treated compared to their untreated controls. (R) Western blot analysis of PDE4D in liver lysates of *Vhl*^{F/F} and *Vhl*^{LivKO} mice. Experiments using primary hepatocytes were done in triplicates and repeated at least three times.

Supplement Figure 6 *Global change in PDEs results in blunted glucagon signaling in Vhl^{LivKO} mice (Related to Figure 6).* (A) Glucagon stimulated *G6pase* expression in primary hepatocytes of $Vhl^{F/F}$ and Vhl^{LivKO} mice infected with 100 MOI Adeno-GFP or Adeno-MEK1DN virus for 48 hours. #####, $p < 0.0001$ compared to no glucagon; ****, $p < 0.0001$ compared to $Vhl^{F/F}$; \$, $p < 0.05$ compared to Ad-GFP; \$\$, $p < 0.01$ compared to Ad-GFP. (B) Glucose production in hepatocytes pretreated with 10 μ M SU6656 for 16-hours and then treated with glucagon for 4-hours. Each bar represents the mean value \pm S.E.M. *, $p < 0.01$ compared to $Vhl^{F/F}$; ##, $p < 0.001$ compared to no glucagon; ^, $p < 0.05$ compared to substrate. (C) Western blot analysis of p42/p44 ERK phosphorylation in liver lysates from mice with compound disruption of VHL/ARNT ($Vhl/Arnt^{LivKO}$). (D) Blood glucose at fed state and (E) qPCR analysis for *G6pase* and *Pepck* in the livers of $Vhl/Arnt^{LivKO}$ mice. Four mice were used per group and experiments with primary hepatocytes were done in triplicates and repeated at least three times.

Supplement Figure 7 *Insulin signaling in STZ treated mice (Related to Figure 7).* (A) Serum insulin levels assessed in STZ treated mice. * $p < 0.05$ compared to untreated mice. (B) Co-immunostaining for insulin and glucagon performed in the pancreas of Vehicle or STZ treated mice. (C) Serum glucagon level and (D) Insulin:glucagon ratio assessed in STZ treated mice. Five to seven mice were assessed per group. * $p < 0.05$, *** $p < 0.001$ compared to Vhl^{LivKO} mice.

Supplement Figures

Figure S1

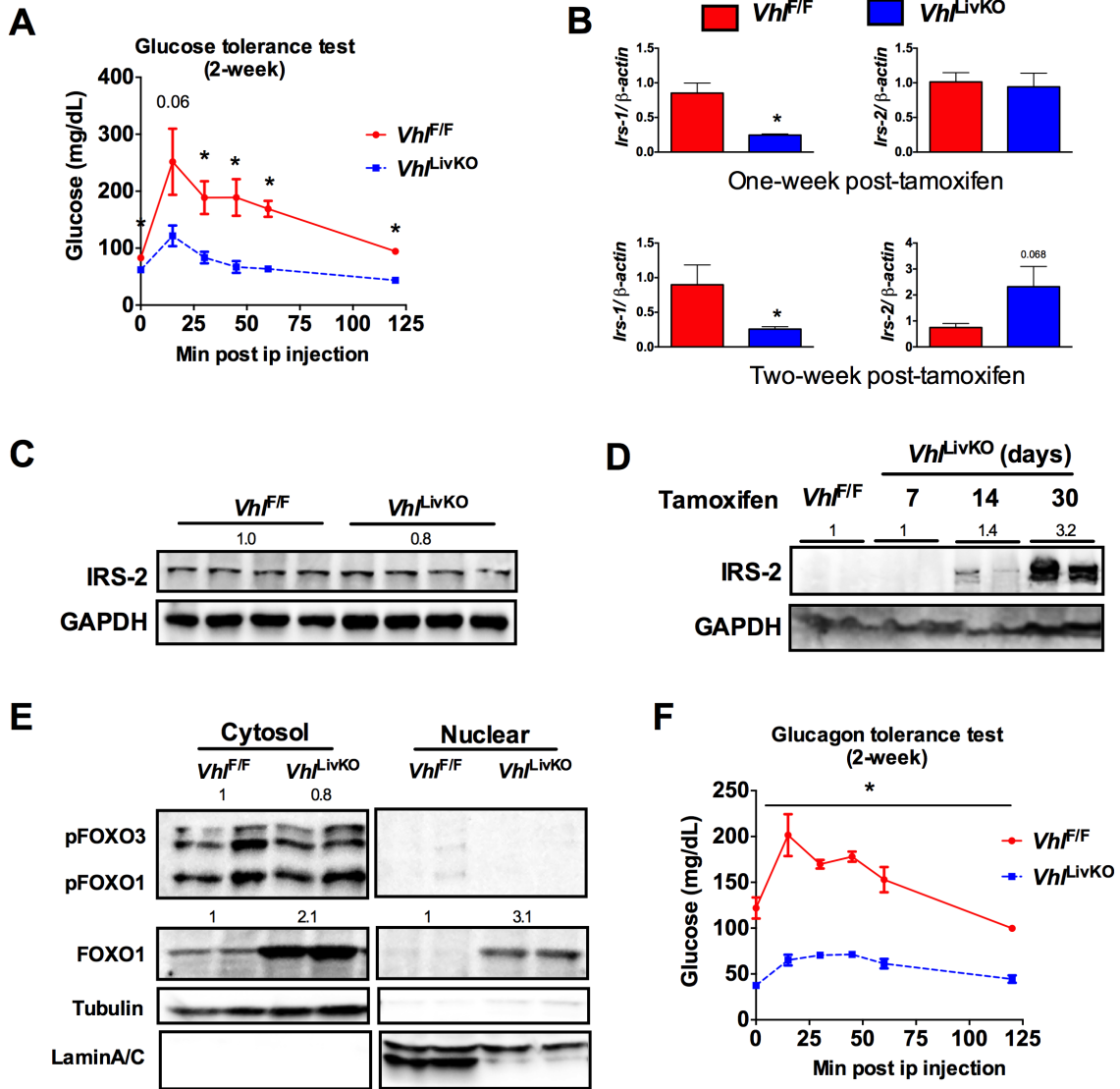


Figure S2

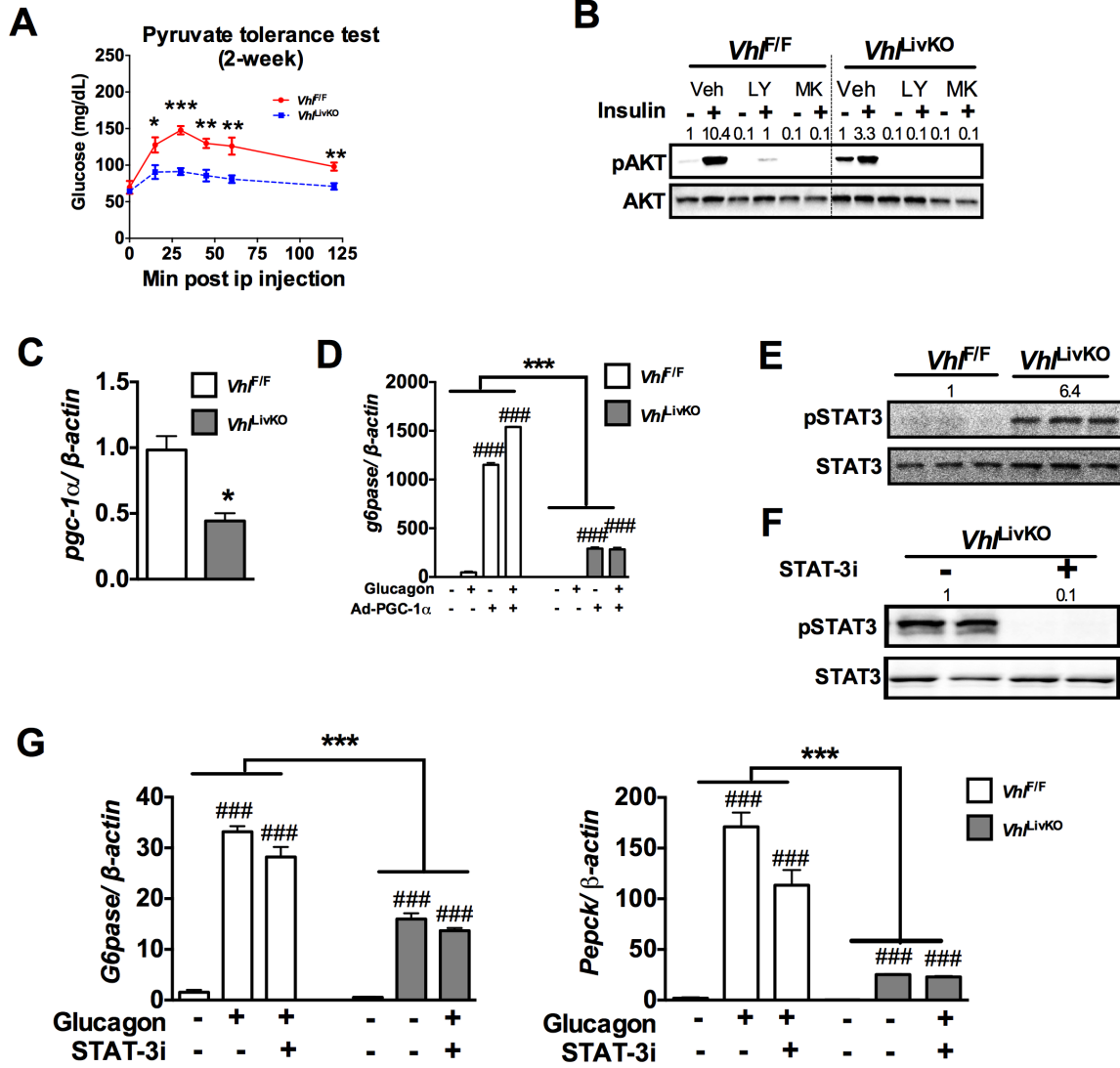
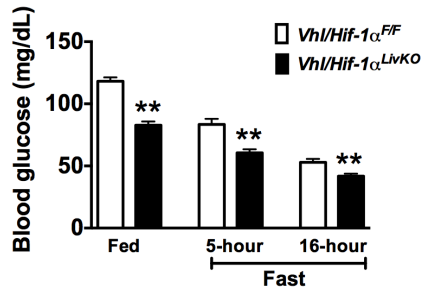


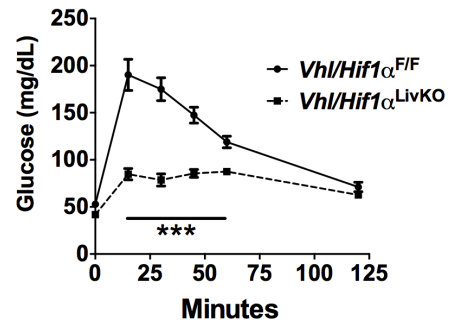
Figure S3

A



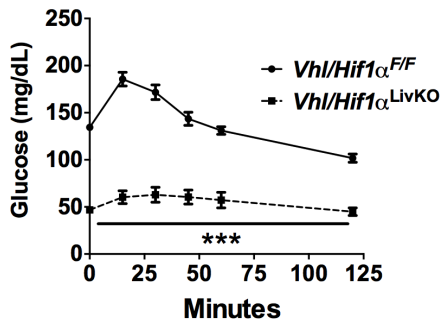
B

Glucose tolerance test



C

Glucagon tolerance test



D

Pyruvate tolerance test

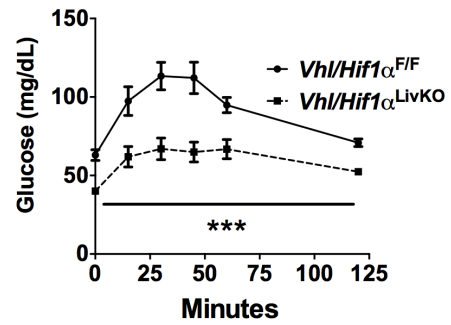


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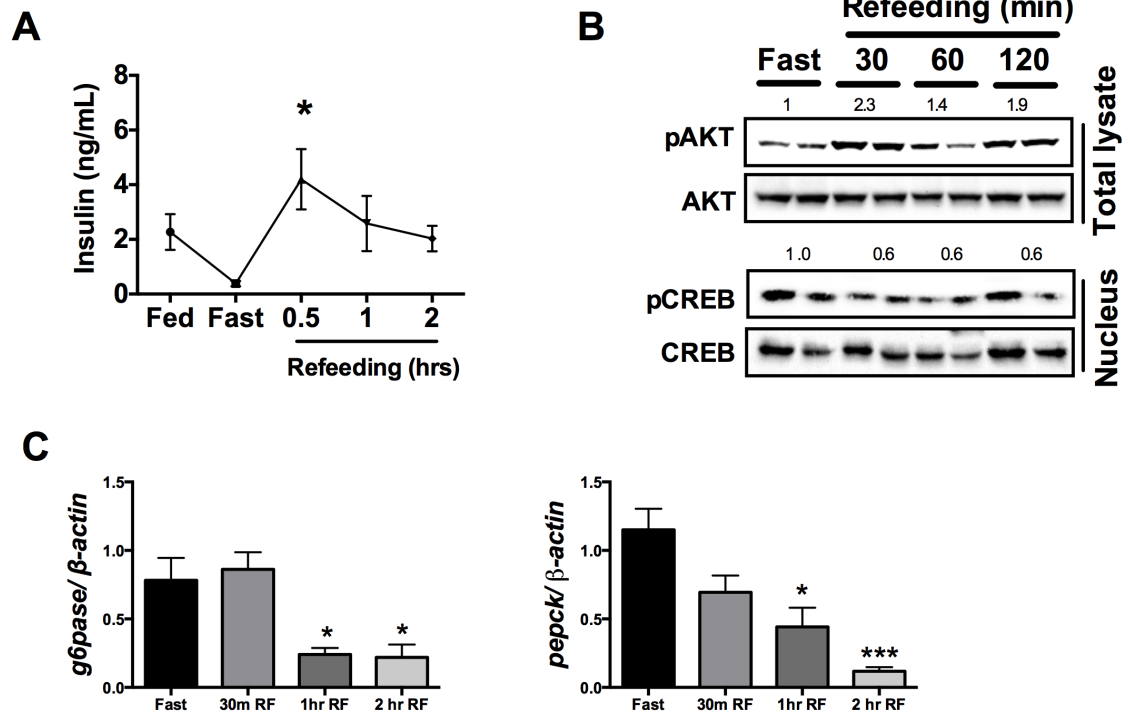


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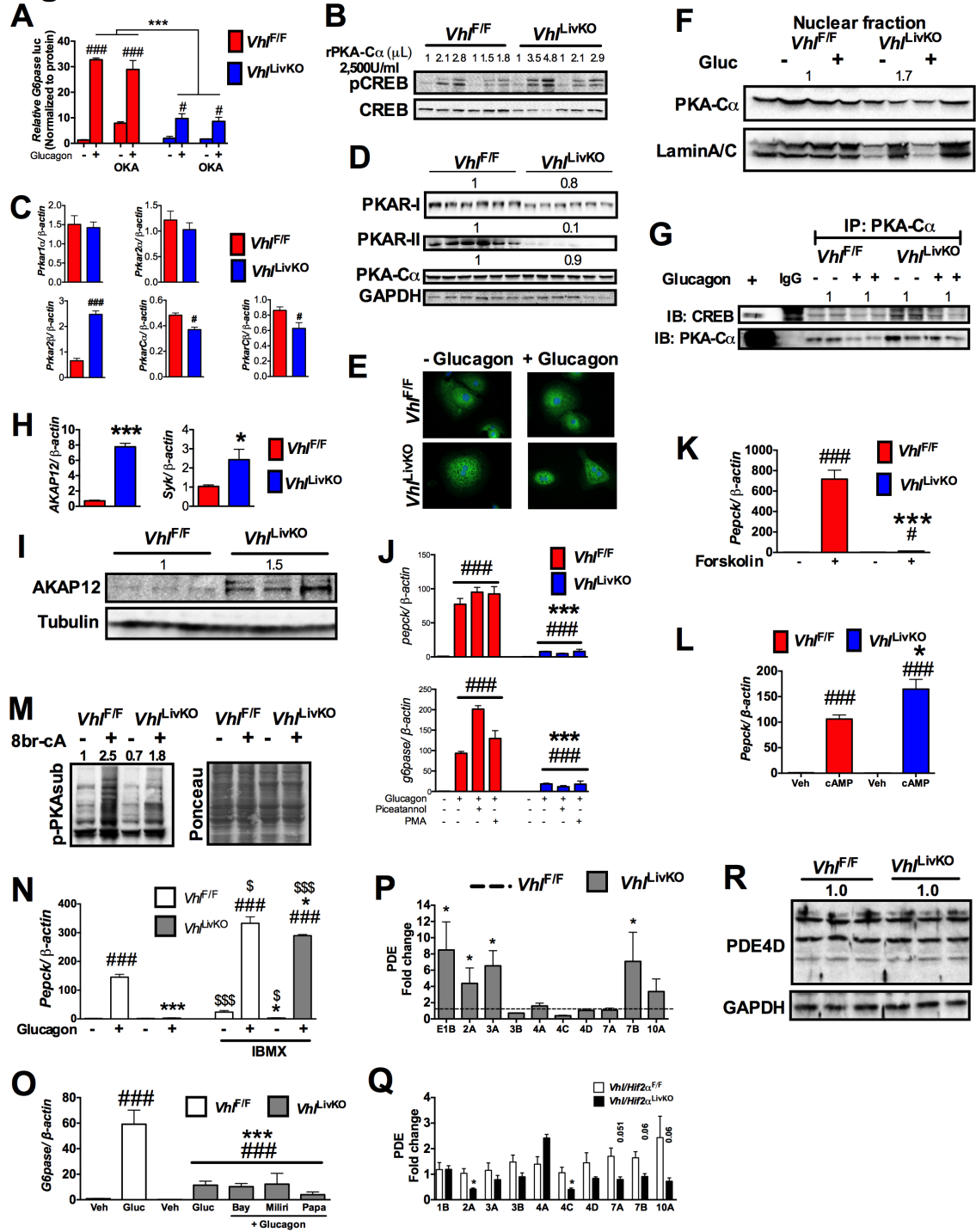


Figure S6

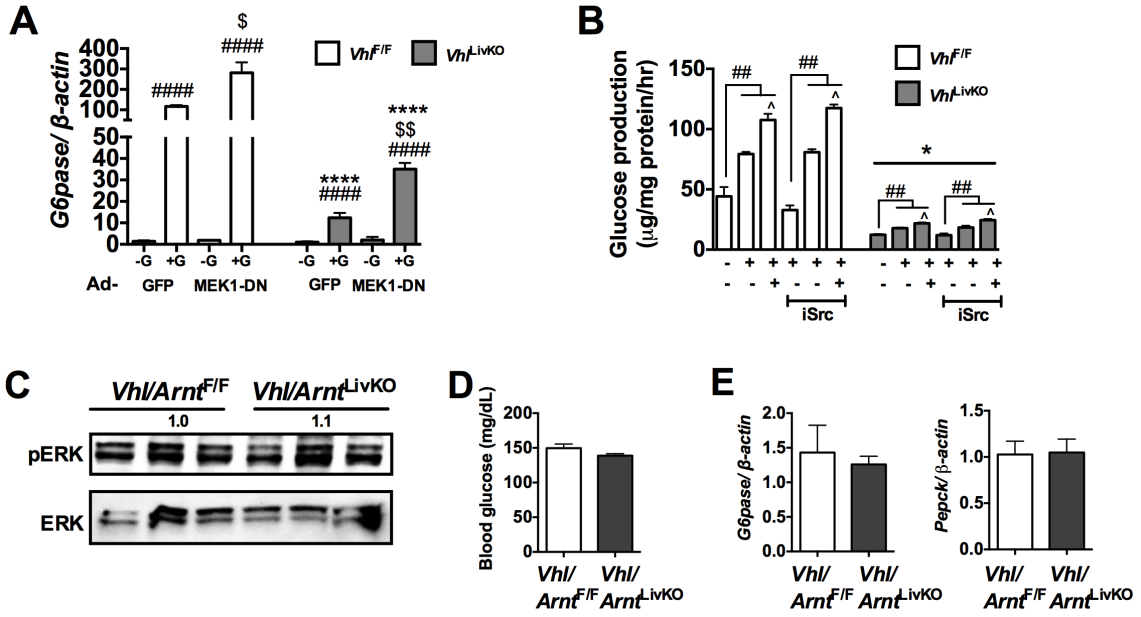
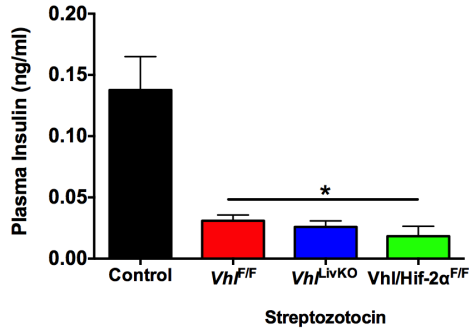
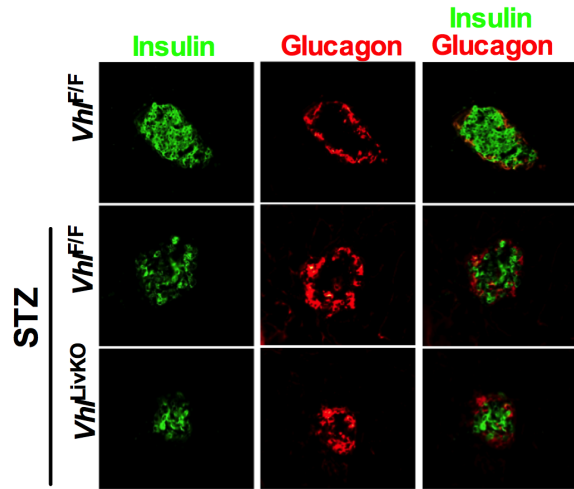


Figure S7

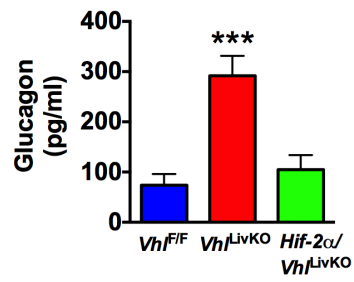
A



B



C



D

