

Hypoxia reduces HNF4 α /MODY1 protein expression in **pancreatic β-cells by activating AMP-activated protein kinase**

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Hypoxia plays a role in the deterioration of β -cell function. Hepatocyte nuclear factor 4 α (HNF4 α) has an important role in β -cells, and mutations of the human *HNF4A* gene **cause a type of maturity-onset diabetes of the young (MODY1). However, it remains unclear whether hypoxia affects the expression of HNF4**- **in -cells. Here, we report that hypoxia reduces HNF4α protein expression in β-cells. Hypoxia-inducible factor** was not involved in the down-regulation of $HNF4\alpha$ under ${\rm hypoxic\ conditions.}$ The down-regulation of ${\rm HNF4}\alpha$ was depen**dent on the activation of AMP-activated protein kinase (AMPK),** and the reduction of HNF4 α protein expression by metformin, an **AMPK activator, and hypoxia was inhibited by the overexpression of a kinase-dead (KD) form of AMPK**-**2. In addition, hypoxia decreased the stability of the HNF4**- **protein, and the down-regu**lation of HNF4α was sensitive to proteasome inhibitors. Adenovirus-mediated overexpression of KD-AMPKα2 improved insulin **secretion in metformin-treated islets, hypoxic islets, and** *ob***/***ob* mouse islets. These results suggest that down-regulation of HNF4 α could be of importance in β -cell dysfunction by hypoxia.

Pancreatic β -cells sense glucose and secrete insulin to maintain normal glucose levels. Because pancreatic β -cells are highly dependent on oxidative phosphorylation for ATP production, they require large amounts of oxygen, especially at increased glucose levels (1, 2). Indeed, we and others demonstrated that pancreatic islets and insulin-producing β -cell lines become hypoxic under high glucose conditions and that the pancreatic islets of diabetic mice are hypoxic (3–5). Furthermore, we recently indicated that moderate hypoxia leads to β -cell dysfunction with a selective down-regulation of genes including *Mafa*, *Pdx1*, and *Ins1*, which play important roles in pancreatic

 β -cells (6). These results suggest that hypoxia is involved in the deterioration of β -cell function.

Hepatocyte nuclear factor 4α (HNF4 α)³ is a transcription factor belonging to the nuclear receptor superfamily and is expressed in the pancreas (including β -cells), liver, kidney, and intestine (7). We found that mutations of the human *HNF4A* gene cause a particular form of maturity-onset diabetes of the young (MODY), that is, MODY1, which is characterized by autosomal dominant inheritance, early age of onset, and pancreatic β -cell dysfunction (8). In addition, targeted disruption of HNF4 α in pancreatic β -cells leads to defective insulin secretion in mice (9, 10). These findings demonstrate the important role of HNF4 α in pancreatic β -cells.

In the present study, we investigated the impact of hypoxia on HNF4 α expression in MIN6 cells and mouse islets. We demonstrated that hypoxia decreases $HNF4\alpha$ protein expression via proteasome-mediated degradation. The hypoxia-induced down-regulation of HNF4 α was regulated by the activation of AMP-activated protein kinase (AMPK). This reduction of HNF4 α protein expression was recovered by inactivation of AMPK and re-oxygenation. Our results suggest that down-regulation of HNF4 α is a novel mechanism of β -cell dysfunction by hypoxia.

Results

Down-regulation of HNF4- *protein expression by hypoxia*

MIN6 cells were cultured under moderately hypoxic conditions (3–7% oxygen tension) for 24 h, and HNF4 α expression levels were examined by Western blot analysis. Hypoxia significantly decreased HNF4 α protein levels, but not β -actin, in a dose-dependent manner (Fig. 1, *A* and *B*). Hypoxia did not affect $HNF1\alpha/MODY3$ (11) and $HNF1\beta/MODY5$ (12) protein levels in MIN6 cells (Fig. 1*C*). Similar to MIN6 cells, decreased $HNF4\alpha$ protein levels by hypoxia were observed in pancreatic

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³ The abbreviations used are: HNF, hepatocyte nuclear factor; AICAR, 5-Aminoimidazole-4-carboxamide-1-β-D-ribofuranoside; AMPK, adenosine monophosphate-activated protein kinase; CHX, cycloheximide; HIF, hypoxia inducible factor; MODY, maturity-onset diabetes of the young; Nedd4l, developmentally down-regulated 4-like; TBP, TATA-binding protein; UTR, untranslated region; ERK, extracellular signal-regulated kinase; qRT, quantitative PCR; KD, kinase-dead.

 ${\sf Figure~1.}$ ${\sf Effect~of}$ ${\sf hypoxia}$ on ${\sf HNF4}\alpha$ ${\sf expression}$ in β -cells. A , MIN6 cells were exposed to the indicated oxygen tension (% O₂) for 24 h, and ${\sf HNF4}\alpha$ expression was examined by Western blotting. *B,* relative HNF4 α protein levels were calculated ($n=3$). C , effect of hypoxia for 24 h on HNF1 α and HNF1 β expression in MIN6 cells ($n = 3$). D, isolated mouse islets were incubated at either 5% O₂ or 20% O₂ for 24 h, and HNF4 α protein levels were evaluated by Western blot analysis ($n = 3$). E, MIN6 cells were cultured at either 5% O₂ or 20% O₂ for 12 h and 24 h, and *Hnf4a* mRNA levels were analyzed by qPCR ($n = 3$). The *Hnf4* α mRNA level was normalized to that of *TBP*. *F* and *G*, MIN6 cells were cultured at the same conditions as in *E*, and HNF4 α protein levels were examined by Western blot analysis ($n = 3$). All data are presented as mean \pm S.E. (S.E.; *error bars*). *N.S.*, not significant; *, p < 0.05; **, p < 0.01; ***, p < 0.001.

mRNA in MIN6 cells (6). We then examined the expression levels of *Hnf4a* mRNA. Hypoxia for 12 h had no effect on *Hnf4* mRNA expression in MIN6 cells (Fig. 1*E*). However, a significant down-regulation of HNF4 α protein (61.3% of control; $p < 0.01$) was detected in MIN6 cells following 5% oxygen tension for 12 h (Fig. 1, *F* and *G*), suggesting the post-transcriptional regulation of $HNF4\alpha$ expression. In this

study, we focused on the hypoxia-induced down-regulation of HNF4 α at the protein level.

E ffect of down-regulation of HNF4 α expression on β -cells

 $HNF4\alpha$ plays an important role in glucose-stimulated insulin secretion by β-cells (8). Suppression of endogenous HNF4 α consistently reduced insulin secretion in MIN6 cells [\(supple-](http://www.jbc.org/cgi/content/full/M116.767574/DC1)

Figure 2. Role of HIF-1 α **in HNF4** α **expression by hypoxia. A, MIN6 cells** were incubated at either 5% O₂ or 20% O₂ for the indicated time, and HIF-1 α protein levels were detected by Western blotting. *B* and *C,*the effect of*HIF-1* CA on HNF4 α protein levels. MIN6 cells expressing the pMX empty vector or p MX-*HIF-1* α constitutively active were cultured in normoxia and cell lysates were recovered ($n = 3$). D and *E*, MIN6 cells expressing either control shRNA or *HIF-1* α shRNA were incubated at either 5% O₂ or 20% O₂ for 24 h, and HNF4 α protein levels were examined ($n = 3$). All data are presented as mean \pm S.E. (*error bars*). *N.S*., not significant.

[mental Fig. S1,](http://www.jbc.org/cgi/content/full/M116.767574/DC1) A and B). In addition, HNF4 α knockdown significantly increased the number of propidium iodide-positive dead cells [\(supplemental Fig. S1](http://www.jbc.org/cgi/content/full/M116.767574/DC1)*C*).We previously reported that hypoxia induces an insulin secretion defect and cell death in MIN6 cells (6). Down-regulation of $HNF4\alpha$ may be involved in this dysfunction by hypoxia. In contrast, cell proliferation rate was unaffected by the down-regulation of $HNF4\alpha$ [\(supplemen](http://www.jbc.org/cgi/content/full/M116.767574/DC1)[tal Fig. S1](http://www.jbc.org/cgi/content/full/M116.767574/DC1)*D*).

Effect of HIF on HNF4 α *expression by hypoxia*

Hypoxia-inducible factor (HIF) is a transcription factor that regulates the expression of genes mediating adaptive responses to hypoxia. HIF-1 α protein is degraded under normoxic conditions, whereas it is stabilized under hypoxic conditions and binds to HIF-1 β to act as a transcription factor (13). Hypoxia (5% O_2) increased the expression of HIF-1 α protein in a timedependent manner in MIN6 cells (Fig. 2*A*). We then investigated the role of HIF-1 in the hypoxia-induced down-regulation of HNF4 α . Overexpression of a constitutively active form of HIF-1 α by retrovirus infection did not affect HNF4 α expression (Fig. 2, *B* and *C*). Next, we introduced HIF-1 α short hairpin RNA (shRNA) into MIN6 cells, and successful HIF-1 α knockdown was confirmed under hypoxic conditions (Fig. 2*D*). Suppression of HIF-1 α did not affect the down-regulation of HNF4 α at the 5% oxygen condition (Fig. 2, *D* and *E*). In addition to HIF-1 α , suppression of HIF-2 α , HIF-3 α , or HIF-1 β failed to reverse the down-regulation of HNF4 α by hypoxia [\(supplemental Fig. S2\)](http://www.jbc.org/cgi/content/full/M116.767574/DC1). These findings indicate that HIF is not involved in the down-regulation of HNF4 α under hypoxic conditions.

Effect of AMPK on HNF4- *expression by hypoxia*

In addition to the stabilization of HIF-1 α , hypoxia is known to activate AMPK (14). Indeed, the phosphorylation of both AMPK and acetyl-CoA carboxylase, a substrate of AMPK, was induced after exposure to hypoxia for 1 h in MIN6 cells [\(sup](http://www.jbc.org/cgi/content/full/M116.767574/DC1)[plemental Fig. S3\)](http://www.jbc.org/cgi/content/full/M116.767574/DC1). Down-regulation of HNF4 α was subsequently detected after exposure to hypoxia for 12 h. We then examined whether activation of AMPK could affect HNF4 α expression. Both 5-aminoimidazole-4-carboxamide-1-β-Dribofuranoside (AICAR), a cell-permeable activator of AMPK, and metformin decreased the expression of $HNF4\alpha$ in MIN6 cells (Fig. 3A). Similar to MIN6 cells, metformin also decreased HNF4 α protein levels in pancreatic islets (Fig. 3, *B* and *C*).

The kinase-dead (KD) form of $AMPK\alpha2$ (Lys-45 was changed to Arg) reportedly functions as a dominant inhibitory protein that eliminates AMPK activity (15). We examined the effect of KD-AMPK α 2 overexpression by retrovirus on the metformin-induced down-regulation of $HNF4\alpha$. Phosphorylation of acetyl-CoA carboxylase was inhibited by KD-AMPK α 2 overexpression (Fig. $3D$). In addition, HNF4 α down-regulation was significantly suppressed by KD -AMPK α 2 overexpression (Fig. 3, *D* and *E*). We next examined the effect of AMPK activity on insulin secretion in MIN6 cells. Consistent with the decreased expression of HNF4 α , metformin reduced insulin secretion at 22 mM glucose (Fig. 3, *F* and *G*). In accordance with the increased expression of HNF4 α , KD-AMPK α 2 improved insulin release by 22 mM glucose in metformin-treated MIN6 cells. Similarly, adenovirus-mediated overexpression of KD-AMPK α 2 increased $HNF4\alpha$ protein expression and improved glucose-stimulated insulin secretion in metformin-treated islets (Fig. 3, *H*–*J*).

We also investigated the effect of KD -AMPK α 2 overexpression on the hypoxia-induced down-regulation of HNF4 α . Phosphorylation of AMPK by hypoxia was inhibited by KD- $AMPK\alpha2$ overexpression [\(supplemental Fig. S4\)](http://www.jbc.org/cgi/content/full/M116.767574/DC1). Overexpression of KD-AMPK α 2 in MIN6 cells suppressed the reduction of $HNF4\alpha$ protein expression by hypoxia, as with metformin (Fig. 4, *A* and *B*). Collectively, these findings indicate that hypoxia decreases HNF4 α protein levels via the activation of AMPK. Insulin secretion was stimulated 3.4-fold by 22 mM (*versus* 2.2 mm) glucose in MIN6 cells under 20% $O₂$ tension, whereas hypoxic MIN6 cells exhibited dysregulated insulin secretion (increased insulin secretion at low glucose and blunted insulin secretion at high glucose) (6) (Fig. 4, *C* and *D*). However, the diminished glucose-stimulated changes in insulin secretion under 5% O_2 tension were significantly increased by overexpression of KD-AMPK α 2 (pMX control, 1.9-fold; pMX-KD- $\text{AMPK}\alpha2$, 2.4-fold, $p < 0.01$) (Fig. 4*D*). Overexpression of KD -AMPK α 2 also increased HNF4 α expression and enhanced insulin secretion in response to high glucose in islets under

Figure 3. Role of AMPK in HNF4 α **expression.** A, effect of AMPK activators on HNF4 α protein levels. MIN6 cells were cultured at the indicated concentration of AICAR or metformin for 24 h, and Western blotting was performed. B and C, isolated mouse islets were treated with 2 mm metformin for 24 h, and HNF4 α protein levels were examined.*D*and *E,* MIN6 cells expressing the pMX empty vector or pMX-KD-*AMPK2* vector were treated with 2 mM metformin for 20 h, and HNF4 α protein levels were examined ($n = 3$). F and G, an insulin secretion assay was performed ($n = 4$ -5), and insulin concentration was determined by an insulin ELISA. Fold-change in glucose-stimulated insulin secretion (insulin level at 22 mm glucose divided by that at 2.2 mm glucose) is shown ($n = 4$ –5). *H*, isolated mouse islets expressing either LacZ or KD-AMPKα2 were treated with 2 mm metformin for 20 h, and HNF4α protein levels were examined by Western blot analysis. *I* and *J,* islet insulin secretion was examined. Insulin levels are expressed as absolute values or as fold-change (*n* 11–12). All data are presented as mean \pm S.E. (*error bars*). *N.S.,* not significant; *, p $<$ 0.05; **, p $<$ 0.01; ***, p $<$ 0.001.

hypoxic conditions (Fig. 4, *E*–*G*). These results suggest the functional significance of AMPK- and hypoxia-dependent regulation of HNF4 α expression on insulin secretion.

HNF4- *protein stability under hypoxic conditions*

Hypoxia reportedly leads to the degradation of proteins, such as estrogen receptor α and Na,K-ATPase (16, 17). We next evaluated whether HNF4 α protein stability is influenced by hypoxia. MIN6 cells were cultured under 20 or 5% oxygen tension in the presence of cycloheximide (CHX), a translational inhibitor. During incubation with CHX, HNF4 α protein decayed rapidly in the 5% oxygen condition (Fig. 5*A*), indicating that hypoxia decreases the stability of $HNF4\alpha$ protein. We also investigated HNF4 α stability by metformin treatment. Metformin promoted HNF4 α degradation (Fig. 5B). Then, $HNF4\alpha$ protein stability was tested in the presence or absence of the proteasome inhibitor MG132 (Fig. 5*C*). As shown in Fig. 5, *C*–*F*, proteasome inhibition resulted in the stabilization of

Figure 4. Role of AMPK in hypoxia-induced HNF4 α **down-regulation.** A and B, MIN6 cells expressing the pMX empty vector or pMX-KD-AMPK α 2 vector were cultured at either 5% O₂ or 20% O₂ for 20 h, and HNF4 α protein levels were examined (*n* = 3). C and *D*, an insulin secretion assay was performed (*n* = 6–9), and insulin concentration was determined by an insulin ELISA. Fold-change in glucose-stimulated insulin secretion (insulin level at 22 mm glucose divided by that at 2.2 mm glucose) is shown ($n = 6$ -9). *E*, mouse isolated islets expressing either LacZ or KD-*AMPK* α 2 were treated with 5% O₂ for 20 h, and HNF4 α protein levels were examined by Western blot analysis. *F* and*G,* islet insulin secretion was examined, and insulin levels are expressed as absolute values or asfold-change (*n* 5–10). Data are presented as mean \pm S.E. (*error bars*). *N.S.*, not significant; *, p $<$ 0.05; **, p $<$ 0.01; ***, p $<$ 0.001.

 $HNF4\alpha$ in both hypoxic and metformin-treated conditions (compare *lanes 3* and *4* of Fig. 5*D*). Another proteasome-specific inhibitor, epoxomicin, also stabilized $HNF4\alpha$ protein expression under hypoxia [\(supplemental Fig. S5\)](http://www.jbc.org/cgi/content/full/M116.767574/DC1). These findings indicate that proteasome-mediated degradation is involved in the hypoxia- and AMPK activation-induced down-regulation of HNF4 α .

AMPK activation leads to decreased levels of proteins by regulating the binding of RNA-binding proteins within the 3'-untranslated region (3-UTR) of target mRNAs (18). To address the role of the $3'$ -UTR in HNF4 α mRNA, MIN6 cells were infected with a retroviral HNF4 α -FLAG expression vector (lacking both the $3'$ -UTR and $5'$ -UTR of the $HNF4\alpha$ gene). Both hypoxia and metformin decreased HNF4 α protein levels (Fig. 6, A and *B*), indicating that the 3-UTR is not involved in down-regulation of HNF4 α by hypoxia/AMPK activation in MIN6 cells.

Phosphorylation of HNF4α by hypoxia

The phosphorylation of proteins triggers protein degradation (19, 20). AMPK activation reportedly leads to the phosphorylation of the liver type of HNF4 α at serine 304 (serine 291 of HNF4 α 7) in vitro (21). Thus, we examined whether this phosphorylation is essential for the hypoxia-induced down-regulation of HNF4 α in HNF4 α 7-FLAG-expressing MIN6 cells. As shown in Fig. 7A, HNF4 α 7 with a mutation of serine 291 to alanine (S291A) was still down-regulated by hypoxia. Database screening (ppsp.biocuckoo.org/index.php) identified another potential AMPK phosphorylation site $(^{117}TRRSSYEDS^{125}$, potential phosphorylation serine 121 is underlined) in HNF4 α 7. However, the replacement of serine 121 with alanine (S121A) also did not restore HNF4 α 7 instability (Fig. 7A). Because both S121A and S291A mutants were as sensitive to hypoxia-induced down-regulation as wild-type HNF4 α 7, we investigated whether HNF4 α 7 was phosphorylated by hypoxia and AMPK activation in MIN6 cells. Phosphorylated proteins show slower migration in Phos-tag SDS-PAGE due to the selective binding of phosphorylated amino acids to the Phos-tag reagent (22). After treatment with metformin or hypoxia, whole cell lysates of MIN6 cells were separated by SDS-PAGE with Phos-tag, and

Figure 5. Regulation of HNF4 α **protein stability by hypoxia or metformin.** A **, MIN6 cells were incubated at 5%** $\rm O_2$ **or 20%** $\rm O_2$ **for 18 h, and HNF4** α **protein** stability was examined for another 6 h using CHX (10 μ g/ml). Relative HNF4 α protein levels are indicated as percentage values ($n = 3-4$). Half-lives of HNF4 α protein are indicated. *B*, MIN6 cells were incubated with or without metformin (2 mm) for 18 h, and HNF4 α protein stability was examined. Relative HNF4 α protein levels are indicated as percentage values ($n = 3-4$). Half-lives of HNF4 α protein are shown. *C*, experimental scheme using the MG132 proteasome inhibitor. *D*, effect of MG132 (10 μM) on HNF4α expression by hypoxia or metformin treatment. *E* and *F*, relative HNF4α protein levels in 5% O₂ (hypoxia) (*E*) or metformin (*F*) were calculated (*n* = 3). All data are presented as mean \pm S.E. (error bars). *N.S.*, not significant; * , p < 0.05; ** , p < 0.01.

blotted with anti-AMPK and anti-HNF4 α antibodies. Treatment with hypoxia or metformin led to AMPK phosphorylation (Fig. 3). Phosphorylated AMPK following hypoxia or metformin treatment was detected as a band with slower migration using an anti-AMPK antibody (Fig. 7*B*). It has been reported that protein kinase A phosphorylates $HNF4\alpha$ (23). Consistently, a band shift of HNF4 α was observed by treatment with forskolin, which activates the protein kinase A pathway via cAMP. In contrast, Western blot analysis with an anti-HNF4 α antibody did not show a detectable band shift by

treatment with hypoxia or metformin, suggesting that the HNF4 α protein is not phosphorylated by metformin and 5% hypoxia in MIN6 cells.

The effect of ERK5 and E3 ubiquitin ligases on HNF4 α *expression by hypoxia*

A recent study showed that extracellular signal-regulated kinase 5 (ERK5)/mitogen-activated protein kinase 7 (MAPK7) up-regulates proteasome levels upon mTOR (mechanistic target of rapamycin) signaling inhibition (24). Because hyp-

Figure 6. Role of the 3'-UTR in HNF4 α **protein stability. A, MIN6 cells** expressing HNF4 α 7-FLAG by retrovirus transfection were incubated at either 5% O_2 or 2 mm metformin treatment for 24 h, and exogenous HNF4 α protein expression was detected using an anti-FLAG antibody. B , relative HNF4 α protein levels were calculated ($n = 3$). All data are presented as mean \pm S.E. (*error bars*). *N.S.,* not significant; **, $p < 0.01$; ***, $p < 0.001$.

Figure 7. Phosphorylation of HNF4 α **protein by hypoxia.** A, MIN6 cells expressing wild-type (WT) or mutant (S121A or S291A) HNF4α7-FLAG by retrovirus transfection were cultured at either 5% O_2 or 20% O_2 for 24 h. Both exogenous and endogenous HNF4 α proteins were examined using an anti-FLAG antibody and anti-HNF4 α antibody, respectively. *B*, MIN6 cells were treated with metformin (2 mm) or incubated in 5% $O₂$ for 12 and 24 h. Phosphorylation status of both AMPK and HNF4 α was evaluated using Phos-tag SDS-PAGE. For a positive control, MIN6 cells were treated with either DMSO or forskolin (FSK , 100 μ _M) for 3 h.

oxia increased the expression of ERK5 protein (Fig. 8*A*), we examined the effect of ERK5 on HNF4 α expression. The hypoxia-induced down-regulation of $HNF4\alpha$ was similar between control and ERK5 knockdown MIN6 cells (Fig. 8, *B* and *C*, [supplemental Fig. S6,](http://www.jbc.org/cgi/content/full/M116.767574/DC1) *A* and *B*), indicating that ERK5 does not contribute to the down-regulation of $HNF4\alpha$ by hypoxia.

Activation of AMPK causes proteasomal degradation of proteins via E3 ubiquitin ligases, including atrogin-1, neural precursor cell expressed, developmentally down-regulated gene

Figure 8. Role of ERK5 and E3 ubiquitin ligases in HNF4 α expression by **hypoxia.** A, MIN6 cells were incubated in either 5 or 20% O₂ for 18 h, and expression of ERK5 or HNF4 α protein was examined by Western blot analysis. *B,* MIN6 cells expressing either control shRNA or Erk5 shRNA were incubated in either 5 or 20% O₂ for 24 h, and HNF4 α protein levels were examined by Western blot analysis. C, relative HNF4 α protein levels were calculated (*n* 3). *D–G,* effect of overexpression of atrogin-1 shRNA (*D* and *E*) or Nedd4l shRNA (*F* and *G*) on HNF4 expression was examined. All data are presented as mean \pm S.E. (*error bars*). *N.S.*, not significant; *, p $<$ 0.05; ** , *p* $<$ 0.01; ***, *p* $<$ 0.001.

4-like (Nedd4l), muscle RING finger-1 (MuRF1), and Smad ubiquitylation regulatory factor-1 (Smurf1) (25–27). Because the expression levels of MuRF1 and Smurf1 mRNA were low in MIN6 cells (data not shown), we examined the effect of atrogin-1 and Nedd4l on HNF4 α expression. Suppression of these E3 ubiquitin ligases did not reverse the down-regulation of HNF4 α protein under hypoxia (Fig. 8, *D*–*G*, [supplemental](http://www.jbc.org/cgi/content/full/M116.767574/DC1) [Fig. S6,](http://www.jbc.org/cgi/content/full/M116.767574/DC1) *C* and *D*), suggesting that these ubiquitin ligases are also not involved in the down-regulation of $HNF4\alpha$.

Restoration of HNF4- *protein expression by re-oxygenation or inactivation of AMPK*

We next examined the effect of re-oxygenation on HNF4 α expression. Expression of HNF4 α protein was increased in MIN6 cells that were re-oxygenized for more than 3 h (R3–R12) (Fig. 9A). Re-oxygenation for 12 h (R12) also increased HNF4 α expression in pancreatic islets (Fig. 9, *B* and *C*). Consistent with our previous finding that pancreatic islets of diabetic *ob*/*ob* mice are hypoxic (3), HNF4 α protein expression was significantly

Figure 9. Impact of re-oxygenation on HNF4 α **expression.** A, MIN6 cells were cultured at 5% O₂ for 24 h and then cultured at 20% O₂ (re-oxygenation) for the indicated time (0.5–12 h). HNF4 α expression was examined by Western blot analysis. *B* and *C*, isolated mouse islets were cultured at 5% O₂ for 20 h and then cultured at 20% O₂ for 12 h. HNF4 α protein levels were examined ($n = 3$). *D* and *E*, islets were isolated from C57BL/6J or *ob/ob* mice. HNF4 α protein levels were examined immediately after islet isolation and after incubation at 20% O₂ for 12 h ($n = 4 - 6$). *F*, islets from *ob/ob* mice were infected with adenovirus expressing LacZ or KD-AMPK_Q2 for 2 h, and cultured for 36 h. Expression of AMPK_Q and HNF4_Q was examined by Western blotting. G, insulin secretion assay was performed using *ob/ob* mouse islets expressing either LacZ (*n* = 13) or KD-*AMPKα2* (*n* = 24). Insulin concentration was determined by an insulin ELISA. *H*, fold-change in glucose-stimulated insulin secretion (insulin level at 22 mm glucose divided by that at 2.2 mm glucose) is shown. All data are presented as mean \pm S.E. (*error bars*). *N.S*., not significant; *, *p* - 0.05; **, *p* - 0.01; ***, *p* - 0.001.

decreased in *ob*/*ob* mouse islets (Fig. 9,*D*and *E*). However, expression levels of the HNF4 α protein became similar between control and *ob*/*ob* islets after incubation at 20% oxygen tension.

Finally, we investigated the effect of KD-AMPK α 2 on insulin secretion in *ob*/*ob* islets. Insulin secretion from *ob*/*ob* islets was unresponsive to glucose stimulation (28). Consistently, insulin

Figure 10. Schematic representation of the hypoxia-induced down-reg u lation of HNF4 α in β -cells.

release by high glucose did not differ from that by low glucose in control ob/ob islets. However, overexpression of KD-AMPK α 2 increased the expression of $HNF4\alpha$ in ob/ob islets, and insulin secretion by high glucose was significantly improved (Fig. 9, *F*–*H*).

Discussion

HNF4 α 7 and HNF4 α 8 are major isoforms of HNF4 α in pancreatic β -cells, and HNF4 α plays important roles in the function of these cells (8, 29, 30). In the present study, we demonstrated that hypoxia leads to the down-regulation of HNF4 α protein in β -cells. HNF4 α down-regulation was dependent on AMPK activation and was sensitive to proteasome inhibitors, indicating that $HNF4\alpha$ proteasomal degradation is regulated, at least in part, by AMPK activation (Fig. 10). A reduction of hepatic HNF4 α (HNF4 α 1 and HNF4 α 2 are major hepatic forms) by AICAR treatment has also been reported (31). HNF4 α plays an essential role in glucose-stimulated insulin secretion by pancreatic β -cells (8, 9) [\(supplemental Fig. S1\)](http://www.jbc.org/cgi/content/full/M116.767574/DC1). Metformin treatment attenuated high glucose-stimulated insulin secretion with the decreased expression of $HNF4\alpha$. Conversely, overexpression of KD-AMPK α 2 improved insulin secretion with the increased expression of $HNF4\alpha$ (Fig. 3). Collectively, these results suggest that the regulation of $HNF4\alpha$ expression by AMPK has a functional consequence on insulin secretion. Hypoxia and AMPK activation inhibit multiple steps in the process of insulin secretion (6, 32–36). Our findings uncover a new role for AMPK in impaired insulin secretion by β-cells. Decreased HNF4α expression in β-cells could reduce oxygen consumption by reducing insulin secretion under hypoxic conditions. Thus, the down-regulation of $HNF4\alpha$ may be a fail-safe mechanism against hypoxia. However, prolonged \exp osure of β -cells to hypoxia, in turn, would have deleterious effects on insulin secretion by reducing $HNF4\alpha/MODY1$ transcription factor levels and could contribute to the development of type 2 diabetes.

AMPK activation reportedly induces the phosphorylation of hepatic HNF4 α in vitro (21), but the replacement of potential AMPK phosphorylation sites (Ser^{121} and Ser^{291}) did not affect HNF4 α instability in MIN6 cells (Fig. 7). A mobility shift of $HNF4\alpha$ after hypoxia or metformin treatment was not detected using Phos-tag SDS-PAGE. In addition, we could not detect a phosphorylation signal in HNF4 α immunoprecipitates that were prepared from hypoxic MIN6 cells using the commercially available anti-phospho-AMPK substrate motif MultiMab antibody (data not shown). Although the possibility that β -celltype HNF4 α is a direct target of AMPK cannot be excluded, these results suggest that $HNF4\alpha$ is not phosphorylated directly by metformin and/or hypoxia in MIN6 cells.

We have demonstrated that hypoxia increases the expression of ERK5, and ERK5 has been shown to up-regulate proteasome levels (24). Moreover, AMPK activation leads to proteasomal degradation of target proteins via the ubiquitin ligases atrogin-1 and Nedd4l (25, 26). However, these molecules do not appear to account for the mechanism of $HNF4\alpha$ degradation by AMPK (Fig. 8). Further studies are necessary to clarify the role of AMPK on HNF4 α expression in β -cells.

We found that re-oxygenation restored $HNF4\alpha$ expression in both control and *ob*/*ob* islets (Fig. 9). In addition, restoration of HNF4 α in *ob/ob* islets significantly improved insulin secretion. Restoration of HNF4 α alone may not be sufficient to normalize completely the impaired secretion of insulin because *ob*/*ob* islets exhibit additional abnormalities (37). However, our findings suggest that methods to enhance oxygenation in hypoxic β -cells could be an effective therapeutic approach to remedy insulin secretion defects in type 2 diabetes by restoring $HNF4\alpha$.

There are limitations in this study. First, the mechanisms by which AMPK activation leads to the down-regulation of HNF4 α are unclear. Second, we do not have direct evidence that hypoxia leads to pancreatic β -cell HNF4 α protein degradation *in vivo*. Investigation of $HNF4\alpha$ protein levels of mouse β -cells in a low oxygen environment would be a major undertaking.

In conclusion, we demonstrated that hypoxia decreased HNF4 α protein expression via proteasome-mediated protein degradation in β -cells. Furthermore, the hypoxia-induced down-regulation of HNF4 α was regulated by the activation of AMPK. Because a small perturbation of HNF4 α activity in β -cells results in MODY1 (38), our findings suggest that downregulation of HNF4 α could be of importance in β -cell dysfunction by hypoxia.

Experimental procedures

Reagents

AICAR and 1,1-dimethylbiguanide hydrochloride (metformin) were purchased from Sigma. CHX was obtained from Nacalai Tesque (Kyoto, Japan), MG132 from Calbiochem, and epoxomicin from the Peptide Institute (Osaka, Japan).

Cell culture

The MIN6 β -cell line (39) was a gift from Jun-ichi Miyazaki (Osaka University). MIN6 cells were maintained in Dulbecco's modified Eagle's medium supplemented with 10% (v/v) fetal

bovine serum (FBS), 0.1% (v/v) penicillin/streptomycin, and 50 μм β-mercaptoethanol at 37 °C in 5% CO₂, 95% air. A multigas incubator (APM-300; ASTEC, Fukuoka, Japan) and an INVIVO₂400 hypoxia work station (Ruskin Technology, Leeds, UK) were used to render the cells chronically hypoxic (6).

Animals

C57BL/6J and B6.Cg-Lep*ob*/J (*ob*/*ob*) mice were purchased from KBT Oriental Co., Ltd. (Saga, Japan). The mice were kept under specific pathogen-free conditions in a 12-h light (7:00– 19:00)/12-h dark (19:00–7:00) cycle with free access to water and normal mouse chow (CE-2; CLEA, Tokyo, Japan). Room temperature was maintained at $22 \pm 1-2$ °C. Handling and killing of the mice by cervical dislocation were in compliance with the animal care guidelines of Kumamoto University. This study was approved by the animal research committee of Kumamoto University and all animal experimental protocols were approved by the Kumamoto University Ethics Review Committee for Animal Experimentation.

Isolation of pancreatic islets

Pancreatic islets were isolated from C57BL/6J or *ob*/*ob* mice (20–30 weeks old) as described previously (40). Briefly, after bile duct cannulation and digestion of the pancreas using a mixture of 1.3 mg/ml of collagenase L (Nitta Gelatin, Osaka, Japan), 1.3 mg/ml of hyaluronidase (H3506; Sigma), and 0.1% (v/v) protease inhibitor mixture (Nacalai Tesque), isolated islets were manually picked up and they were maintained in RPMI-1640 medium supplemented with 10% (v/v) FBS, 0.1% (v/v) penicil- $\text{lin}/\text{streptomycin}, 50 \ \mu\text{M}$ β -mercaptoethanol, 10 mM HEPES (Nacalai Tesque), and 1 mm sodium pyruvate (Nacalai Tesque) at 37 °C in 5% CO_2 , 95% air.

Western blotting

Western blotting was performed as described previously (41). For specific detection of proteins, the following primary antibodies were used: anti-HNF4 α (H1415; Perseus Proteomics, Tokyo, Japan), anti-HNF1 α (610902; BD Biosciences, San Jose, CA), anti-HNF1 β (H85; Santa Cruz Biotechnology, Santa Crux, CA), anti-β-actin (clone AC15; Sigma), anti-GAPDH (Promega, Madison, WI), anti-HIF-1 α (NB100-479; Novus Biologicals, Littleton, CO), anti-HIF-1 β (#611078; BD Transduction Laboratories, San Jose, CA), anti-phospho-AMPK (Thr^{172}) (40H9) (number 2535; Cell Signaling, Beverly, MA), anti-AMPK (number 2532; Cell Signaling), anti-phosphoacetyl-CoA carboxylase (Ser^{79}) (number 3661; Cell Signaling), anti-acetyl-CoA carboxylase (number 3662; Cell Signaling), anti-ERK5 (number 3372; Cell Signaling), and anti-DYK-DDDDK (FLAG) tag monoclonal (Clone No. 1E6; Wako Pure Chemicals, Osaka, Japan). After incubation with secondary antibodies, the signals were detected by using Chemi-Lumi One L or Chemi-Lumi One Super (Nacalai Tesque).

Quantitative real-time RT-PCR

For quantitative real-time PCR (qPCR), total RNA was extracted from MIN6 cells. qPCR was performed using SYBR Premix Ex TaqII (RR820A; TaKaRa Bio Inc., Shiga, Japan) in an ABI 7300 thermal cycler (Applied Biosystems, Foster City, CA)

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(42). The mRNA level of each gene was normalized to that of the TATA-binding protein (TBP). The specific primers were as follows: HNF4 α : forward, 5'-GCCACAGTTTTCCACCAA-GAG-3', reverse, 5'-AAGGAGGACGTCTGCTTCTGA-3'; HIF-2a: forward, 5'-CCCCGATGAATTCACCCAAA-3', reverse, 5'-TCTTGGCGTTCTCCCCTGAG-3'; HIF-3a: forward, 5'-CACTTTGTTGAGCAAGGGCC-3', reverse, 5'-TGCGAGTATGTTGCTCCGTT-3'; atrogin-1: forward, 5'-GTGGCATCGCCCAAAAGAAC-3', reverse, 5'-CTTGC-CGACTCTCTGCACCA-3'; Nedd4l: forward, 5'-TCGGCCA-ATCATGCAGCTTGC-3', reverse, 5'-TGTGTTGTGGTTT-GGGGGAGTTG-3; Erk5: forward, 5-TTGCGCCCCACCT-TTTGACTTTG-3, reverse, 5-AAGCGGATTTGTTGG-CGGATGC-3'; TBP: forward, 5'-CCCCTTGTACCCTTCAC-CAAT-3', reverse, 5'-GAAGCTGCGGTACAATTCCAG-3'.

Vector constructs, retrovirus-mediated transduction, and adenovirus-mediated transduction

The pMXs-HIF-1 α CA5 and pSUPER.retro-HIF-1 α plasmids have been described previously (3, 6). A kinase-dead α 2 isoform of K45R AMPK (KD-AMPK α 2) was a gift from Morris Birnbaum (Addgene plasmid number 15992, Addgene, Cambridge, MA) (15) and cloned into the pMXs-puro retroviral vector. To introduce an in-frame FLAG tag at the C-terminal of HNF4 α 7, PCR was performed using pcDNA3-HNF4 α 7 (42) as a template and the following primers: 5'-CCCCTGCACCCT-CACCTGATGCAG-3' and 5'-GAATTCCTACTTGTCATC-**GTCGTCCTTGTAGTC**GATAACTTCCTGCTTGGTGAT-GGT-3 (the underlined nucleotides indicate an EcoRI cloning site, and the bold letters indicate the FLAG sequence). After verification of the sequence, an NdeI/EcoRI fragment of the PCR product and an EcoRI/NdeI fragment of pcDNA3- $HNF4\alpha$ 7 were subcloned into the EcoRI site of pcDNA3.1 to generate pcDNA3.1-HNF4 α 7-FLAG. S121A and S291A mutations were introduced into the pcDNA3.1-HNF4 α 7-FLAG plasmid using a KOD-Plus Mutagenesis Kit (TOYOBO, Osaka, Japan) according to the manufacturer's protocol. The following primers were used: 5'-TCAGCCTATGAGGACAGCAGC-CTG-3' and 5'-CCTTCGAGTGCTGATCCGGTCCCG-3' (S121A); 5'-CGTGCCCAGGTGCAGGTGAGCTTG-3' and 5-GATCCAGGGAAGATCAAGCGGCTG-3 (S291A) (the underlined nucleotides indicate the mutated nucleotides for alanine). EcoRI fragments of pcDNA3.1-HNF4 α 7 (wild-type, S121A, and S291A)-FLAG were cloned into the pMXs-puro retroviral vector. The pSIREN-RetroQ HNF4 α shRNA expression vector has been described previously (35). For knockdown of HIF-2 α , HIF-3 α , atrogin-1, developmentally down-regulated 4-like (Nedd4l), and Erk5, oligonucleotides encoding respective shRNA was cloned into a pSIREN-RetroQ expression vector (Clontech Laboratories, Mountain View, CA). Target sequences were as follows: 5'-GGGACTTACTCAGGTAGA-ACT-3' for HIF-2a shRNA, 5'-TTTCCGACAGGTAAGCCA-TGT-3' for HIF-3a shRNA, 5'-AAAGTACAGTATCCATGG-CGC-3' for atrogin-1 shRNA, 5'-GGAGACCATTTCAGAG-GAAAT-3' for Nedd4l shRNA, 5'-ATTGTGGCTGAGATTG-AGGACTT-3' for Erk5 shRNA. Retrovirus transfection was performed as described previously (6). Recombinant adenovirus expressing KD-AMPK α 2 or LacZ was prepared using

Adeno-X Expression System 1 (Clontech Laboratories) per the manufacturer's protocol. Briefly, KD-AMPKa2 was subcloned into Adeno-X viral DNA via the shuttle vector pShuttle2. Then, 10 μ g of linearized Adeno-X KD-AMPK α 2 vector was transfected to a low passage of HEK293T using JetPRIME transfection reagent (Polyplus-transfection SA, Illkirch, France). Recombinant adenovirus was amplified in HEK293T cells and viral titers were determined by Adeno-X qPCR Titration Kit according to the manufacturer's protocol (Clontech Laboratories).

Insulin secretion assay

Control MIN6 cells or $HNF4\alpha$ knockdown MIN6 cells were seeded in a 24-well plate at 5.0×10^5 cells/well and maintained for 2–3 days. They were preincubated for 1 h in Krebs-Ringerbicarbonate HEPES (KRBH) buffer (120 mm NaCl, 4.7 mm KCl, 1.2 mm KH_2PO_4 , 2.4 mm $CaCl_2$, 1.2 mm $MgCl_2$, 20 mm $NaHCO₃$, and 10 mm HEPES) containing 2.2 mm glucose and 0.5% (v/v) bovine serum albumin (BSA) at pH 7.4. They were incubated in 2.2 or 22 mm glucose containing KRBH buffer for 1 h, and the culture supernatant was recovered.

Then, mouse pancreatic islets from C57BL/6J mice or *ob*/*ob* mice were cultured in RPMI-1640 medium for 2 days, and islets of similar size were randomly picked up, divided into 2 groups (20–30 islets), and separately infected with recombinant adenovirus (LacZ or KD-AMPK α 2, 2.0 \times 10⁵ infectious units (IFU)) for 2 h. After 24-h incubation in RPMI-1640 medium, these islets were further cultured in 2 mM metformin or 5% $O₂$ for 20 h, and then the insulin secretion assay was performed until 48 h after adenoviral infection. The islets were preincubated for 30 min in KRBH buffer containing 2.2 mm glucose and 0.5% (v/v) BSA. They were incubated in 2.2 or 22 mM glucose containing KRBH buffer for 30 min, and the culture supernatant was recovered to evaluate insulin secretion. Insulin concentration was determined by a mouse insulin ELISA (TMB) kit (AKRIN-011T; Shibayagi Co., Ltd., Gunma, Japan). For MIN6 cells, insulin levels were standardized by whole cell protein content.

Apoptosis assay

Control MIN6 cells or HNF4 α knockdown MIN6 cells were incubated for 48 h, and then trypsinized and collected by centrifugation at 6,000 rpm for 10 min. Cells were stained with propidium iodide for 5 min at room temperature in the dark, and stained cells were immediately analyzed using a FACSCalibur flow cytometer (BD Biosciences) and FlowJo software (Tomy Digital Biology, Tokyo, Japan).

Cell proliferation assay

Control or HNF4 α knockdown MIN6 cells were seeded in a 96-well plate at 1.5×10^4 cells/well. After 2 days, a cell proliferation assay was performed for 3 consecutive days using cell proliferation reagent WST-1 (Roche Diagnostic, Mannheim, Germany) and the absorbance (450/655) was measured by iMark microplate reader (Bio-Rad).

Protein degradation assay

MIN6 cells were seeded at 2.0 \times 10⁵ cells in a 35-mm dish. After MIN6 cells were cultured at 20% O_2 or 5% O_2 condition for 18 h, they were then treated with 10 μ g/ml of CHX for the indicated time $(0 - 6 h)$. To examine the effects of MG-132 or epoxomicin on HNF4 α protein expression, MIN6 cells were cultured at either 20% O_2 or 5% O_2 for 20 h and then treated with CHX in the presence or absence of MG132 (or epoxomicin) for 4 h. MIN6 cells were also cultured with or without 2 mm metformin for 20 h and then treated with CHX $(+)$ /MG132 $(+)$ or CHX (+)/MG132 (-) for 4 h. HNF4 α protein levels were analyzed by Western blotting.

Phos-tag SDS-PAGE

MIN6 cells were treated with 2 mm metformin or cultured under the hypoxic 5% O_2 condition for 12 and 24 h. Whole cells were lysed in EDTA $(-)$ RIPA buffer (50 mm Tris-HCl, pH 8.0, 150 mM NaCl, 0.1% SDS, 1% Nonidet P-40, 0.5% sodium deoxycholate, 20 mg/ml of Na_3VO_4 , 10 mm NaF, 1 mm PMSF, and 1% (v/v) protease inhibitor mixture). Phos-tag-containing polyacrylamide gels were made with Phos-tag acrylamide (40 μ M; Wako Pure Chemicals, Osaka, Japan) and MnCl₂ (40 μ M) according to the manufacturer's recommendation. Twenty micrograms of proteins were subjected to 10% SDS-PAGE. Dr. Western (Oriental Yeast Co., Ltd, Tokyo, Japan) was used for a protein ladder marker. After electrophoresis, the Phos-tag gel was soaked in transfer buffer (25 mm Tris and 192 mm glycine) containing 1 mm EDTA for 10 min to remove the Mn^{2+} before transferring the proteins to a membrane, and then Western blotting was performed.

Statistical analysis

The significance of differences was assessed with an unpaired t test and a value of $p < 0.05$ was considered to be statistically significant.

Author contributions—Y. S. and K. Y. designed the study and wrote the paper. Y. S., T. T., C. S., M. F. K., T. Y., and M. I. contributed to the acquisition of data, the statistical data analyses, and drafting of the manuscript. All authors have approved the final version of the manuscript.

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