1

A kidney-hypothalamus axis promotes compensatory glucose production in response to
glycosuria

Tumininu S. Faniyan[1], Xinyi Zhang[2], Donald A. Morgan[3], Jorge Robles[1], Siresha
Bathina[1], Paul S. Brookes[4], Kamal Rahmouni[3], Rachel J. Perry[2], Kavaljit H. Chhabra[1*]

[1]Department of Medicine, Division of Endocrinology, Diabetes, and Metabolism,
University of Rochester Medical Center; [2]Department of Cellular and Molecular
Physiology, Yale School of Medicine;[3]Department of Neuroscience and Pharmacology,
University of Iowa Carver College of Medicine; [4]Department of Anesthesiology,
University of Rochester Medical Center.

18 *Corresponding author. Email: Kavaljit_Chhabra@URMC.Rochester.Edu

19
20

Abstract

The kidneys facilitate energy conservation through reabsorption of nutrients including glucose. Almost all of the filtered blood glucose is reabsorbed by the kidneys. Loss of glucose in urine (glycosuria) is offset by an increase in endogenous glucose production to maintain normal energy supply in the body. How the body senses this glucose loss and consequently enhances glucose production is unclear. Using renal *Glut2* knockout mice, we demonstrate that elevated glycosuria activates the hypothalamic-pituitary-adrenal axis, which in turn drives endogenous glucose production. This phenotype was attenuated by selective afferent renal denervation, indicating the involvement of the afferent nerves in promoting the compensatory increase in glucose production. In addition, through plasma proteomics analyses we observed that acute phase proteins - which are usually involved in body's defense mechanisms against a threat – were the top candidates which were either upregulated or downregulated in renal *Glut2* KO mice. Overall, afferent renal nerves contribute to promoting endogenous glucose production in response to elevated glycosuria and loss of glucose in urine is sensed as a biological threat in mice. These findings may be useful in improving efficiency of drugs like SGLT2 inhibitors that are intended to treat hyperglycemia by enhancing glycosuria, but are met with a compensatory increase in endogenous glucose production.

Introduction

The kidneys help conserve energy by reabsorbing nutrients and preventing their loss in urine. Almost all of the filtered glucose is reabsorbed by the kidneys. The renal threshold for glucose reabsorption in humans is 180 mg/dl and in rodents is about 400 mg/dl[1]. In conditions such as overt hyperglycemia (beyond the renal glucose threshold), genetic loss of glucose transporters like GLUT2[2-4] or SGLT2[5], and/or renal dysfunction, glucose is excreted in urine. Enhancing glycosuria, through SGLT2 inhibition[6-8], is now used to reduce blood glucose levels in humans with diabetes. However, the glucose loss is offset by a compensatory increase in endogenous glucose production, consequently compromising the efficacy of this strategy in treatment of diabetes[9, 10]. We recently produced renal *Glut2* knockout mice[3], which show massive glycosuria and are protected from diabetes and diet-induced obesity. Despite the loss of glucose in urine, renal *Glut2* knockout mice maintain normal fasting blood glucose levels. Collectively, these observations indicate the presence of a fundamental mechanism involved in sensing glucose loss and triggering a compensatory increase in endogenous glucose production in rodents and humans. This mechanism is also aligned with the established survival strategies that reduce utilization and increase production of glucose (energy) during a flight or fight response against a threat[11, 12].

Understanding how the kidney senses glucose loss and consequently activates compensatory pathways to make up for this loss will enhance our knowledge about integrative mechanisms that regulate glucose homeostasis. Answering such research

questions may also be useful in medicine to increase efficiency of drugs such as SGLT2
inhibitors that are used to reduce blood glucose levels in diabetes by elevating
glycosuria.

In this study, we used renal *Glut2* knockout mice to identify mechanisms involved in
activating a compensatory increase in glucose production in response to glucose loss in
urine. We determined whether neural connections between the kidneys and
hypothalamus can explain the compensatory increase in glucose production. In
addition, we performed plasma proteomics analyses to identify secreted proteins or
endocrine factors that may provide insights into pathways involved in conserving energy
and producing endogenous glucose in response to elevated glycosuria in renal *Glut2*
knockout mice.

Methods

Mouse husbandry

All animal procedures were approved by the Institutional Animal Care and Use
Committee at the University of Rochester, Yale University, or University of Iowa, and
were performed according to the US Public Health Service guidelines for the humane
care and use of experimental animals. Mice were housed in ventilated cages under
controlled temperature (~23°C) and photoperiod (12 h light / dark cycle, lights on from
06:00 hours to 18:00 hours or 07:00 hours to 19:00 hours at Yale) conditions with free
access to Hydropac water (Lab Products, USA) and regular laboratory chow (5010,
LabDiet, USA or Global 2018, Harlan Teklad, USA). We used male and female renal
Glut2 knockout[3] and their littermate control mice in this study. After genotyping, mice
were randomly assigned to different experimental groups. Renal *Glut2* deficiency was
induced using tamoxifen as described previously[3].

Glucose production and liver metabolomics

Mice underwent surgery under isoflurane anesthesia to place a catheter in the jugular
vein, and were allowed to recover for 7 days prior to flux studies. They were fasted
overnight (16 h) to deplete liver glycogen prior to study. Mice were administered a 3X
primed-continuous infusion of [3-^{13}C] sodium lactate (continuous infusion rate 20
μmol/kg/min) and ^{2}H$_7$ glucose (continuous infusion rate 2.2 μmol/kg/min) for 120 min,
and euthanized with IV Euthasol. Glucose production from pyruvate was measured in
vivo in the liver and kidney (both freeze-clamped in situ within 30 second of euthanasia)
using mass isotopomer distribution analysis as described previously[13]. Briefly, we
measured the whole-body ratio of pyruvate carboxylase flux to total glucose production
by gas chromatography/mass spectrometry-mass spectrometry. We compared the
whole-body gluconeogenesis from phosphoenolpyruvate (PEP) and all substrates
upstream of PEP (i.e., pyruvate, lactate, amino acids) to that measured locally in the
liver and kidney. As the whole-body fraction of gluconeogenesis from PEP represents

an integrated contribution of both liver and kidney to whole-body glucose production
from PEP, this allowed us to –calculate the fractional contribution of the kidney to
whole-body glucose production.

For measuring liver metabolites involved in glycolysis, flash-frozen liver was
homogenized to extract metabolites in 80% methanol, which was evaporated under
liquid nitrogen followed by resuspension of metabolites in 50% methanol. Liquid
Chromatography Tandem Mass Spectrometry (LC-MS/MS) analysis was performed with
reverse-phase LC on a Synergi Fusion-RP column (Phenomenex, Torrence CA) at 35
°C. Samples were analyzed by single reaction monitoring on a Thermo Quantum triple-
quadrupole mass spectrometer. Metabolites were identified against a library of validated
standards based on retention time, intact mass, collision energy, and fragment masses.
Data were collected and analyzed using Thermo XCaliber 4.0 software. Post-hoc data
was analyzed using Metaboanalyst 5.0. Data were median-normalized (normalized to
the median peak area per sample, which correlated well with the total protein
concentration). 73 metabolites were measured, out of which 9 were significantly
($p<0.05$) different (1.5 fold-change threshold, Supplementary data table 1).

Measurement of genes and hormones involved in regulating hypothalamic-
pituitary-adrenal axis

We measured the levels of *Crh* mRNA in the mouse hypothalamus using RNA
fluorescence in situ hybridization (*Crh* probe, 316091, ACD) according to the
manufacturer's instructions. DAPI was used to stain the nucleus and verify the regions
of interest in the mouse brain. Images were captured using Keyence fluorescence
microscope BZ-X800. To further quantify the number of cells expressing *Crh* or DAPI,
we used CellProfiler 4.4.1 (https://cellprofiler.org/releases).

We used the following primers to measure gene expression using RT-qPCR: *Crh*: 5'-
GGAATCTCAACAGAAGTCCCGC-3' and 5'-CTGCAGCAACACGCGGAAAAAG-3',
Hrprt: 5′-AACAAAGTCTGGCCTGTATCC-3′ and 5′-CCCCAAAATGGTTAAGGTTGC-3′.
All primers were used at a final concentration of 500 nmol/l. The relative quantity of
each mRNA was calculated from standard curves and normalized to the internal control
Hprt, and then normalized to the mean of corresponding controls.

Plasma levels of adrenocorticotropic hormone (ab263880, Abcam) and corticosterone
(80556, Crystal Chem) were measured using ELISA per the manufacturers' instructions.

Ablation of afferent renal nerve

After anesthetizing mice using 1 – 5% isoflurane, a dorsal midline incision was made to
access both the kidneys. We selectively ablated afferent renal nerves using capsaicin.
Gauze, soaked in capsaicin (35 mM in 10% ethanol and 90% saline solution), was
wrapped around the renal blood vessels for 15 min. Other surrounding tissues were
covered with parafilm to avoid their exposure to capsaicin. After the surgery, muscle
layers and skin were closed with nylon sutures. This procedure ablates afferent renal

nerves without affecting the blood vessels or efferent renal nerves[14]. Control mice
underwent the same procedure without capsaicin (sham surgery using 10% ethanol and
90% saline solution). The mice were allowed to recover for a week before determining
the effects of the afferent denervation on blood glucose and other parameters.

Measurement of renal nerve activity, blood pressure, and heart rate

The mouse renal nerve activity was measured using multifiber recording as described
previously [15]. After anesthesia, the nerve innervating the left kidney was identified,
dissected free, and placed on a bipolar 36-gauge platinum–iridium electrode (A-M
Systems; Carlsborg, WA). The electrode was connected to a high impedance probe
(HIP-511; Grass Instruments Co., Quincy, MA), and the nerve signal was amplified 10^5
times with a Grass P5 AC pre-amplifier and filtered at low and high frequency cutoffs of
100 Hz and 1000 Hz, respectively. This nerve signal was directed to a speaker system
and to an oscilloscope (54501A, Hewlett–Packard Co., Palo Alto, CA) for auditory and
visual monitoring of the nerve activity. The signal was then directed to a resetting
voltage integrator (B600C, University of Iowa Bioengineering) that sums the total
voltage output in units of 1 V * sec before resetting to zero and counting the number of
spikes per second. The final neurograms were continuously routed to a MacLab
analogue–digital converter (8S, AD Instruments Castle Hill, New South Wales,
Australia) for permanent recording and data analysis on a Macintosh computer. Renal
sympathetic nerve activity was measured during 30 min. To measure the afferent renal
nerve activity, the nerve was cut distally and the signal was recorded for 30 min. Nerve
activity was corrected for post-mortem background activity to eliminate background
electrical noise in the measurements.

In addition to the nerve activity, arterial pressure and heart rate were measured using a
tapered MRE-040 tubing inserted into the left carotid artery and the other end
connected to a transducer (BP-100; iWorks System, Inc., Dover, NH) that led to an
ETH-250 Bridge/Bio Amplifier (CB Sciences; Milford, MA). Core body temperature of the
mouse was measured using a rectal probe (YSI 4000A Precision Thermometer; Yellow
Springs, OH) and maintained constant at 37.5 °C using a custom-made heated surgical
platform.

Two-dimensional difference gel electrophoresis (2D-DIGE)

Plasma proteins from control and renal *Glut2* knockout mice were labeled with Cy3 and
Cy5 dyes respectively. We also used an internal standard (labeled with Cy2 dye)
containing equal amounts of protein of each plasma sample as a quality control. Using
2D gel electrophoresis, we separated plasma proteins (30 µg in 2D sample buffer) on a
gel by both isoelectric point and molecular weight. We then imaged the gel using
Typhoon scanner. DeCyder software was used to identify protein spots of interest
(downregulated or upregulated by at least 25%), which were automatically picked from
the 2D gel with Ettan Spot Picker (Amersham BioSciences). The proteins were

identified by MALDI TOF/TOF mass spectrometry (AB SCIEX TOF/TOF™ 5800
System) using Swiss-Prot database.

Statistics

Data are shown as mean ± s.e.m. Results were analyzed by two-tailed Student's
unpaired t-test or two-way ANOVA followed by a Tukey's post hoc multiple comparison
test when appropriate. All analyses were performed using Prism version 8.0.1
(GraphPad, USA) and differences were considered statistically significant at $p < 0.05$.

Results and Discussion

Increased hepatic and renal glucose production in renal *Glut2* knockout mice

Renal *Glut2* knockout mice exhibit normal fasting blood glucose levels despite massive
glycosuria[3]. This observation suggests a compensatory increase in endogenous
glucose production, similar to that observed in humans following SGLT2 inhibition[9, 10,
16]. Therefore, we measured hepatic and renal glucose production in renal *Glut2*
knockout and their littermate control mice. We observed that both tissues contribute to
the compensatory increase in glucose production in renal *Glut2* knockout mice (Figure
1a-c). This was accompanied by a decrease in hepatic glucose 6-phosphate, glucose 1-
phosphate, and fructose 6-phosphate (Figure 1d-f), suggesting a decline in glucose
metabolism in favor of increasing glucose production. List of all measured metabolites is
included in Supplementary data table 1.

Renal *Glut2* knockout mice have an activated hypothalamic-pituitary-adrenal
(HPA) axis

The HPA axis is a major stress response system involved in enhancing glucose
production. Therefore, we measured plasma corticosterone and adrenocorticotropin
hormone coupled with gene expression of hypothalamic corticotropic releasing hormone
(*Crh*). We observed an increase in all these factors (Figure 2) that comprise the HPA
axis. These findings suggest that the HPA axis may contribute to the compensatory
increase in glucose production in renal *Glut2* knockout mice.

Afferent renal nerves contribute to enhancing glucose production in renal *Glut2*
knockout mice

To determine the role of afferent renal nerves in triggering a compensatory glucose
production in response to glycosuria, we used capsaicin to selectively suppress afferent
renal nerve activity in renal *Glut2* knockout mice. Afferent denervation was confirmed by
measuring renal pelvic CGRP (pg/mg protein) at the end of this study (Sham: 86.3 ±6.4
(control mice), 93.7 ±8.2 (renal *Glut2* knockout mice); Capsaicin: 4.7 ±0.06 (control
mice), 6.4 ±0.7 (renal *Glut2* knockout mice), using ELISA kit, Cayman Chemical,
589001). Mice were allowed to recover for a week before we fasted them to measure
their blood glucose levels. Because reinnervation may occur following the denervation,

we completed this study within 14 days after the denervation procedure. We observed
that fasting and fed (random) blood glucose levels were about 50% decreased in renal
Glut2 knockout mice compared to their control littermates (Figure 3a,b) after the afferent
renal denervation. In addition, the denervation reversed the activation of the HPA axis in
renal *Glut2* knockout mice (Figure 3c,d). Fasting plasma insulin levels were not
changed (Figure 3e) after the denervation. These observations suggest that renal
nerves contribute to the compensatory increase in glucose production.

Although the afferent renal denervation attenuated the compensatory increase in
glucose production in renal *Glut2* knockout mice, there was no change in their total or
afferent renal nerve activity (Supplementary Figure 1a,b). The *Glut2* knockout mice also
had similar blood pressure, heart rate coupled with similar weights of brown and white
adipose tissue, kidney, and liver (Supplementary Figure 1c-j). These findings suggest
the role of integrative neuroendocrine mechanisms independently of changes in nerve
activity, fat, kidney, or liver mass, in triggering a compensatory increase in glucose
production in renal *Glut2* knockout mice. Overall, the results indicate that afferent renal
nerves contribute to a compensatory increase in glucose production to make up for
glucose loss in renal *Glut2* knockout mice. These findings support data from
humans[16] that renal nerves may be involved in mediating a compensatory increase in
glucose production in response to elevated glycosuria by SGLT2 inhibition.

Changes in circulating acute phase proteins in renal *Glut2* knockout mice

To further investigate underlying mechanisms compensating for glucose loss in *Glut2*
knockout mice, we measured levels of secreted proteins that may explain the increase
in endogenous glucose production. We performed 2D-DIGE followed by proteomic
analyses with plasma samples collected from renal *Glut2* knockout mice and their
littermate controls. We observed that acute phase proteins like mannose binding lectin,
albumin, haptoglobin, ferritin were either upregulated or downregulated in renal *Glut2*
knockout mice (Figure 4 and Supplementary table 2). Usually levels of these proteins
are changed in response to threat, infection, and/or injury to activate body's defense
systems[17]. Similarly, major urinary proteins (MUP) like MUP18, which belong to the
lipocalin protein family, were elevated in renal *Glut2* knockout mice (Figure 4 and
Supplementary table 2). These proteins are involved in regulating inter-organ chemical
signaling, pheromonal communication, and energy metabolism[18]. We also observed
that secreted glutathione peroxidase 3 (Gpx3) was the most downregulated protein in
plasma in renal *Glut2* knockout mice (Figure 4 and Supplementary table 2). Gpx3 is
predominantly expressed in the kidneys and is an anti-oxidant[19]. We validated these
findings using qRT-PCR. Renal *Gpx3* gene expression was about 50% lower
(Supplementary Figure 2a) in renal *Glut2* knockout mice compared to their littermate
controls. Moreover, renal - but not hepatic - *Mup18* gene expression was higher
(Supplementary Figure 2b,c) in renal *Glut2* knockout mice relative to their controls,
suggesting that the kidneys were responsible for increasing plasma MUP18 in the
knockout mice. Altogether, these results suggest that multiple pathways - including local

(renal) stress and general (HPA axis) stress response combined with the acute phase
proteins - compensate for glucose loss and defend against a perceived biological threat
in renal *Glut2* knockout mice.

This study has limitations. Higher urine volume (polyuria) observed in renal *Glut2*
knockout mice may contribute to increase in glucose production to some extent,
however that contribution would likely be minor[20, 21] based on previous studies about
the role of diuretics (which increase urine volume) in glucose production. The extent to
which the changes in circulating secretory proteins observed in renal *Glut2* knockout
mice contribute to their compensatory increase in glucose production is unknown.
Further studies are necessary to address the role of these secretory proteins in glucose
and energy homeostasis. It is possible that afferent renal denervation in the present
study attenuated only hepatic glucose production through the hypothalamus without
affecting the compensatory increase in renal (local) glucose production. If validated, this
would explain the source of remaining endogenous glucose after the denervation in
renal *Glut2* knockout mice.

In summary, we report that a kidney-hypothalamus axis contributes to triggering a
compensatory increase in endogenous glucose production in response to elevated
glycosuria in mice. Glucose loss in urine appears to be sensed as a biological threat
and therefore activates body's defense systems including some of the acute phase
proteins. These findings may explain why SGLT2 inhibitors don't achieve their full
potential in lowering blood glucose levels during treatment of diabetes mellitus.

Author contributions

TSF designed and performed experiments including mouse genotyping, analysed
results, prepared graphs and figures, and edited the manuscript. XZ and RJP measured
glucose production, analysed results, and edited the manuscript. DAM and KR
performed electrophysiological recordings, analysed the results, and edited the
manuscript. JR and SB performed microscopy, qRT-PCR or ELISA, and edited the
manuscript. PB performed metabolomics, analysed the results, and edited the
manuscript. KHC conceived the study, designed and performed experiments, analysed
results, wrote and edited the manuscript. All authors approved the final version of the
manuscript.
299
Disclosures: Authors declare they have no conflict of interests.

Acknowledgments

We thank V. Kaye Thomas, Julie Zhang, URMC Center for Advanced Light Microscopy
and Nanoscopy, for help with microscopy; Applied Biomics for proteomics analyses.

Funding

National Institutes of Health grant CA258261 to RJP
National Institutes of Health grant HL071158 PSB.
National Institutes of Health grant DK124619 to KHC.

Startup funds, Department of Medicine, University of Rochester, NY to KHC.

Data and materials availability

References

[1]   Noonan WT, Banks RO (2000) Renal function and glucose transport in male and female mice with diet-induced type II diabetes mellitus. Proc Soc Exp Biol Med 225(3): 221-230. 10.1046/j.1525-1373.2000.22528.x

[2]   de Souza Cordeiro LM, Elsheikh A, Devisetty N, et al. (2021) Hypothalamic MC4R regulates glucose homeostasis through adrenaline-mediated control of glucose reabsorption via renal GLUT2 in mice. Diabetologia 64(1): 181-194. 10.1007/s00125-020-05289-z

[3]   de Souza Cordeiro LM, Bainbridge L, Devisetty N, McDougal DH, Peters DJM, Chhabra KH (2022) Loss of function of renal Glut2 reverses hyperglycaemia and normalises body weight in mouse models of diabetes and obesity. Diabetologia 65(6): 1032-1047. 10.1007/s00125-022-05676-8

[4]   Sakamoto O, Ogawa E, Ohura T, et al. (2000) Mutation Analysis of the GLUT2 Gene in Patients with Fanconi-Bickel Syndrome. Pediatric Research 48(5): 586-589. 10.1203/00006450-200011000-00005

[5]   Kleta R, Stuart C, Gill FA, Gahl WA (2004) Renal glucosuria due to SGLT2 mutations. Mol Genet Metab 82(1): 56-58. 10.1016/j.ymgme.2004.01.018

[6]   Rossetti L, Smith D, Shulman GI, Papachristou D, DeFronzo RA (1987) Correction of hyperglycemia with phlorizin normalizes tissue sensitivity to insulin in diabetic rats. J Clin Invest 79(5): 1510-1515. 10.1172/jci112981

[7]   Oku A, Ueta K, Arakawa K, et al. (1999) T-1095, an inhibitor of renal Na+-glucose cotransporters, may provide a novel approach to treating diabetes. Diabetes 48(9): 1794-1800. 10.2337/diabetes.48.9.1794

[8]   Han S, Hagan DL, Taylor JR, et al. (2008) Dapagliflozin, a selective SGLT2 inhibitor, improves glucose homeostasis in normal and diabetic rats. Diabetes 57(6): 1723-1729. 10.2337/db07-1472

[9]   Solis-Herrera C, Daniele G, Alatrach M, et al. (2020) Increase in Endogenous Glucose Production With SGLT2 Inhibition Is Unchanged by Renal Denervation and Correlates Strongly With the Increase in Urinary Glucose Excretion. Diabetes Care 43(5): 1065-1069. 10.2337/dc19-2177

[10]   Liu JJ, Lee T, DeFronzo RA (2012) Why Do SGLT2 inhibitors inhibit only 30-50% of renal glucose reabsorption in humans? Diabetes 61(9): 2199-2204. 10.2337/db12-0052

[11]   Surwit RS, Schneider MS, Feinglos MN (1992) Stress and diabetes mellitus. Diabetes Care 15(10): 1413-1422. 10.2337/diacare.15.10.1413

[12]   Marik PE, Bellomo R (2013) Stress hyperglycemia: an essential survival response! Crit Care 17(2): 305. 10.1186/cc12514

[13]   Qing H, Desrouleaux R, Israni-Winger K, et al. (2020) Origin and Function of Stress-Induced IL-6 in Murine Models. Cell 182(2): 372-387.e314. 10.1016/j.cell.2020.05.054

[14]   Foss JD, Wainford RD, Engeland WC, Fink GD, Osborn JW (2015) A novel method of selective ablation of afferent renal nerves by periaxonal application of capsaicin. Am J Physiol Regul Integr Comp Physiol 308(2): R112-122. 10.1152/ajpregu.00427.2014

[15]   Morgan DA, Rahmouni K (2010) Differential effects of insulin on sympathetic nerve activity in
agouti obese mice. J Hypertens 28(9): 1913-1919. 10.1097/HJH.0b013e32833c2289
[16]   Daniele G, Solis-Herrera C, Dardano A, et al. (2020) Increase in endogenous glucose production
with SGLT2 inhibition is attenuated in individuals who underwent kidney transplantation and bilateral
native nephrectomy. Diabetologia 63(11): 2423-2433. 10.1007/s00125-020-05254-w
[17]   Gabay C, Kushner I (1999) Acute-phase proteins and other systemic responses to inflammation.
N Engl J Med 340(6): 448-454. 10.1056/nejm199902113400607
[18]   Zhou Y, Rui L (2010) Major urinary protein regulation of chemical communication and nutrient
metabolism. Vitam Horm 83: 151-163. 10.1016/s0083-6729(10)83006-7
[19]   Whitin JC, Bhamre S, Tham DM, Cohen HJ (2002) Extracellular glutathione peroxidase is secreted
basolaterally by human renal proximal tubule cells. Am J Physiol Renal Physiol 283(1): F20-28.
10.1152/ajprenal.00014.2001
[20]   Hall JJ, Eurich DT, Nagy D, Tjosvold L, Gamble J-M (2020) Thiazide Diuretic–Induced Change in
Fasting Plasma Glucose: a Meta-analysis of Randomized Clinical Trials. Journal of General Internal
Medicine 35(6): 1849-1860. 10.1007/s11606-020-05731-3
[21]   Perry RJ, Rabin-Court A, Song JD, et al. (2019) Dehydration and insulinopenia are necessary and
sufficient for euglycemic ketoacidosis in SGLT2 inhibitor-treated rats. Nat Commun 10(1): 548.
10.1038/s41467-019-08466-w

Figure 1. Renal *Glut2* knockout mice exhibit increased glucose production. In vivo increase in total **(a)**, hepatic **(b)**, and renal **(c)** glucose production through gluconeogenesis with pyruvate as a substrate in 28 weeks old male renal *Glut2* knockout mice. Decreased hepatic glucose 6-phosphate **(d)**, glucose 1-phosphate **(e)**, and fructose 6-phosphate **(f)** in renal *Glut2* knockout mice. *p<0.05, ***p<0.001, unpaired 2-tailed Student's t-test. Data are presented as mean ±sem.

Figure 2. Enhanced activity of hypothalamic-pituitary-adrenal axis in renal *Glut2* knockout mice. Representative images from fluorescence RNA in situ hybridization showing an increase in expression of corticotropin releasing hormone (Crh) in the paraventricular nucleus of the hypothalamus in 28 weeks old male renal *Glut2* knockout mice **(a)**. Scale, 100 μm. For the quantification shown next to the images, 4 sections per mouse and 3 areas of interest per section were analyzed in 4 mice. qRT-PCR analysis showing an increase in hypothalamic *Crh* **(b)**, data from ELISA demonstrating an increase in plasma adrenocorticotropic hormone (ACTH) **(c)** and corticosterone **(d)** in 12 weeks old male renal *Glut2* knockout mice. *p<0.05, **p<0.01, ***p<0.001, unpaired 2-tailed Student's t-test. Data are presented as mean ±sem.

Figure 3. Effects of afferent renal denervation on blood glucose levels and hypothalamic-pituitary-adrenal axis in renal *Glut2* knockout mice. Afferent renal denervation decreases fed (random) and fasting (overnight, 6:00 pm – 9:00 am) blood glucose levels **(a,b)**, restores expression of hypothalamic corticotropin releasing hormone (Crh) **(c)** measured using RT-qPCR, and plasma corticosterone **(d)** without affecting plasma insulin levels **(e)** in 30 weeks old female renal *Glut2* knockout mice. **p<0.01, ***p<0.001, two-way ANOVA followed by a Tukey's post hoc multiple comparison test. Data are presented as mean ±sem.

Figure 4. Changes in levels of plasma proteins in renal *Glut2* knockout mice. Representative image of two-dimensional difference gel electrophoresis with numbered protein spots of interest is shown in **(a)**. Internal standard was prepared using equal amounts of protein of each plasma sample as a quality control (Cy2 labeled, pseudo blue), plasma proteins from control group were labeled using Cy3 dye (shown in pseudo green), and plasma proteins from renal *Glut2* knockout mice were labeled using Cy5 dye (shown in pseudo red). The identified proteins and their fold change in 28 weeks old male renal *Glut2* knockout mice compared to the control group is shown in **(b)**. The number on each bar graph in **(b)** represents the corresponding protein spot on the gel shown in **(a)**. MBL1, mannose binding lectin 1; Gpx3, glutathione peroxidase 3; MUP18, major urinary protein 18.