E3 Ubiquitin Ligase, WWP1, Interacts with AMPK2 and Down-regulates Its Expression in Skeletal Muscle C2C12 Cells*

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Background: The role of the ubiquitin-proteasome pathway under high glucose conditions is unclear. **Results:** AMPK α 2 interacts with WWP1, and its expression is down-regulated by the ubiquitin proteasome pathway in high glucose culture conditions in C2C12 cells.

Conclusion: WWP1 down-regulates ΑΜΡΚα2 expression through direct interaction in high glucose culture conditions of skeletal muscle C2C12 cells.

Significance: The ubiquitin proteasome pathway may involve high glucose-induced AMPK α 2 down-regulation.

It is known that the activity of AMP-activated protein kinase (AMPK2) was depressed under high glucose conditions. However, whether protein expression of AMPK2 is also down-regulated or not remains unclear. In this study, we showed that the expression of AMPK2 was down-regulated in cells cultured under high glucose conditions. Treatment of proteasome inhibitor, MG132, blocked high glucose-induced AMPKα2 down**regulation. Endogenous AMPK2 ubiquitination was detected by immunoprecipitation of AMPK2 followed by immunoblotting detection of ubiquitin. The yeast-two hybrid (YTH) approach identified WWP1, an E3 ubiquitin ligase, as the AMPK2-interacting protein in skeletal muscle cells. Interaction between AMPK2 and WWP1 was validated by co-immunoprecipitation. Knockdown of WWP1 blocked high glucoseinduced AMPK2 down-regulation. The overexpression of WWP1 down-regulated AMPK2. In addition, the expression of WWP1 is increased under high glucose culture conditions in both mRNA and protein levels. The level of AMPK2 was downregulated in the quadriceps muscle of diabetic animal model db/db mice. Expression of WWP1 blocked metformin-induced glucose uptake. Taken together, our results demonstrated that WWP1 down-regulated AMPK2 under high glucose culture conditions via the ubiquitin-proteasome pathway.**

The metabolic syndrome is a combination of metabolic risk factors, including obesity, insulin resistance, hyperglycemia, and dyslipidemia, and it increases the risk of developing cardiovascular disease and diabetes. Maintaining energy balance depends on the efficiency of tightly regulated mechanisms of energy intake and expenditure. Hyperglycemia is a condition in which an excess amount of glucose circulates in the blood

plasma. Glucose homeostasis is maintained by a balance between hepatic glucose production and glucose uptake by peripheral tissues. Activation of AMP-activated protein kinase $(AMPK)^2$ enhances glucose uptake through the translocation of the glucose transporter 4 (GLUT4) to the cell membrane and also through regulation of Glut4 gene expression (1–3). The AMPK activator enhances glucose uptake into muscle by increasing cell surface $GLUT4$ levels $(4-6)$. AMPK may regulate GLUT4 endocytosis (7, 8) and GLUT4 translocation (9, 10). Thus, AMPK is a well-accepted therapeutic target for the metabolic syndrome and type 2 diabetes (11, 12).

AMPK is an enzyme that plays a role in cellular energy homeostasis. AMPK, a heterotrimeric complex comprised of a catalytic subunit and two regulatory subunits, is activated when cellular energy is depleted (13). Activated AMPK switches on ATP-generating catabolic pathways, including glucose and fatty acid oxidation (14–16), while it simultaneously switches off ATP-consuming anabolic pathways (cholesterol, fatty acid, and triacylglycerol synthesis) (17). Thus, AMPK plays as a counter-regulatory mechanism and restores the AMP:ATP ratio. In addition to its roles in energy homeostasis, AMPK has also been shown to be regulated by ubiquitination (18). This fact suggests that the expression level of AMPK may play an important role in in specific biological conditions; however, the molecular link by which ubiquitination acts in high glucose conditions is obscure. It is reported that elevated glucose downregulated the activity of AMPK in pancreatic cells (19) and rat skeletal muscle (20).

Protein ubiquitination is a mechanism for targeting proteins for rapid proteasomal degradation (21). Ubiquitin is attached covalently to target proteins in a series of reactions driven by ubiquitin ligase. E1 ligases load ubiquitin onto E2 ligases, which in turn modify target proteins. Substrate specificity is generated by E3 ligases, which serve as adaptors. Recent studies demon-

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 2 The abbreviations used are: AMPK, AMP-activated protein kinase; AICAR, 5-aminoimidazole-4-carboxy-amide-1-D-ribofuranoside; ACC, acetyl-CoA carboxylase; ACO, acyl-CoA oxidase; GLUT4, glucose transporter 4, YTH, yeast two hybrid; Cide, cell death-inducing DFFA-like effector.

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strated that the ubiquitin proteasome system may regulate AMPK activity. The de-ubiquitination enzyme USP9X interacts with NUAK1 (AMPK-related kinase 5) (18), implicating a role of AMPK in ubiquitination. This group identified Cidea (cell death-inducing DFFA-like effector a) as an E3 ubiquitin ligase for AMPK (22). Although the basic regulation of AMPK activity by kinases has been delineated, regulation by posttranslational modifications such as ubiquitination remains to be determined.

In this study, we found that $AMPK\alpha2$ directly interacted with E3 ubiquitin ligase WWP1. We also showed that the expression level of AMPK α 2 was down-regulated in C2C12 cells cultured under high glucose conditions, and further demonstrated that the ubiquitination is involved in high glucose-induced $AMPK\alpha2$ down-regulation. These findings provide novel insights into the manner in which high glucose contributes to insulin resistance through down-regulation of $AMPK\alpha2$ in skeletal muscle cells.

MATERIALS AND METHODS

 $$ Cell Signaling Technology (New England Biolabs, Beverly, MA). Anti-WWP1 antibody was purchased from ProteinTech Group, Inc. (Chicago, IL). Anti-ubiquitin antibody, anti-Flag antibody, anti- β -actin antibody, and AICAR were purchased from Cell Signaling Technology (New England Biolabs). Horseradish peroxidase-conjugated secondary antibodies were obtained from Assay Designs and Stressgen (Ann Arbor, MI). Metformin, MG132, and lactacystin were obtained from Calbiochem (San Diego, CA). L -(-)Glucose, D -(+)glucose, and mannitol were purchased from Sigma.

Cell Culture and Differentiation—Mouse myoblast C2C12 and human embryonic kidney (HEK) 293 cells were maintained in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% fetal bovine serum (FBS) and antibiotics at 37 °C in an incubator with 5% $CO₂$. Cells were grown in culture medium consisting of 500 μ l of DMEM (GIBCOTM, Auckland, NZ) containing 0.584 g/liter of L-glutamate and 4.5 g/liter of glucose, and mixed with 500 ml of F-12 medium containing 0.146 g/liter of L-glutamate, 1.8 g/liter of glucose, 100 μ g/ml of gentamicin, 2.5 g/liter of sodium carbonate, and 10% heat-inactivated FBS. L6 myoblasts were maintained in cell monolayers in α -MEM supplemented with 10% FBS, 50 units of penicillin per ml, and 50 μ g of streptomycin per ml in a 5% CO₂ humidified atmosphere at 37 °C. Two days after the myoblasts achieved confluence, differentiation to myotubes was induced by incubating the cells for 6~7 days in α -MEM supplemented with 2% FBS, which was changed every 2 days. The monolayer myotubes were finally serum-starved in DMEM for 2 h, then used in the following experiments.

Animal Tissue—Male C57BLKS/J-db/db mice and wild-type C57BL/6J mice were obtained from the Central Lab. Animal Inc. (Seoul, Korea) at age 6 weeks. All mice were maintained under standard light (12 hours light/dark), temperature (22 \pm 2 °C), and humidity (40 \pm 1 0%) conditions. The mice were anesthetized with dry ice, and quadriceps muscle was prepared. All procedures were approved by the Institutional Animal Care and Utilization Committee for Medical Science of Korea University.

Immunoblot Analysis—Cells were grown on 6-well plates and serum-starved for 24 h prior to treatment with the indicated agents. Following treatment of the cells, the media was aspirated, and the cells were washed twice in ice-cold PBS and lysed in 100 μ l of lysis buffer. The samples were then briefly sonicated, heated for 5 min at 95 °C, and centrifuged for 5 min. The supernatants were electrophoresed on SDS-PAGE (8%) gels, and transferred to polyvinylidene difluoride membranes. The blots were incubated overnight at room temperature with primary antibodies and then washed six times in Tris-buffered saline/0.1% Tween 20 prior to 1 h incubation with horseradish peroxidase-conjugated secondary antibodies at room temperature. The blots were then visualized via ECL (Amersham Biosciences, Buckinghamshire, UK). In some cases, the blots were stripped and reprobed using other antibodies.

Immunoprecipitation—The amount of proteins from C212 cells or Flag-WWP1-transfected HEK293 cells was determined by the Bradford method. Cellular protein $(100 \mu g)$ was mixed with 1μ g of anti-Flag or anti-AMPK antibodies and incubated overnight at 4° C. Then, 10 μ l of protein A-Sepharose (Amersham, Uppsala, Sweden) was added to these samples and incubated for another 3 h at 4 °C. After the incubation, samples were washed three times with wash buffer (25 mm HEPES, 5 mm EDTA, 1% Triton X-100, 50 mM NaF, 150 mM NaCl, 10 mM phenylmethylsulfonyl fluoride, 1μ M leupeptin, 1μ M pepstatin, and 1 μ M aprotinin A, pH 7.2). The washed samples were resuspended in SDS sample buffer (125 mm Tris-HCl, pH 6.8, 20% [v/v] glycerol, 4% [w/v] SDS, 100 mm dithiothreitol, and 0.1% [w/v] bromphenol blue), and heated at 100 °C for 5 min prior to electrophoresis.

Silencing AMPK2 and WWP1—C2C12 cells were seeded in 6-well plates and allowed to grow to 70% confluence for 24 h. Transient transfections were performed with transfection reagent (Lipofectamine 2000; Invitrogen) according to the manufacturer's protocol. Briefly, both $AMPK\alpha2$ siRNA (NM_001013367; Dharmacon, CO), WWP1 siRNA (Genolution Pharmaceuticals, Inc.) and non-targeted control siRNAs (Non-targeting pool; Dharmacon CO) were designed. 5 μ l of siRNA and 5 μ l of transfection reagent (Lipofectamine 2000) were each diluted with 95 μ l of reduced serum media (Opti-MEM; Invitrogen), then mixed. The mixtures were allowed to incubate for 30 min at room temperature and then were added dropwise to each culture well containing 800 μ l of reduced serum media (Opti-MEM; final siRNA concentration, 100 nm). Four hours after transfection, the medium was changed with fresh complete medium. The cells were cultivated for 24 h and lysed, and expression of $AMPK\alpha2$ protein was assayed with Western blotting.

Immunohistochemistry—Seven micrometer-thick cryosections from db/db and control mouse quadriceps femoris muscle were fixed with 4% freshly depolymerized paraformaldehyde for 15 min at 4 °C. After blocking with normal goat serum, sections were immunostained with antibodies against $AMPK\alpha$ for overnight at room temperature. After extensive washing with phosphate-buffered saline (PBS), sections were incubated with FITC-labeled secondary antibody.

Reverse Transcription-Polymerase Chain Reaction (RT-PCR)— First-strand cDNA synthesis was performed using 1μ g of total RNA isolated from C2C12 at 55 °C for 20 min using the Thermoscript II one-step RT-PCR Kit (Invitrogen, Paisly, UK). Amplification of cDNA was carried out in the same tube using the Gene Amp System 9700 thermocycler (Applied Biosystems, Warrington, UK). Heating to 94 °C for 5 min inactivated the reverse transcriptase. The following PCR conditions were used: 27 cycles of 30 s at 94 °C, 30 s at 56 °C, and 30 s at 72 °C, followed by 7 min at 72 °C. The number of PCR cycles used was optimized to ensure amplification at the exponential phase. $10-\mu$ l samples from each RT-PCR product were removed and analyzed by agarose gel electrophoresis. Bands were stained with ethidium bromide and visualized under ultraviolet light. Band intensity quantification was determined by a gel documentation system (Gene Genius, Syngene, UK). The following primers were used: WWP1-sense (5-TCC TGG TGA CGG AAG AAA AC-3) and WWP1-antisense (5-AAC CAG GTG TCT TTG CCA AC-3'); and $AMPK\alpha2$ -sense (5'-CGC CTC TAG TCC TCC ATC AG-3') and AMPK α 2-antisense (5'-CAG CTG TGC TGG AAT CAA AA-3'); and β -actin-sense (5'-CAG GAG GAG CAA TGA TCT TGA-3') and β -actin antisense (5-ACT ACC TCA TGA AGA TCC TCA-3).

GST Pull-down Assay—GST fusion proteins containing different domains of $AMPK\alpha2$ were engineered by PCR. The fusion proteins were expressed in *Escherichia coli* bacteria by incubation with 0.2 mm isopropyl β -D-galactopyranoside for 6 h at 37 °C. The bacteria were lysed by sonication, and the fusion proteins were purified with glutathione-agarose beads with buffer A followed by immunoblotting with anti-WWP1 antibody.

Glucose Uptake—Glucose uptake was measured using the 2-deoxy-D-[³H]glucose. Briefly, L6 cells were serum-starved overnight before each experiment, and glucose uptake was measured on day 7. Medium was replaced in the morning (serum-free medium), and cells were exposed to drugs for 1 h. Cells were washed twice in warm PBS before media and drugs were placed in DMEM devoid of glucose for 20 min. 2-Deoxy-D-[³H]glucose (50 nM) was added for 15 min at 37 °C, and the reactions were terminated by washing twice in ice-cold PBS. Cells were lysed in 10% SDS or 50 mM NaOH, and samples were transferred to scintillation vials with scintillant and allowed to sit at room temperature for 1 h before being counted.

Plasmid Construction of Flag-wwp1—The open reading frames of human WWP1 was cloned into the Pcmv-Tag2C vector (Stratagene) between EcoRI and SalI sites. cDNA from Hela cell line was amplified by PCR (forward primer, 5'-C GGA ATT CGA ATG GCC ACT GCT TCA CCA AG-3; reverse primer 5-A CGC GTC GAC TCA TTC TTG TCC AAA TCC CTC TG-3). Kpn1- and Sac1-digested products were ligated into a linearized pCMV-Tag2C vector. The construct was verified by direct sequencing.

Data Analysis—Data are expressed as the means \pm S.E. Image Gauge (version 3.12; Fujifilm, Tokyo, Japan) was used for analysis of band intensity. One-way ANOVA was used followed by a Holm-Sidak multiple-range test for comparison between groups. p values ≤ 0.05 were considered statistically significant.

FIGURE 1. **AMPK2 down-regulated in C2C12 cells cultured under high glucose conditions.** *A*, C2C12 cells were stimulated for the indicated agents, such as D-glucose (30 mm), L-glucose (30 mm), and mannitol (30 mm) for 6 days. The cell lysates (20 μ g) were analyzed via Western blotting for anti-AMPK α 2 antibody. The anti-β-actin antibody was a protein loading control. *B*, C2C12 cells were stimulated for the indicated times with 30 mm high glucose. The cell lysates (20 μ g) were analyzed via Western blotting for anti-AMPK α 2 antibody. The anti- β -actin antibody was a protein loading control.

RESULTS

AMPK2 Down-regulated in C2C12 Cells Cultured under High Glucose Conditions—High glucose concentrations are known to have detrimental effects on many cells types. To gain insight into the effect of high glucose conditions on skeletal muscle system, we evaluated the effect of high glucose concentration on the expression of $AMPK\alpha2$ in skeletal muscle C2C12 cells. The expression of $AMPK\alpha2$ significantly down regulated in high glucose concentration in cell culture medium. Mannitol is used as the isotonic control. L-Glucose is used as the D-glucose specificity control. The level of $AMPK\alpha2$ was not affected by these two agents, suggesting that $AMPK\alpha2$ down-regulation is D-glucose-specific (Fig. 1*A*). We also assessed the effects of high concentrations of glucose in time dependence. Exposure to 30 mM concentration of glucose for 3 days dramatically down-regulated AMPK2 expression (Fig. 1*B*). Together, these results demonstrate that high glucose conditions down-regulates $AMPK_α2$ expression in skeletal muscle cells.

Proteasome Degradation Pathway Is Involved in High Glucose-induced AMPK2 Degradation—To check the involvement of proteasome degradation in $AMPK\alpha2$ down-regulation, we analyzed the effect of proteasome inhibition on $AMPK\alpha2$ expression. MG132, a proteasome inhibitor, blocked high glucose-induced AMPKα2 down-regulation (Fig. 2A). This effect was also observed by another proteasome inhibitor, lactacystin (Fig. 2*B*), suggesting the involvement of the ubiquitin-proteasome pathway. Next, we tested whether $AMPK\alpha2$ was ubiquitinated in cells. We immunoprecipitated $AMPK\alpha2$ from cell lysates, and then analyzed with antibody against ubiquitin. As can be seen, the multi-ubiquitination of $AMPK\alpha2$ was detected in the immunoprecipitated sample (Fig. 2*C*). These results demonstrate that proteasome pathway is involved in high glu- \cos e-mediated AMPK α 2 down-regulation.

AMPK2 Interacts with WWP1—To explain the mechanism of high glucose-mediated AMPK α 2 down-regulation, we tried to identify $AMPK\alpha2$ -interacting protein. To identify protein

FIGURE 2. **Proteasome degradation pathway is involved in high glucoseinduced AMPK** α **2 degradation.** A, C2C12 cells were pre-treated 5 μ M MG132 for 1 h and stimulated with 30 mm glucose incubation for 48 h. The cell lysates (20 μ g) were analyzed via Western blotting for anti-AMPK α 2 antibody. The anti-β-actin antibody was a protein loading control. *B*, C2C12 cells were pretreated 5 μ M lactacystin for 1 h and incubated with 30 mm glucose for 48 h. The cell lysates (20 μ g) were analyzed via Western blotting for anti-AMPK α 2 antibody. The anti-ß-actin antibody was a protein loading control. C, C2C12 cells were immunoprecipitated with anti-AMPK α 2 antibody. The immunoprecipitates were analyzed via Western blotting for anti-ubiquitin and $AMPK\alpha$ 2 antibodies. *TCL*, total cell lysates.

that interacts with the catalytic domain of $AMPK\alpha2$, we carried out a YTH screen of a skeletal muscle C2C12 cDNA library. In our results, WWP1, a E3 ubiquitin ligase, was identified as an $AMPK\alpha$ 2-interacting protein. To determine whether WWP1 interacts with AMPK α 2 in mammalian cells, Flag-AMPK α 2 was transfected into HEK293 cells. Lysates from transfected cells were immunoprecipitated with antibody against Flag. A band corresponding to WWP1 was detected in the immunoprecipitates, demonstrating interaction (Fig. 3*A*). Moreover, to assess *in vivo* interaction, we performed immunoprecipitation of skeletal muscle C2C12 cell protein using an AMPK α 2 antibody, followed by Western blotting with an antibody to WWP1. $AMPK\alpha2$ was co-immunoprecipitated with WWP1 from endogenous C2C12 cells (Fig. 3*B*). Immunoprecipitation of Flag-WWP1-transfected HEK293 cells using a Flag antibody showed that $AMPK\alpha2$ was co-immunoprecipitated with WWP1 (Fig. 3*C*). These results demonstrated that the interaction detected in the YTH also occurs *in vivo* and *in vitro*.

Expression Level of AMPK2 Is Regulated by Expression Status of WWP1—To investigate the relationship between WWP1 and $AMPK\alpha2$, we first examined the effect of WWP1 knockdown on high glucose-induced $AMPK\alpha2$ down-regulation. High glucose-induced down-regulation of $AMPK\alpha2$ was blocked in WWP1 knockdown conditions. However, $AMPK\alpha2$ down-regulation was apparent in scramble siRNA, implying that WWP1 is important to down-regulate $AMPK\alpha2$ (Fig. 4*A*). Interestingly, the anti-WWP1 antibody recognized two protein bands in the C2C12 cell lysates. The intensity of the two bands changed in similar patterns, indicating that WWP1 may be recognized by both protein bands. Next, to further study the function of WWP1, we transiently transfected with Flag-WWP1. The level of $AMPK\alpha2$ was down-regulated in cells in which WWP1 was overexpressed. This down-regulation was blocked by treatment with proteasome inhibitor (Fig. 4*B*), suggesting that ubiquitin-proteasome pathway is critical for $AMPK\alpha2$ down-regulation. To determine the interaction domain, we constructed GST-fused whole AMPK α 2, C-terminal AMPK α 2, and N-terminal AMPK α 2. We incubated GST fusion proteins

FIGURE 3. **AMPK2 interacts with WWP1.** *A*, HEK293 cells were transiently transfected with Flag-AMPK α 2 for 48 h and immunoprecipitated with anti-Flag antibody. The immunoprecipitates were analyzed via Western blotting for anti-WWP1 and Flag antibodies. *B*, cell lysates (1 mg) of C2C12 cells were immunoprecipitated with anti-AMPK α 2 antibody. The immunoprecipitates were analyzed via Western blotting for anti-WWP1 and AMPK α 2 antibodies. *C*, HEK293 cells were transiently transfected with Flag-WWP1 for 48 h and immunoprecipitated with anti-Flag antibody. The immunoprecipitates were analyzed via Western blotting for anti-AMPK α 2 and Flag antibodies. *TCL*, total cell lysates.

with cell lysates from C2C12 cells. Targeted $AMPK\alpha$ 2 proteins (asterisk) were detected by Coomassie Blue staining. Immunoblotting with anti-WWP1 antibody showed that C-terminal AMPK2 interacted with WWP1 (Fig. 4*C*). Interaction between $AMPK\alpha2$ and WWP1 was enhanced by high glucose incubation (Fig. 4*D*). Together, these results indicate that WWP1 play an important role in $AMPK\alpha2$ down-regulation in high glucose culture conditions.

WWP1 Increased in High Glucose Conditions—To gain insight into the role of WWP1 in high glucose culture conditions, we examined the effect of high glucose concentrations on expression of WWP1 both in protein and mRNA levels. When cells were cultured with high concentrations of glucose, Western blot analysis showed that the level of WWP1 protein expression increased in dose-dependently (Fig. 5*A*). In experiments with RT-PCR, mRNA levels of WWP1 were also increased by incubation of C2C12 cells under a 30 mM concentration glucose condition (Fig. 5*B*). Together, these results suggest that WWP1 may be regulated by glucose concentration status.

AMPK2 Decreased in db/db Mouse—To examine the effect of high glucose on $AMPK\alpha2$ *in vivo*, we used the type 2 diabetes animal model, db/db mouse. Proteins of the quadriceps from wild type and db/db mice were prepared. Each group had 5 mice. The non-fasting blood glucose levels in db/db mice were significantly higher than those of wild type mice (data not shown). Insulin levels of db/db mice were also increased (data not shown). Hyperinsulinemic-hyperglycemia of db/db mice indicates that these animals show insulin resistance, a primary characteristic of type 2 diabetes. The expression of $AMPK\alpha2$ was down-regulated in quadriceps muscle of these mice (Fig. $6A$). Furthermore, the level of $AMPK\alpha2$ mRNA was also downregulated (Fig. $6B$). To confirm whether $AMPK\alpha$ 2 is regulated at the transcriptional level, we performed RT-PCR experiments in C2C12 cells. The level of $AMPK\alpha2$ mRNA was slightly down-regulated at 12 h after incubation with high glucose con-

FIGURE 4. Expression level of AMPKa² is regulated by expression status of WWP1.A, C2C12 cells were transiently transfected with 50 nm siRNA WWP1 and scramble siRNA for 48 h and then exposed to 30 mm glucose incubation for 48 h. The cell lysates (20 μ g) were analyzed via Western blotting for anti-AMPK α 2, anti-WWP1 antibodies. The anti-*ß*-actin antibody was a protein loading control. *B*, C2C12 cells were pre-treated 5 μ m MG132 for 1 h and then cells were transiently transfected with Flag-WWP1 for 48 h. The cell lysates (20 µg) were analyzed via Western blotting for anti-AMPK α 2, anti-Flag antibodies. The anti-*ß*-actin antibody was a protein loading control. C, C2C12 cells were prepared and incubated with GST fusion proteins of various AMPK α 2. Formed protein complexes were isolated by glutathione beads, washed three times with washing buffer, and analyzed by SDS/PAGE and subsequent Coomassie staining. The pull-down sample was also immunoblotted with anti-WWP1 antibody. D, C2C12 cells were exposed to 30 mm glucose incubation for 48 h. The cell lysates (1 mg) of normal and high glucose conditions C2C12 cells were immunoprecipitated with anti-AMPKa2 antibody. The immunoprecipitates were analyzed via Western blotting for anti-WWP1 and AMPK α 2 antibodies.

FIGURE 5. **WWP1 increased in high glucose conditions.** *A*, C2C12 cells were stimulated for the indicated doses of glucose for 48 h. The cell lysates (20 μ g) were analyzed via Western blotting for anti-WWP1 antibody. The anti- β -actin antibody was a protein loading control. *B*, total mRNA was prepared for these cells after 30 mm glucose incubation, and RT-PCR was conducted using specific WWP1 primers. The PCR product was then gel-run in 1% agarose, and visualized in UV. β -Actin was employed as a positive control.

centration, suggesting that $AMPK\alpha2$ may be regulated by transcriptional level (Fig. 6*C*). Immunostaining of db/db mouse and wild type mouse identified $AMPK\alpha2$ in quadriceps muscle. In hyperglycemic db/db mouse, the level of $AMPK\alpha$ 2 was downregulated compared with euglycemic wild type mouse. (Fig. 6*D*). The results from the diabetic animal model were correlated with biochemical data. These results indicate that the WWP1 may play a key function in $AMPK\alpha2$ down-regulation under high glucose conditions.

Expression of WWP1 Suppressed the Metformin-induced Glucose Uptake—Finally, to provide physiological relevance of WWP1, we investigated the effect of WWP1 expression on glucose uptake. Metformin-induced glucose uptake was blocked in WWP1 overexpression (Fig. 7*A*). To gain insight into the role of high glucose-mediated $AMPK\alpha2$ down-regulation, we examined the effect of metformin on glucose uptake under high glucose culture condition. Attenuation of glucose uptake under high glucose culture conditions was recovered by treatment with metformin, a first choice for the treatment of type 2 diabetes (Fig. 7*B*). These results indicated that expression of WWP1 is important for metformin-mediated glucose uptake through regulating $AMPK\alpha2$ expression.

DISCUSSION

The principal finding of this study was that WWP1 is involved in high glucose-induced $AMPK\alpha2$ down-regulation. Specifically, we demonstrated that direct interaction between AMPK α 2 and WWP1 is instrumental in AMPK α 2 down-regulation.

The anti-diabetic role of AMPK has previously been evaluated in conjunction with its activity. Dysfunction of AMPK activity may lead to metabolic syndrome. There is correlation between the low activation state of AMPK with metabolic disorders associated with insulin resistance and obesity (23, 24). AMPK activity is also suppressed in muscle and liver by sustained hyperglycemia (25). Moreover, many agents that are useful in treating diabetes, including TZDs (26, 27), metformin (28) have been shown to stimulate AMPK activity. The contribution of AMPK activity to the anti-diabetic role has raised questions as to which aspect of the protein expression of AMPK may be relevant to its metabolic role. In the present study, we have established that high glucose culture conditions down-regu-

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FIGURE 6. AMPKa2 down-regulated in db/db mouse. A, Western blot analysis of quadriceps muscles of wild type and db/db mice. The tissue lysates (40 µg) were analyzed via Western blotting for anti-AMPK α 2 antibody. The anti-ß-actin antibody was a protein loading control. Densitometry analysis (n = 5) of values are means ± S.E. values of the ratios of densities (AMPKa2/ß-actin). *, $p < 0.05$ *versus* wild type mouse. B, RT-PCR analysis of quadriceps muscles of wild type and db/db mice. Total mRNA was prepared for these mice, and RT-PCR was conducted using specific AMPKa2 primers. The PCR product was then gel-run in 2% agarose, and visualized in UV. β-Actin RNA was employed as a positive control. Densitometry analysis (*n* = 5) of values are means ± S.E. values of the ratios of densities (ΑΜΡΚα2 mRNA/β-actin mRNA). *, *p* < 0.05 *versus* wild type mouse. C, total mRNA was prepared for these cells after 30 mm glucose incubation, and RT-PCR was conducted using specific AMPK α 2 primers. The PCR product was run in 1% agarose gel, and visualized in UV. β -Actin was employed as a positive control. *D*, histochemistry of the quadriceps muscle of wild type and db/db mice. Each cryosection was stained with anti-AMPK α 2 antibody.

FIGURE 7. **Expression of WWP1 blocked metformin-induced glucose uptake.** *A*, L6 myoblast cells were differentiated for 7 days. Cells were transiently transfected with Flag-WWP1 for 48 h and then treated with metformin for 16 h. Glucose uptake was measured using 2-deoxy-D-[3 H]glucose. *, *p* 0.05 *versus* basal. *B*, L6 myoblast cells were differentiated for 7 days. Cells were incubated to 30 mm glucose for 48 h in the presence of or absence of metformin for 16 h. Glucose uptake was measured using 2-deoxy-D-[3 H]glucose. *, *p* 0.05 *versus* high glucose.

lated AMPK expression. Additionally, we have demonstrated that WWP1 down-regulated $AMPK\alpha$ 2 through direct interaction. The relationship of AMPK with ubiquitination has been suggested. Collectively, our results indicate that AMPK may down-regulate in high glucose culture conditions via the ubiquitination pathway.

The main goal of the present study was to ascertain whether or not $AMPK\alpha2$ is directly regulated by high glucose, and if so, to determine which mechanisms are involved in this process. Our data have revealed a novel role for WWP1 of $AMPK\alpha2$ down-regulation in high glucose conditions. Our identification of an AMPK α 2-WWP1 interaction in high glucose-treated

skeletal muscle cells led us to hypothesize that WWP1 might play a role in the high glucose-mediated down-regulation of $AMPK\alpha2$. Indeed, this is the first report of a link between $AMPK\alpha$ 2 and WWP1 in skeletal muscle cells. It is thus tempting to speculate that WWP1-mediated regulation of $AMPK\alpha2$ plays a role in diabetes, since $AMPK\alpha2$ expression has been implicated in insulin resistance. Overall, although the mechanism by which high glucose influences $AMPK\alpha$ 2 remains unknown, the findings in this report suggest that high glucose may down-regulate $AMPK\alpha2$ expression through the WWP1 as part of the process of high glucose-mediated insulin resistance. In the present study, we further demonstrated that high

glucose induces expression of WWP1, and overexpression of WWP1 decreased $AMPK\alpha2$ expression. In addition, wild type WWP1 blocks metformin-mediated GLUT4 translocation. These results demonstrate that WWP1 has an important role in glucose homeostasis through regulating $AMPK\alpha2$ expression. Regulation of AMPK by ubiquitination has been suggested (29, 30). The AMPK kinases NUAK1 and MARK4 are polyubiquitinated, and expression of NUAK and MARK4 mutants did not interact with the de-ubiquitinating enzyme USP9X de-ubiquitinating enzyme (DUB), thus resulting in AMPK activity (18). This result implies an important role for de-ubiquitination in AMPK activity regulation and further raises the question of what E3 ubiquitin ligases are involved. Because E3 ubiquitin ligases function to recognize specific substrates and to transfer ubiquitin to target protein, the specificity of the ubiquitin process is found in the E3 ubiquitin ligases. In the present study, we found an interaction of $AMPK\alpha2$ with E3 ubiquitin ligase WWP1. This interaction is presumably required for high glu- \cos e-mediated AMPK α 2 down-regulation. Combined with the recent report of interaction between proteasome itself and AMPK (31), protein-protein interaction has become a model for ubiquitin-mediated AMPK regulation.

On the basis of our present results, we propose a novel mechanism to explain how the expression of AMPK is down-regulated in high glucose culture conditions. This study revealed that $AMPK\alpha2$ is down-regulated in hyperglycemic animal db/db mice, while its interaction partner WWP1 is up-regulated. In conclusion, WWP1 appears to be involved in insulin resistance through suppressing $AMPK\alpha2$ expression. In conclusion, the present study supports the hypothesis that WWP1 may have an important role in hyperglycemic situations, such as diabetes and insulin resistance, through direct interaction with $AMPK\alpha2$ and down-regulation of its expression.

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