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Article

Metabolism

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# The AMP-activated protein kinase $\alpha 2$ catalytic subunit controls whole-body insulin sensitivity

Benoit Viollet,<sup>1</sup> Fabrizio Andreelli,<sup>1</sup> Sebastian B. Jørgensen,<sup>2</sup> Christophe Perrin,<sup>3</sup> Alain Geloën,<sup>4</sup> Daisy Flamez,<sup>5</sup> James Mu,<sup>6</sup> Claudia Lenzner,<sup>1</sup> Olivier Baud,<sup>1</sup> Myriam Bennoun,<sup>1</sup> Emmanuel Gomas,<sup>1</sup> Gaël Nicolas,<sup>1</sup> Jørgen F.P. Wojtaszewski,<sup>2</sup> Axel Kahn,<sup>1</sup> David Carling,<sup>7</sup> Frans C. Schuit,<sup>5</sup> Morris J. Birnbaum,<sup>6</sup> Erik A. Richter,<sup>2</sup> Rémy Burcelin,<sup>3</sup> and Sophie Vaulont<sup>1</sup>

<sup>1</sup>Department of Genetic, Development, and Molecular Pathology, Institut Cochin, Institut National de la Santé et de la Recherche Médicale (INSERM), Centre National de la Recherche Scientifique (CNRS), René Descartes University, Paris, France

<sup>2</sup>Copenhagen Muscle Research Centre, Department of Human Physiology, Institute of Exercise and Sport Sciences, University of Copenhagen, Copenhagen, Denmark

<sup>3</sup>Unité Mixte de Recherche 5018, CNRS, Paul Sabatier University, Rangueil Hospital, Toulouse, France

<sup>4</sup>INSERM, Institut National des Sciences Appliquées-Lyon, Villeurbanne, France

<sup>5</sup>Diabetes Research Center, Laboratory of Biochemistry, Vrije Universiteit Brussel, Brussels, Belgium

<sup>6</sup>Howard Hughes Medical Institute, University of Pennsylvania School of Medicine, Philadelphia, Pennsylvania, USA

<sup>7</sup>Cellular Stress Group, Medical Research Council Clinical Sciences Centre, Hammersmith Hospital, London, United Kingdom

AMP-activated protein kinase (AMPK) is viewed as a fuel sensor for glucose and lipid metabolism. To better understand the physiological role of AMPK, we generated a knockout mouse model in which the AMPK $\alpha 2$  catalytic subunit gene was inactivated. AMPK $\alpha 2^{-/-}$  mice presented high glucose levels in the fed period and during an oral glucose challenge associated with low insulin plasma levels. However, in isolated AMPK $\alpha 2^{-/-}$  pancreatic islets, glucose- and L-arginine-stimulated insulin secretion were not affected. AMPK $\alpha 2^{-/-}$  mice have reduced insulin-stimulated whole-body glucose utilization and muscle glycogen synthesis rates assessed *in vivo* by the hyperinsulinemic euglycemic clamp technique. Surprisingly, both parameters were not altered in mice expressing a dominant-negative mutant of AMPK in skeletal muscle. Furthermore, glucose transport was normal in incubated isolated AMPK $\alpha 2^{-/-}$  muscles. These data indicate that AMPK $\alpha 2$  in tissues other than skeletal muscles regulates insulin action. Concordantly, we found an increased daily urinary catecholamine excretion in AMPK $\alpha 2^{-/-}$  mice, suggesting altered function of the autonomic nervous system that could explain both the impaired insulin secretion and insulin sensitivity observed *in vivo*. Therefore, extra-muscular AMPK $\alpha 2$  catalytic subunit is important for whole-body insulin action *in vivo*, probably through modulation of sympathetic nervous activity.

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## Introduction

The AMP-activated protein kinase (AMPK) has been proposed to act as a fuel sensor capable of mediating the cellular adaptation to nutritional and environmental variations (1). AMPK is a ubiquitous serine/threonine protein kinase activated in response to environmental or nutritional stress factors that deplete intracellular ATP levels. AMPK is a heterotrimeric enzyme consisting of an

$\alpha$  catalytic subunit and noncatalytic regulatory  $\beta$  and  $\gamma$  subunits. Two isoforms have been identified for both the  $\alpha$  subunit ( $\alpha 1$  and  $\alpha 2$ ) and  $\beta$  subunit ( $\beta 1$  and  $\beta 2$ ), and three isoforms have been reported for the  $\gamma$  subunit ( $\gamma 1$ ,  $\gamma 2$ , and  $\gamma 3$ ). Regulation of AMPK activity is complex, involving allosteric activation by AMP and covalent modification by phosphorylation via an upstream AMPK kinase (2). Phosphorylation of threonine 172 in the  $\alpha$  subunit of AMPK is essential for its activation (2).

Metabolic actions of AMPK have been extensively studied in skeletal muscle. Activation of AMPK by 5-aminoimidazole-4-carboxamide-1- $\beta$ -D-ribofuranoside (AICAR) has been demonstrated to increase muscle glucose uptake *in vivo* and *in vitro* via an acute insulin-independent mechanism (3, 4) and to decrease the glycogen synthase activity ratio (5). Implication of AMPK in the physiology of tissues other than skeletal muscle has also been suggested by findings in liver (6–8), pancreas (9, 10), and adipose tissue (4, 7, 11, 12). Lastly, AMPK is also expressed in the CNS (13), but the role of AMPK in this tissue remains to be determined.

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**Address correspondence to:** B. Viollet, Institut Cochin, Département GDPM, 24 Rue du Faubourg Saint-Jacques 75014 Paris, France. Phone: 33-1-44-41-2408; Fax: 33-1-44-41-2421; E-mail: viollet@cochin.inserm.fr.

Benoit Viollet and Fabrizio Andreelli contributed equally to this work.

**Conflict of interest:** The authors have declared that no conflict of interest exists.

**Nonstandard abbreviations used:** AMP-activated protein kinase (AMPK); 5-aminoimidazole-4-carboxamide-1- $\beta$ -D-ribofuranoside (AICAR); kinase-dead AMPK $\alpha 2$  transgenic (Tg-KD-AMPK $\alpha 2$ ); oral glucose tolerance test (OGTT); acetyl-CoA carboxylase (ACC).

In skeletal muscle, the actions of AMPK are thought to be mediated mainly through the AMPK $\alpha$ 2 isoform (5, 14, 15). In other tissues, the physiological role of this isoform *in vivo* is less clear. To get more insights into this issue, we knocked out the *AMPK $\alpha$ 2* gene. In the present study, we show that *AMPK $\alpha$ 2*<sup>-/-</sup> mice exhibit glucose intolerance that is associated with low plasma insulin levels and reduced muscle glycogen synthesis during hyperinsulinemic clamp conditions. Reduction of insulin sensitivity was not dependent on muscle AMPK activity, since mice expressing a dominant-negative mutant of AMPK in skeletal muscle are not insulin resistant. Finally, *AMPK $\alpha$ 2*<sup>-/-</sup> mice have increased catecholamine urinary excretion, suggesting increased sympathetic tone. Consequently, our data suggest that the  $\alpha$ 2 catalytic subunit of AMPK plays a major role *in vivo* as a fuel metabolic sensor by modulating the activity of the autonomous nervous system.

## Methods

**Gene targeting and generation of AMPK $\alpha$ 2 gene knockout mice.** *AMPK $\alpha$ 2* genomic clones were isolated after screening a mouse 129 strain genomic library (Stratagene, Loyola, California, USA). The targeting construct was generated by flanking exon C, which encodes the AMPK $\alpha$ 2 catalytic domain (corresponding to amino acids 189–260), with *loxP* sites for the Cre recombinase and inserting a PGK promoter-driven neomycin selection cassette flanked by an additional *loxP* site. The targeting construct was linearized and electroporated into 129/Sv MPI-I embryonic stem cells. Targeted clones were identified by Southern blot analysis of *Hind*III-digested DNA using a flanking 5' DNA fragment as a hybridization probe and PCR analysis. Cells expanded from targeted clones were injected into C57BL/6 blastocysts, and germline-transmitting chimeric animals were mated with C57BL/6 mice. The resulting heterozygous offspring were bred with "deleter" EIIa-Cre transgenic mice, in which Cre is expressed in the germ cells, to produce *AMPK $\alpha$ 2*<sup>-/-</sup> mice (16). These mice were crossed to generate control and mutant mice. Southern blot analysis of *Hind*III-digested DNA with an internal probe (Figure 1a) and PCR analysis were used to genotype mice.

**Animals.** Control, *AMPK $\alpha$ 2*<sup>-/-</sup>, and kinase-dead AMPK $\alpha$ 2 transgenic (Tg-KD-AMPK $\alpha$ 2) male mice were studied at 12 weeks of age. The establishment and characterization of Tg-KD-AMPK $\alpha$ 2 mice were previously reported (17). Animals were housed under controlled temperature (21°C) and lighting, with 12 hours of light (7:00 am to 7:00 pm) and 12 hours of dark (7:00 pm to 7:00 am), and had free access to water and a standard mouse chow diet. All procedures were performed in accordance with the principles and guidelines established by the European Convention for the Protection of Laboratory Animals.

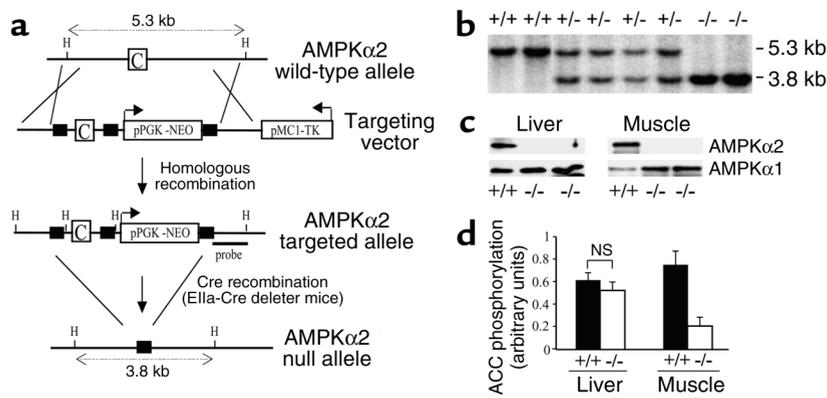
**Body weight, food intake, and body composition.** Mice were weighed weekly after 3 weeks of age. Food intake was measured daily for a period of 3 days using metabolic

cages (Marty Technologie, Paris, France) that allow free access to tap water and food. Body composition analysis was performed on anesthetized living mice by a dual-energy x-ray absorptiometry method using a small animal densitometer (PIXImus Lunar; GE Medical Systems, Madison, Wisconsin, USA). Careful dissection and weighing of individual organs was also performed. Tissue triglycerides were extracted with acetone, and triglyceride content was measured with a commercial kit (Sigma-Aldrich, St. Louis, Missouri, USA).

**Blood and urinary parameters.** Blood was withdrawn from the tail using EDTA-aptotinin as the anticoagulant. In the fed state, blood was collected at 9:00 pm. For fasting experiments, food was removed at 5:00 pm, and the mice were kept in a clean new cage for 5 hours before blood collection. During the oral glucose tolerance test (OGTT) and AICAR tolerance test, blood glucose levels were assessed using a glucometer (GlucoTrend II; Roche Diagnostics, Meylan, France). Serum insulin concentrations were assessed using a rat insulin ELISA kit with a mouse insulin standard (Crystal Chem Inc., Chicago, Illinois, USA). Serum leptin concentrations were assessed with a mouse leptin ELISA kit (Crystal Chem Inc.). Serum concentrations of potassium, sodium, triglycerides, FFAs, glycerol, total and HDL cholesterol, creatinine, and urea were determined using an automated Monarch device (Instrumentation Laboratory Co., Lexington, Massachusetts, USA). For determination of urinary levels of catecholamines, animals were housed in individual metabolic cages, and daily urinary catecholamine excretion during 3 consecutive days was determined by reverse-phase HPLC (performed at Cochin Hospital, Paris, France).

**Glucose and AICAR tolerance tests.** During OGTT and AICAR tolerance testing, blood was collected from the tail vein at 0, 20, 40, 60, 80, 100, and 120 minutes for determination of glucose levels. OGTTs were performed in the morning after an overnight fast by oral administration of 3 g glucose per kg body weight. Blood was also collected at time 20 minutes in chilled heparinized tubes for determination of plasma insulin levels. To study the effect of  $\alpha$ -adrenergic antagonist on glucose tolerance, mice were injected intraperitoneally with 10 mg/kg phentolamine 30 minutes before glucose challenge. To study the effect of  $\beta$ -adrenergic antagonist on glucose tolerance, mice were injected intraperitoneally with 10 mg/kg propranolol 30 minutes before glucose challenge. AICAR tolerance tests were performed in the morning by intraperitoneal injection of 0.25 g AICAR per kg body weight into overnight-fasted mice (17).

**Hyperinsulinemic euglycemic clamp procedures.** To determine the rate of glucose utilization, an indwelling catheter was inserted into the femoral vein under anesthesia, sealed under the back skin, and glued on top of the skull (18). The mice were then housed individually. After 2 days, they showed normal feeding behavior and motor activity. The mice were allowed to recover for 4–6 days. On the day of the experiment, the mice were fast-



**Figure 1**

Generation of mice lacking AMPK $\alpha$ 2. (a) Schematic representation (not to scale) of genomic structure of AMPK $\alpha$ 2 wild-type allele, AMPK $\alpha$ 2 gene-targeting construct, AMPK $\alpha$ 2 targeted allele, and AMPK $\alpha$ 2 null allele. Squares indicate *loxP* sites and H's indicate *Hind*III restriction sites. C corresponds to the exon encoding the AMPK $\alpha$ 2 catalytic domain (amino acids 189–260) (b) Southern blot analysis after *Hind*III digestion of tail DNA from offspring derived from heterozygous intercrosses. Expected fragment sizes of the AMPK $\alpha$ 2 wild-type (+/+; 5.3 kb) and null (-/-; 3.8 kb) alleles after *Hind*III digestion and hybridization with the indicated probe (solid bar in a) are shown. (c) Western blot analysis of AMPK $\alpha$ 1 and AMPK $\alpha$ 2 proteins in liver and gastrocnemius muscle from control (+/+) and AMPK $\alpha$ 2<sup>-/-</sup> mice. (d) Phosphorylation level of ACC in liver and gastrocnemius muscle from control and AMPK $\alpha$ 2<sup>-/-</sup> mice.

ed for 6 hours. The whole-body glucose utilization rate was determined under basal and hyperinsulinemic euglycemic conditions. In the basal state, HPLC-purified D-(3-<sup>3</sup>H)-glucose (NEN Life Science Products Inc., Boston, Massachusetts, USA) was continuously infused through the femoral vein at a rate of 10  $\mu$ Ci/kg.min for 3 hours. In the hyperinsulinemic condition, insulin was infused at a rate of 6 mU/kg.min for 3 hours; D-(3-<sup>3</sup>H)-glucose was infused at rate of 30  $\mu$ Ci/kg.min — a higher rate than for the basal condition, to ensure a detectable plasma D-(3-<sup>3</sup>H)-glucose enrichment. Throughout the infusion, blood glucose was assessed from blood samples and euglycemia was maintained by periodically adjusting a variable infusion of 16.5% (wt/vol) glucose. Plasma glucose concentrations, D-(3-<sup>3</sup>H)-glucose specific activity, and <sup>3</sup>H<sub>2</sub>O and D-(3-<sup>3</sup>H)-glucose enrichments were determined as previously described (18). Glucose turnover and whole-body glycolysis rate were calculated as described (18). Whole-body glycogen synthesis was calculated by subtracting the whole-body glycolytic rate from the glucose turnover rate. The mean was calculated for each timepoint for this period. The mean value of mice from the same group was reported. Mice showing variations in specific activity larger than 15% were excluded from the study.

**Determination of total glycogen content and glycogen synthesis rate in skeletal muscle and liver.** At completion of the infusion, the hind limb and the main lobe of the liver were removed and ground into liquid nitrogen for determination of total glycogen content and glycogen synthesis rate. Total glycogen was extracted from 20 mg of liver or 100 mg of muscle by digesting tissue samples in KOH. An aliquot was neutralized with HCl and the

glycogen content was assessed using amyloglucosidase (19). Muscleder glycogen synthesis rate was determined by extracting glycogen with perchloric acid (6% wt/vol) and precipitating glycogen with ethanol as described previously (20). The radioactive glycogen was counted and divided by the D-(3-<sup>3</sup>H)-glucose specific activity to determine the rate of synthesis.

**Deoxyglucose uptake in isolated muscles.** Mice were anesthetized with pentobarbital and soleus and extensor digitorum longus muscles were exposed. Ligatures were placed around the tendon at each end of the muscle. Muscles were suspended at resting tension (4–5 m Newtons) and incubated at 30°C for 10 minutes at basal conditions in incubation chambers (Multi myograph system organ bath 700MO; Danish Myo Technology A/S, Aarhus, Denmark) oxygenated with 95% O<sub>2</sub> and 5% CO<sub>2</sub> gas. Basal

medium consisted of 0.01% BSA, 8 mM mannitol, and 2 mM pyruvate in Krebs-Henseleit buffer (118.5 mM NaCl, 24.7 mM NaHCO<sub>3</sub>, 4.74 mM KCl, 1.18 mM MgSO<sub>3</sub>, 1.18 KH<sub>2</sub>PO<sub>4</sub>, and 2.5 CaCl<sub>2</sub>, pH 7.4). After basal incubation the medium was aspirated and exchanged with medium containing 600  $\mu$ U insulin/ml (100 IU/ml Actrapid; Novo Nordisk A/S, Bagsv er, Denmark) for 30 minutes. 2-deoxy-D-glucose uptake measurements were initiated by incubation in a medium containing 600  $\mu$ U insulin/ml, 1 mM 2-[2,6-<sup>3</sup>H]deoxy-D-glucose with a specific activity in the medium of 0.128 mCi/ml, and 8 mM [1-<sup>14</sup>C]mannitol with a specific activity in the medium of 0.083 mCi/ml (Amersham Biosciences Europe, Freiburg, Germany). Uptake was measured for 10 minutes before the muscle was washed in ice-cold Krebs-Henseleit buffer and quickly frozen with aluminum tongs precooled in liquid nitrogen. Frozen muscles were digested in 250  $\mu$ l 1 M NaOH at 80°C for 10 minutes. The solution was neutralized with 250  $\mu$ l 1 M NaCl and spun at 13,000 g for 2 minutes. Radioactivity in the supernatant was measured using liquid scintillation (Tri-Carb 2000; Packard Instruments, Greve, Denmark).

**Insulin release from isolated islets of Langerhans.** The insulin-secretory response of AMPK $\alpha$ 2<sup>-/-</sup>  $\beta$  cells was studied in collagenase-isolated islets of Langerhans. After isolation, islets were cultured for 16 hours in Ham's F10 medium containing 10 mM glucose. Insulin release was measured by perfusion experiments in a multiple microchamber module (Endotronics Inc., Coon Rapids, Minnesota, USA) as previously described (21). Approximately 200 islets were loaded onto Biogel P2 (Biorads Labs, Richmond, California, USA) in a col-

**Table 1**

Body composition, food intake, and triglyceride content of liver, gastrocnemius muscle, and pancreas of *AMPK $\alpha$ 2*<sup>-/-</sup> mice

	Control	<i>AMPK<math>\alpha</math>2</i> <sup>-/-</sup>
Body weight (g)	28.8 ± 0.9	29.6 ± 0.8
Fat-free mass (g) <sup>A</sup>	21.6 ± 0.6	21.7 ± 0.6
% fat mass <sup>A</sup>	7.1 ± 0.3	6.7 ± 0.2
Food intake (g/d)	3.7 ± 0.1	3.8 ± 0.1
Liver TG content (mg/g tissue)	11.2 ± 0.7	14.0 ± 3.2
Muscle TG content (mg/g tissue)	11.7 ± 2.7	8.1 ± 0.9
Pancreas TG content (mg/g tissue)	6.1 ± 1.2	5.2 ± 0.6

<sup>A</sup>Body composition was assessed by biphotonic absorptiometry. All data are expressed as mean ± SEM (*n* = 6–10 mice/group). *P* not significant between groups by unpaired, two-tailed Student *t* test. TG, triglyceride.

umn and preperfused for 20 minutes in Ham's F10 medium supplemented with 0.5% (wt/vol) BSA, 2 mM glutamine, 2 mM CaCl<sub>2</sub>, and 3 mM glucose, equilibrated with 95% O<sub>2</sub> and 5% CO<sub>2</sub>. At a flow rate of 0.5 ml/min, 15-minute pulses of 3 mM glucose were alternated with 10-minute pulses of 7.5 mM glucose, 10 mM glucose, 20 mM glucose, and 10 mM glucose with 20 mM L-arginine (Sigma-Aldrich). Samples were collected every minute and assayed for immunoreactive insulin with guinea pig anti-insulin serum (Linco Research Inc., St. Charles, Missouri, USA). Results were expressed as percentages of the insulin content measured in each individual batch of islets after finishing the experiment by sonicating the Biogel P2 containing the islets in 5 ml of 2 mM acetic acid and 0.25% (wt/vol) BSA.

**Western blot analysis.** Liver AMPK was obtained by polyethylene glycol precipitation (2), and muscle AMPK was prepared as previously described (5). AMPK catalytic subunits were detected using specific anti-AMPK $\alpha$ 1 and anti-AMPK $\alpha$ 2 antibodies (kindly provided by G. Hardie, Dundee University, Dundee, Scotland, United Kingdom). The phosphorylation status of acetyl-CoA carboxylase (ACC) was visualized using an anti-phospho-ACC (serine 79) antibody (Upstate Biotechnology Inc., Lake Placid, New York, USA).

**Statistical analysis.** Data are expressed as mean ± SEM. Statistical analyses were carried out using an unpaired, two-tailed Student *t* test, and the null hypothesis was rejected at 0.05.

## Results

**Animal model: *AMPK $\alpha$ 2*<sup>-/-</sup> mice.** *AMPK $\alpha$ 2* gene knockout was obtained using a targeting vector inserting *loxP* sites flanking the sequence containing the enzymatic catalytic site (Figure 1a). Deletion of the targeted allele was carried out by mating to EIIa-Cre transgenic mice to cause germline excision (16). Mice harboring the deleted allele were identified by

Southern blot analysis (Figure 1b). Using RT-PCR analysis, no amplification of the region corresponding to the catalytic domain of AMPK $\alpha$ 2 was observed in the liver, skeletal muscle, or pancreas of *AMPK $\alpha$ 2*<sup>-/-</sup> mice (data not shown). Western blot analysis of protein extracts prepared from *AMPK $\alpha$ 2*<sup>-/-</sup> liver and skeletal muscle showed undetectable AMPK $\alpha$ 2 protein (Figure 1c). The AMPK $\alpha$ 1 protein level was similar in the liver of *AMPK $\alpha$ 2*<sup>-/-</sup> and control mice (Figure 1c). In contrast, in *AMPK $\alpha$ 2*<sup>-/-</sup> mice, AMPK $\alpha$ 1 protein was increased 1.8-fold in gastrocnemius muscle (Figure 1c). Phosphorylation of ACC was decreased in gastrocnemius muscle 3.5-fold, while it was unchanged in the liver compared with controls (Figure 1d).

**Physical appearance, body weight, body composition, and triglyceride content in tissues of *AMPK $\alpha$ 2*<sup>-/-</sup> mice.** *AMPK $\alpha$ 2*<sup>-/-</sup> mice from heterozygous mating were born with the expected mendelian ratio. *AMPK $\alpha$ 2*<sup>-/-</sup> mice are fertile, appear indistinguishable from their wild-type littermates, and have a normal life span. No significant difference in body weight or food intake was observed between the two groups (Table 1). Body composition was similar for both groups as assessed both by biphotonic absorptiometry (Table 1) and careful dissection and weighing of individual organs (data not shown). Triglyceride content in liver, gastrocnemius muscle, and pancreas was similar in *AMPK $\alpha$ 2*<sup>-/-</sup> mice and controls (Table 1).

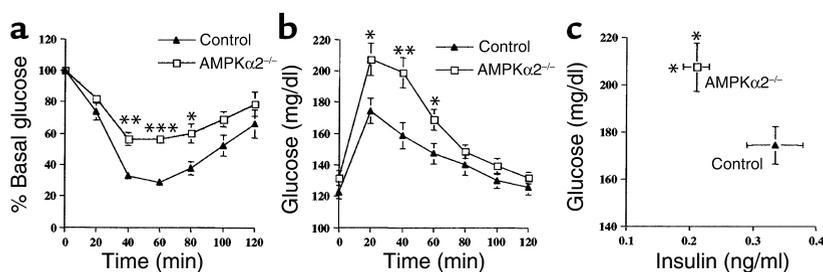
**Metabolic parameters.** In fasted conditions, plasma glucose and insulin levels were similar in *AMPK $\alpha$ 2*<sup>-/-</sup> mice and controls, while plasma FFA concentrations were increased in *AMPK $\alpha$ 2*<sup>-/-</sup> mice (Table 2). Glycerol plasma levels have a tendency to be higher in *AMPK $\alpha$ 2*<sup>-/-</sup> mice but do not reach statistical significance (Table 2). In the fed state, *AMPK $\alpha$ 2*<sup>-/-</sup> mice were characterized by higher plasma glucose and plasma FFA concentrations than were found in control mice, while plasma insulin and leptin levels were lower (Table 2). No difference between plasma levels of total and HDL cholesterol, sodium, potassium, creatinine, or urea were noted between *AMPK $\alpha$ 2*<sup>-/-</sup> and control mice (Table 2 and data not shown).

**Table 2**

Metabolic parameters of *AMPK $\alpha$ 2*<sup>-/-</sup> mice in fasted and fed periods

	Fasted		Fed	
	Control	<i>AMPK<math>\alpha</math>2</i> <sup>-/-</sup>	Control	<i>AMPK<math>\alpha</math>2</i> <sup>-/-</sup>
Glycemia (mg/dl)	102 ± 5	101 ± 13	141 ± 10	171 ± 9 <sup>B</sup>
Insulin (ng/ml)	0.46 ± 0.17	0.29 ± 0.08	1.35 ± 0.07	1.07 ± 0.12 <sup>B</sup>
Leptin (ng/ml)	0.38 ± 0.10	0.38 ± 0.13	4.31 ± 0.24	2.65 ± 0.49 <sup>A</sup>
FFA (mM)	1.10 ± 0.04	1.37 ± 0.06 <sup>A</sup>	0.73 ± 0.06	0.95 ± 0.07 <sup>B</sup>
Glycerol (μM)	367 ± 14	388 ± 15	281 ± 24	312 ± 31
TG (mM)	0.58 ± 0.02	0.74 ± 0.07	0.72 ± 0.05	0.91 ± 0.13
β-hydroxybutyrate (mM)	0.98 ± 0.09	1.17 ± 0.11	0.07 ± 0.01	0.08 ± 0.01
Total cholesterol (mM)	2.3 ± 0.2	2.9 ± 0.1	2.3 ± 0.2	2.6 ± 0.3
HDL cholesterol (mM)	1.7 ± 0.1	1.9 ± 0.2	1.5 ± 0.3	1.7 ± 0.2

All data are expressed as mean ± SEM (*n* = 6–10 mice/group). <sup>A</sup>*P* < 0.01, <sup>B</sup>*P* < 0.05 vs. control group by unpaired, two-tailed Student *t* test.



**Figure 2**

*AMPKα2<sup>-/-</sup> mice are AICAR-resistant, glucose intolerant, and exhibit impaired glucose-stimulated insulin secretion. (a) AICAR tolerance test. (b) OGTT. (c) Plasma glucose and insulin levels at time 20 minutes during OGTT. All results are expressed as mean ± SEM (n = 6–10). \*P < 0.05, \*\*P < 0.001, \*\*\*P < 0.0001 vs. control group by unpaired, two-tailed Student t test. NS, not significant.*

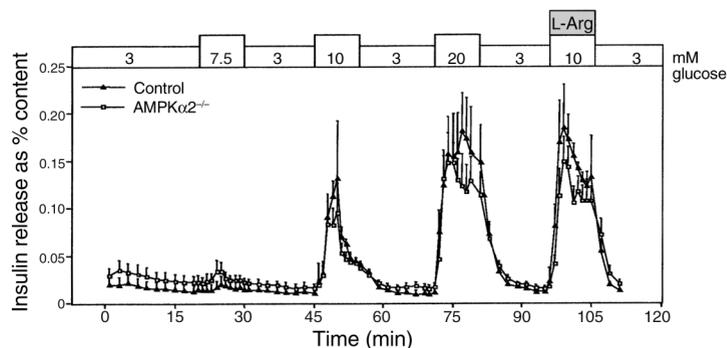
*AMPKα2<sup>-/-</sup> mice are AICAR-resistant, glucose intolerant, and have reduced glucose-stimulated insulin secretion.* To investigate in vivo to what extent remaining AMPKα1 protein levels can substitute for AMPKα2, we performed an AICAR tolerance test. The relative insensitivity to hypoglycemia-mediated AICAR effects observed in *AMPKα2<sup>-/-</sup>* mice (Figure 2a) suggests that AMPKα1 cannot fully compensate for the lack of AMPKα2 in *AMPKα2<sup>-/-</sup>* mice. To investigate the dynamic response of *AMPKα2<sup>-/-</sup>* mice to increased glucose concentrations, an oral glucose challenge was performed. Following glucose absorption, *AMPKα2<sup>-/-</sup>* mice have a markedly increased blood glucose excursion, demonstrating glucose intolerance (Figure 2b). Insulinemia determined at time 20 minutes was significantly lower in *AMPKα2<sup>-/-</sup>* mice (Figure 2c).

*Glucose- and L-arginine-stimulated insulin secretion are normal in isolated pancreatic islets of AMPKα2<sup>-/-</sup> mice.* No significant difference in insulin release was observed between control and *AMPKα2<sup>-/-</sup>* pancreatic islets for basal (3 mM glucose) and stimulatory glucose concentrations (7.5 mM, 10 mM, and 20 mM glucose) or when challenged with 20 mM L-arginine (Figure 3). Furthermore, the insulin content of *AMPKα2<sup>-/-</sup>* and control islets was similar (67.2 ± 16.2 vs. 63.9 ± 16.5 ng/islet in controls; P not significant).

*AMPKα2<sup>-/-</sup> mice are insulin resistant while Tg-KD-AMPKα2 mice are not.* In the basal condition, endogenous glucose production, glycolysis rate, and whole-body glycogen synthesis rate were similar between *AMPKα2<sup>-/-</sup>* and control mice (Table 3). In vivo muscle glycogen synthesis rate and total glycogen content were similar in both groups (Table 3). During hyperinsulinemic

**Figure 3**

*Glucose- and L-arginine-stimulated (L-Arg) insulin secretion in isolated AMPKα2<sup>-/-</sup> islets. All results are expressed as mean ± SEM (n = 4). P not significant between groups by unpaired, two-tailed Student t test.*



euglycemic clamp, plasma glucose and insulinemia were comparable in *AMPKα2<sup>-/-</sup>* and control mice (plasma glucose: 109.8 ± 7.2 mg/dl in *AMPKα2<sup>-/-</sup>* mice vs. 97.2 ± 3.6 mg/dl in controls, P not significant; plasma insulin: 159.7 ± 22.6 mU/l in *AMPKα2<sup>-/-</sup>* mice vs. 146.9 ± 22.5 mU/l in controls, P not significant). Insulin-stimulated whole-body glucose turnover rate was decreased in *AMPKα2<sup>-/-</sup>* mice compared with controls (Table 3). Insulin-stimulated whole-body glycolysis was similar in both groups, while the insulin-stimulated whole-body glycogen synthesis rate was dramati-

cally decreased in *AMPKα2<sup>-/-</sup>* mice (Table 3). Muscle glycogen synthesis rate and total muscle glycogen content were severely reduced in *AMPKα2<sup>-/-</sup>* mice compared with controls at the end of the clamp (Table 3). Because glucose uptake is largely rate limiting for glycogen synthesis in muscle during clamp study, we assayed uptake of deoxyglucose into isolated muscle in the presence of insulin. Basal and insulin-stimulated glucose uptake were similar in isolated extensor digitorum longus and soleus muscles from controls and *AMPKα2<sup>-/-</sup>* mice (Figure 4, a and b). In addition, total glycogen content in skeletal muscle of random-fed animals was similar between *AMPKα2<sup>-/-</sup>* and control mice (0.312 ± 0.106 vs. 0.358 ± 0.072 μg/mg in controls, respectively, P not significant). Lastly, endogenous hepatic glucose production was totally inhibited in both groups, suggesting no hepatic insulin resistance (data not shown). Furthermore, no defect was detected in glycogen synthesis or total glycogen content in the liver of *AMPKα2<sup>-/-</sup>* and control mice in either basal or clamp studies (data not shown).

To further investigate the role of muscle AMPK in the development of insulin resistance, we performed hyperinsulinemic euglycemic clamps in transgenic mice expressing a kinase-dead mutant of AMPK in skeletal muscle (Tg-KD-AMPKα2 mice) and their respective nontransgenic controls (17). Oral glucose tolerance was comparable between the two groups (data not shown). During the clamp, plasma glucose

**Table 3**Metabolic parameters of *AMPK $\alpha$ 2<sup>-/-</sup>* and Tg-KD-*AMPK $\alpha$ 2* mice during basal and hyperinsulinemic euglycemic clamp studies

	Weight (g)	Whole-body glucose turnover (mg/kg/min)	Whole-body glycolysis rate (mg/kg/min)	Whole-body glycogen synthesis rate (mg/kg/min)	Muscle glycogen synthesis rate (ng/mg/min)	Muscle glycogen content ( $\mu$ g/mg)
<b>Basal</b>						
Control	24.7 $\pm$ 1.2	24.2 $\pm$ 1.6	10.3 $\pm$ 0.5	13.9 $\pm$ 1.5	48.4 $\pm$ 14.6	0.035 $\pm$ 0.013
<i>AMPK<math>\alpha</math>2<sup>-/-</sup></i>	24.7 $\pm$ 1.0	25.6 $\pm$ 1.9	9.2 $\pm$ 0.7	16.4 $\pm$ 2.0	15.7 $\pm$ 4.1	0.020 $\pm$ 0.006
<b>Clamp</b>						
Control	22.3 $\pm$ 0.5	90.6 $\pm$ 4.3	19.7 $\pm$ 2.9	71.0 $\pm$ 2.5	611.8 $\pm$ 143.2	0.458 $\pm$ 0.113
<i>AMPK<math>\alpha</math>2<sup>-/-</sup></i>	21.4 $\pm$ 0.5	68.7 $\pm$ 2.9 <sup>A</sup>	21.8 $\pm$ 1.6	46.9 $\pm$ 2.9 <sup>A</sup>	143.9 $\pm$ 17.0 <sup>A</sup>	0.137 $\pm$ 0.030 <sup>B</sup>
NTg	27.2 $\pm$ 0.5	64.2 $\pm$ 7.0	14.8 $\pm$ 1.0	45.9 $\pm$ 6.9	108.9 $\pm$ 50.9	0.175 $\pm$ 0.053
Tg-KD- <i>AMPK<math>\alpha</math>2</i>	25.1 $\pm$ 1.3	72.9 $\pm$ 5.3	16.5 $\pm$ 3.6	56.4 $\pm$ 7.3	89.9 $\pm$ 21.0	0.172 $\pm$ 0.050

All data are expressed as mean  $\pm$  SEM ( $n = 6-12$  mice/group). <sup>A</sup> $P < 0.01$ , <sup>B</sup> $P < 0.05$  vs. control group by unpaired, two-tailed Student *t* test. NTg, nontransgenic.

and insulinemia were comparable in Tg-KD-*AMPK $\alpha$ 2* and nontransgenic mice (plasma glucose: 93.6  $\pm$  1.8 mg/dl in Tg-KD-*AMPK $\alpha$ 2* mice vs. 95.4  $\pm$  9.0 mg/dl in nontransgenic controls, *P* not significant; plasma insulin: 113.7  $\pm$  24.8 mU/l in Tg-KD-*AMPK $\alpha$ 2* mice vs. 145.7  $\pm$  22.9 mU/l in nontransgenic controls, *P* not significant). Insulin-stimulated whole-body glucose turnover, whole-body glycogen synthesis rate, muscle glycogen synthesis rate, and total muscle glycogen content were similar in both groups (Table 3).

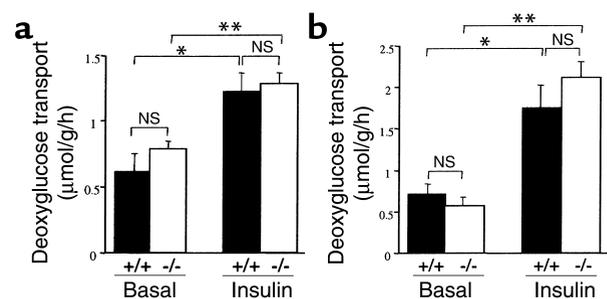
*Increased daily urinary catecholamine excretion in AMPK $\alpha$ 2<sup>-/-</sup> mice.* Daily epinephrine, norepinephrine, and dopamine excretion were significantly higher in *AMPK $\alpha$ 2<sup>-/-</sup>* mice than in controls (Figure 5, a-c). The response to a glucose challenge was dramatically improved in *AMPK $\alpha$ 2<sup>-/-</sup>* mice after treatment with  $\alpha$ -adrenergic blockers, while no change was observed in control mice (Figure 5, d and e). In contrast,  $\beta$ -adrenergic blockers did not modify glucose tolerance of *AMPK $\alpha$ 2<sup>-/-</sup>* mice (Figure 5, d and e).

## Discussion

The primary goal of this study was to assess the role of the *AMPK $\alpha$ 2* isoform in controlling glucose metabolism in vivo. *AMPK $\alpha$ 2<sup>-/-</sup>* mice appear indistinguishable from their control littermates. Importantly, there is no difference in body composition, adiposity, or food intake between mutant and control mice. The most striking observation is reduced glucose tolerance in *AMPK $\alpha$ 2<sup>-/-</sup>* mice. We showed that this defect is linked with both reduced insulin release and decreased insulin sensitivity of peripheral tissues. Few data are available concerning regulation of insulin release by AMPK. Activation of AMPK has been shown to inhibit insulin secretion in vitro (9, 10). Here, we showed that while glucose-induced insulin secretion is clearly impaired in *AMPK $\alpha$ 2<sup>-/-</sup>* mice in vivo, basal and glucose- and L-arginine-stimulated insulin secretion in isolated pancreatic islets are similar in *AMPK $\alpha$ 2<sup>-/-</sup>* mice and controls. Furthermore, insulin content of islets and insulin gene transcription (data not shown) were not affected by *AMPK $\alpha$ 2* gene deletion. Taken together, these data indicate that the lack of insulin secretion observed in

vivo in *AMPK $\alpha$ 2<sup>-/-</sup>* mice is not due to a primary defect in  $\beta$  cell function. Hypokalemia or hyperleptinemia (22), known to modulate insulin secretion negatively in vivo, are not observed in *AMPK $\alpha$ 2<sup>-/-</sup>* mice. In addition, lipid content in pancreas was similar in *AMPK $\alpha$ 2<sup>-/-</sup>* mice and controls, excluding a deleterious effect on insulin release by chronic accumulation of lipids (23, 24). An important regulator of insulin secretion is the fine-tuning of the balance between sympathetic and parasympathetic activity. This is critical for the  $\beta$  cells, since epinephrine and norepinephrine directly and potentially suppress  $\beta$  cell function (25). *AMPK $\alpha$ 2<sup>-/-</sup>* mice exhibited higher catecholamine urinary excretion than did controls. Since catecholamines are known to inhibit insulin release, dysregulation in sympathetic tone observed in *AMPK $\alpha$ 2<sup>-/-</sup>* mice could be implicated in the reduction of insulin secretion. More importantly, no defect in glucose and insulin plasma levels was observed in mutant mice in the fasted state, suggesting that dysregulation of sympathetic tone is predominant in the fed period.

Glucose intolerance is due to both impaired insulin secretion and reduced insulin-stimulated glucose utilization. Therefore, we examined insulin sensitivity in peripheral tissues of *AMPK $\alpha$ 2<sup>-/-</sup>* mice using hyperinsu-

**Figure 4**

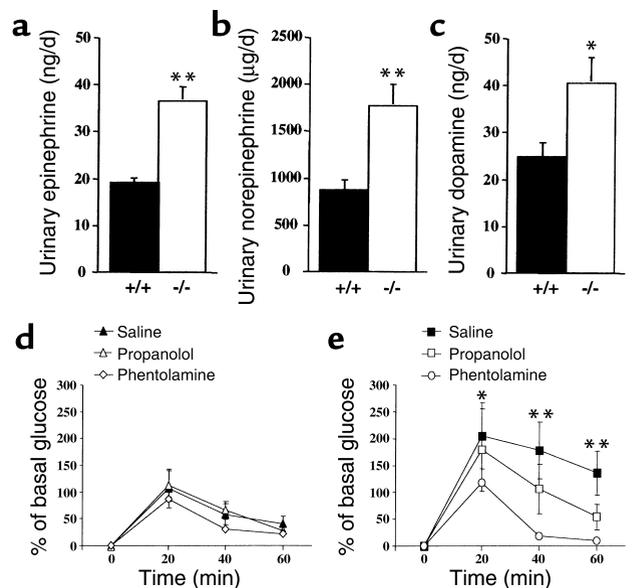
Insulin-stimulated glucose transport in isolated muscles from *AMPK $\alpha$ 2<sup>-/-</sup>* mice. Basal and insulin-stimulated glucose transport in (a) soleus and (b) extensor digitorum longus muscles from control and *AMPK $\alpha$ 2<sup>-/-</sup>* mice. All results are expressed as mean  $\pm$  SEM ( $n = 4-7$ ). \* $P < 0.05$ , \*\* $P < 0.01$  vs. basal conditions by unpaired, two-tailed Student *t* test; *P* not significant (NS) vs. control group.

linemic euglycemic clamp studies. To our surprise, *AMPK $\alpha$ 2*<sup>-/-</sup> mice displayed decreased whole-body insulin sensitivity, with a dramatic reduction in insulin-stimulated glycogen synthesis in skeletal muscle, while hepatic insulin sensitivity was preserved. Does  $\alpha$ 2 deletion in skeletal muscle directly alter insulin sensitivity? This does not seem to be the case since insulin sensitivity is unaltered in Tg-KD-*AMPK $\alpha$ 2* mice, which express a dominant-negative mutant of AMPK in skeletal muscle (17). This points out that lack of AMPK activity in skeletal muscle is unable by itself to alter insulin sensitivity or glycogen synthesis in this tissue. Likewise, comparable muscle glycogen contents of *AMPK $\alpha$ 2*<sup>-/-</sup> and control mice during the random-fed period suggest that the balance between degradation and repletion of glycogen in muscle is intact in this model. In consequence, a defect in muscular glycogen synthesis was specifically observed in *AMPK $\alpha$ 2*<sup>-/-</sup> mice when insulinemia reached supra-physiological levels. Glucose uptake is largely rate limiting for glycogen synthesis in muscle during clamp study. As insulin induced a similar increase in deoxyglucose uptake in incubated extensor digitorum longus and soleus muscles, it is unlikely that primary defective glucose transport in muscle contributes to *AMPK $\alpha$ 2*<sup>-/-</sup> insulin resistance.

Another known important mechanism of insulin resistance in skeletal muscle is increased lipid availability or increased tissue uptake of FFA (26). AMPK regulates lipid oxidation in muscle by reducing ACC activity, which in turn decreases malonyl coenzyme A levels and increases mitochondrial uptake of FFAs. In *AMPK $\alpha$ 2*<sup>-/-</sup> gastrocnemius muscle, we showed that  $\alpha$ 1 protein was increased by 1.8-fold (Figure 1c) but ACC phosphorylation was reduced by 3.5-fold (Figure 1d). These data suggest that in skeletal muscle, the increased  $\alpha$ 1 protein levels in *AMPK $\alpha$ 2*<sup>-/-</sup> mice is not sufficient to ensure normal phosphorylation of ACC. However, the 24-hour respiratory quotient, monitored by indirect calorimetry, was similar in *AMPK $\alpha$ 2*<sup>-/-</sup> and control mice (data not shown), suggesting that whole-body lipid oxidation is not altered by the lack of the AMPK $\alpha$ 2 subunit. Furthermore, comparable skeletal muscle lipid content in both groups argues against any alteration of lipid oxidation rate in *AMPK $\alpha$ 2*<sup>-/-</sup> mice. Finally, we know that adipose tissue operates as an endocrine organ that releases a large number of hormones in response to specific extracellular stimuli or changes in metabolic status. Alteration of secretion of these proteins (which include TNF- $\alpha$ , leptin, adiponectin, resistin, and adiponectin) as a possible mechanism of *AMPK $\alpha$ 2*<sup>-/-</sup> mouse insulin resistance needs further investigation.

So how can we explain insulin resistance in *AMPK $\alpha$ 2*<sup>-/-</sup> mice? The observation that *AMPK $\alpha$ 2*<sup>-/-</sup> mice exhibit altered glucose metabolism in vivo but not in isolated organs suggests that the defect is of central origin. It is known that integration of peripheral and central metabolic signals in the CNS during different metabolic situations such as fasting, fed state, and hyperinsulinemic

clamp activates the autonomous nervous system, which in turn regulates glucose metabolism in peripheral tissues (27–30). As AMPK $\alpha$ 2 is widely expressed in the brain (13), we hypothesized that lack of AMPK $\alpha$ 2 in neurons could contribute to the metabolic defects observed in *AMPK $\alpha$ 2*<sup>-/-</sup> mice. Supporting this hypothesis, *AMPK $\alpha$ 2*<sup>-/-</sup> mice exhibited a significant increase in daily urinary catecholamine excretion, suggesting higher sympathetic activation in these mice. Furthermore, the glucose intolerance of *AMPK $\alpha$ 2*<sup>-/-</sup> mice was dramatically improved after intraperitoneal injection of phentolamine (an  $\alpha$ -adrenergic receptor antagonist), while no change was observed after treatment of *AMPK $\alpha$ 2*<sup>-/-</sup> mice by propranolol (a  $\beta$ -adrenergic receptor antagonist). This suggested that the  $\alpha$ -adrenergic but not the  $\beta$ -adrenergic pathway is involved in the glucose intolerance observed in *AMPK $\alpha$ 2*<sup>-/-</sup> mice. It is well known that epinephrine overproduction inhibits insulin secretion, reduces insulin-stimulated glucose uptake in skeletal muscles, and alters muscle glycogen synthesis (31–35); all of these defects are present in *AMPK $\alpha$ 2*<sup>-/-</sup> mice in vivo. Moreover, normality of some metabolic parameters (fasting glucose and insulin plasma levels or skeletal muscle glycogen content in random-fed animals) suggests that exaggerated sympathetic tone in *AMPK $\alpha$ 2*<sup>-/-</sup> mice is not permanent but is restricted to particular nutritional situations such as glucose challenge and hyperinsulinemic euglycemic clamps.



**Figure 5** Increased daily urinary catecholamine excretion in *AMPK $\alpha$ 2*<sup>-/-</sup> mice. Daily urinary (a) epinephrine, (b) norepinephrine, and (c) dopamine excretion in *AMPK $\alpha$ 2*<sup>-/-</sup> mice. All results are expressed as mean  $\pm$  SEM ( $n = 6$ ). \* $P < 0.05$ , \*\* $P < 0.01$  vs. control group by unpaired, two-tailed  $t$  test. Also shown are effect of phentolamine and propranolol,  $\alpha$ - and  $\beta$ -adrenergic blockers, respectively, on the response of glucose challenge of control (d) and *AMPK $\alpha$ 2*<sup>-/-</sup> (e) mice. All results are expressed as mean  $\pm$  SEM ( $n = 5$ ). \* $P < 0.05$ , \*\* $P < 0.01$ , saline- vs. phentolamine-treated group by unpaired, two-tailed Student  $t$  test;  $P$  not significant for saline- vs. propranolol-treated group.

The data presented here identify AMPK $\alpha$ 2 as a major component in the control of insulin sensitivity and insulin secretion in vivo. We clearly demonstrated that the metabolic function of AMPK $\alpha$ 2<sup>-/-</sup> isolated skeletal muscle and pancreatic islets is normal, suggesting that the origin of the defects observed in vivo is located outside these tissues. In addition, we showed significant increased catecholamine excretion in AMPK $\alpha$ 2<sup>-/-</sup> mice. Thus, we hypothesize that lack of AMPK $\alpha$ 2 in neurons reduces the ability to integrate peripheral metabolic signals into the brain and consequently alters the control of peripheral insulin sensitivity and insulin secretion by increasing sympathetic nervous activity.

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