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Short communication: Protein kinase C regulates glucose uptake and mRNA expression of glucose transporter (GLUT) 1 and GLUT8 in lactating bovine mammary epithelial cells

K. Zhao,*† H.-Y. Liu,*¹ F.-Q. Zhao,*‡ and J.-X. Liu*¹ *Institute of Dairy Science, MOE Key Laboratory of Molecular Animal Nutrition, Zhejiang University, Hangzhou 310058, P. R. China †Shaanxi Institute of Zoology, 88 Xingging Road, Xi'an 710032, P. R. China ‡Laboratory of Lactation and Metabolic Physiology, Department of Animal Science, University of Vermont, Burlington 05405

ABSTRACT

The aim of this study was to determine the role of protein kinase C (PKC) in regulating glucose uptake in lactating bovine mammary epithelial cells (BMEC). The BMEC were cultured and treated with different concentrations of phorbol 12-myristate 13-acetate (PMA; 0, 10, 25, 50, 100, and 200 ng/mL, the classic activator of PKC, for 48 h. Compared with the cells with no PMA treatment, 50 and 100 ng of PMA/mL significantly stimulated the glucose uptake of the BMEC, whereas the glucose uptake by the cells treated with the lowest and the highest amounts of PMA (25 and 200 ng/mL, respectively) did not show a significant difference. Consistently, the mRNA expression of glucose transporter (GLUT) 1 and 8 showed a similar pattern of increase under the treatments of PMA. Furthermore, when the cells were pretreated with GF1090203X (0, 0.25, 0.5, 1, and 2 μ *M*), an inhibitor of PKC, for 30 min before exposed to PMA (50 ng/mL), the PMA-induced glucose uptake and GLUT1 and GLUT8 expression were decreased by GF1090203X in a dose-dependent manner. These results demonstrate that PKC is involved in the regulation of glucose uptake by BMEC, and this function may work, at least partly, through upregulating the expression of *GLUT1* and *GLUT8*.

Key words: bovine, glucose transporter, mammary epithelial cell, protein kinase C

Short Communication

Glucose is an important substrate and energy source in milk synthesis. Large amounts of glucose are required to sustain lactation (Cant et al., 2002). Glucose is taken up from the blood stream by mammary secretory epithelial cells (MEC) through facilitative glucose transporters (**GLUT**), mainly GLUT1 and possibly GLUT8 (Zhao, 2014).

Protein kinase C (**PKC**) is an important cell signaling molecule that modulates a variety of cellular processes, such as secretion, gene expression, proliferation, differentiation, and muscle contraction (Nishizuka, 1986). In addition, it is well known that PKC is a key regulator in glucose uptake in a variety of cells (Watson and Pessin, 2001; Bosch et al., 2004; Luiken et al., 2009). However, whether PKC regulates glucose uptake in MEC is not known. Expression of PKC proteins increases in human mammary epithelial tissue from puberty to pregnancy (Masso-Welch et al., 1999). Activation of PKC enhances proliferation and survival (Grossoni et al., 2007) and plays an important role in prolactininduced milk protein synthesis in mouse MEC (Marte et al., 1994). Furthermore, changes in PKC expression or activity have been observed during mammary carcinogenesis (Urtreger et al., 2012). Therefore, PKC may also be an important regulator in regulating glucose uptake in MEC. The objective of the current study was to determine the role of PKC in regulation of glucose transport and glucose transporter gene expression in lactating bovine mammary epithelial cells (**BMEC**).

The lactating BMEC were isolated, characterized, and cultured as previously described (Zhao et al., 2010; Liu et al., 2013). The cells were starved with serum-free medium for 12 h and then treated with (1) different concentrations (0, 10, 25, 50, 100, and 200 ng/mL) of phorbol 12-myristate 13-acetate (**PMA**; Sigma, St. Louis, MO), a classic activator of PKC, for 48 h and (2) with different concentrations (0, 0.25, 0.5, 1, and 2) μM) of GF1090203X (Merck Biosciences, Darmstadt, Germany), an inhibitor of PKC, for 30 min before being treated with PMA (50 ng/mL) for another 48 h. The difference in glucose content in the culture media before and after the treatments, determined by an enzymatic coloring glucose oxidase/peroxidase assay method (Tiffany et al., 1972) and normalized by total cell protein content, was considered as the net glucose uptake (Accorsi et al., 2005).

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¹Corresponding authors: hyliu@zju.edu.cn and liujx@zju.edu.cn

TRIzol reagent (Invitrogen Corp., Carlsbad, CA) was used to isolate total RNA from the cells. The first strand of cDNA was transcribed using a reverse-transcription PCR kit (Takara, Tokyo, Japan). The mRNA abundance of *GLUT1* and *GLUT8* was analyzed by quantitative reverse-transcription PCR using SYBR Green as described before (Zhao et al., 2012). The relative mRNA changes of each gene were calculated using the $2^{-\Delta\Delta CT}$ (where CT = cycle threshold) method (Livak and Schmittgen, 2001) and normalized to β -actin mRNA in the same sample.

Data were analyzed by one-way ANOVA using SPSS software (version 16.0; IBM Corp., Armonk, NY), and Duncan's multiple range tests were used for multiple comparisons. Each experiment was performed with 3 to 6 replicates and repeated 3 times using pooled BMEC isolated from 3 Holstein lactating cows. Data from 1 representative experiment is reported here. Significant differences were considered at P < 0.05.

In this study, we used a PKC activator (PMA) and a PKC inhibitor (GF1090203X) to study the role of PKC in regulation of glucose uptake and *GLUT* expression in BMEC. Phorbol 12-myristate 13-acetate is a diester of phorbol, which has been widely used in biomedical research to activate classical PKC (Bosch et al., 2004; Goel et al., 2007) because of its structural similarity to one of the natural activators of classic PKC isoforms, diacylglycerol. Compared with the control group (0 ng/ mL), treatment of BMEC with 50 and 100 ng of PMA/ mL significantly increased the cellular glucose uptake (P < 0.05; Figure 1), whereas the glucose uptake in the cells treated with either the lowest concentration (25) ng/mL) or the highest concentration (200 ng/mL) of PMA did not show a significant difference (P > 0.05;Figure 1). In addition, when BMEC were treated with a pan-PKC inhibitor, GF1090203X (Roberts et al., 2005; Nguyen, 2008; Farese and Sajan, 2010), at the concentrations of 0.25, 0.5, 1, and 2 μM before PMA treatment (50 ng/mL), PMA-activated glucose uptake was significantly inhibited by GF1090203X at a concentration of 0.5 μM or higher (P < 0.05; Figure 2). These results demonstrate a regulatory role of PKC in glucose transport in BMEC, consistent with previous observations in adipocytes and intestinal cells (Gibbs et al., 1991; Navé et al., 1996; Zheng and Sarr, 2012).

Furthermore, we found that in BMEC, PKC regulates mRNA expression of *GLUT1* and *GLUT8*, the main glucose transporters expressed in the mammary gland (Zhao et al., 1996, 2004). In particular, 50 and 100 ng of PMA/mL significantly increased the mRNA expression of *GLUT1* and *GLUT8* in BMEC (P < 0.05; Figure 3A and 3B) compared with the control (0 ng of PMA/mL), whereas 25 and 200 ng/mL had no effect (P > 0.05). These observations are in line with the report



Figure 1. Glucose uptake in bovine mammary epithelial cells treated with different concentrations of phorbol 12-myristate 13-acetate (PMA). The glucose uptake in the control cells without PMA treatment (0 ng/mL) is assigned a value of 1.0. Values with different letters (a-c) are significantly different (P < 0.05; n = 6). Error bars indicate the SE.

that *GLUT1* mRNA levels were increased by activation of PKC in bovine brain capillary endothelial cells (Farrell et al., 1992) and suggest that the PKC-activated glucose uptake in BMEC may be mediated by enhanced expression of *GLUT1* and *GLUT8*. In addition, *GLUT8* mRNA was higher at 100 ng/mL of PMA compared with 50 ng/mL (P < 0.05; Figure 3B), whereas no difference was seen in *GLUT1* expression between these 2 groups (P > 0.05; Figure 3A). Similar to the glucose uptake, the PMA-induced *GLUT1* mRNA expression in BMEC was significantly inhibited by GF1090203X at a concentration of 0.5 μM or higher (P < 0.05; Figure 4A), whereas the PMA-induced *GLUT8* mRNA expression was significantly inhibited by GF1090203X at a



Figure 2. Effect of GF1090203X, an inhibitor of protein kinase C, on phorbol 12-myristate 13-acetate (PMA)-induced glucose uptake in bovine mammary epithelial cells. Cells were incubated in media with different concentrations of GF1090203X (0, 0.25, 0.5, 1, and 2 μM) for 30 min and then treated with or without PMA (50 ng/mL). The glucose uptake in the group without PMA and GF1090203X treatments is assigned a value of 1.0. Values with different letters (a–c) are significantly different (P < 0.05; n = 6). Error bars indicate the SE.

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Figure 3. The mRNA abundance of glucose transporter 1 (*GLUT1*; A) and *GLUT8* (B) genes in bovine mammary epithelial cells treated with different concentrations of phorbol 12-myristate 13-acetate (PMA). The mRNA levels of *GLUT1* and *GLUT8* in the cells without PMA treatment are assigned values of 1.0. Values with different letters (a–c) are significantly different (P < 0.05; n = 3). Error bars indicate the SE.

concentration of 0.25 μM or higher (P < 0.05; Figure 4B). It appeared that the mRNA expression of *GLUT8* in BMEC may be more sensitive to PKC activation than the expression of *GLUT1*.

Previous studies have shown that PKC activation induces activator protein-2 (AP-2)-mediated transcriptional activation (Imagawa et al., 1987; Hyman et al., 1989). Both phorbol ester response elements and AP-2 binding sites have been identified in the rat GLUT1 gene in liver cells (Behrooz and Ismail-Beigi, 1997). The GLUT1 gene is highly conserved across mammalian species (Zhao and Keating, 2007), and the rat phorbol ester response element and human AP-2 element binding sites have been found in the bovine GLUT1 gene. Therefore, it is likely that PKC may act on these sites directly or indirectly to activate GLUT1 expression in BMEC. The bovine GLUT8 gene has not been well characterized; thus, it is not known whether these elements are present in bovine GLUT8 gene.

We cannot rule out the possibility that PKC may stimulate glucose uptake by the mechanisms other



Figure 4. Effect of GF1090203X, an inhibitor of protein kinase C, on phorbol 12-myristate 13-acetate (PMA)-induced glucose transporter 1 (*GLUT1*) and *GLUT8* mRNA expression in bovine mammary epithelial cells. Cells were incubated in media with different concentrations of GF1090203X (0, 0.25, 0.5, 1, and 2 μ M) for 30 min and then treated with or without PMA (50 ng/mL). The *GLUT1* and *GLUT8* mRNA levels in the group without PMA and GF1090203X treatments were assigned values of 1.0. Values with different letters (a–d) are significantly different (P < 0.05; n = 3). Error bars indicate the SE.

than enhancing GLUT mRNA expression. Bosch et al. (2003) reported that PMA stimulates glucose uptake in 3T3-L1 adipocytes by increasing the GLUT1 protein content in the total cell homogenate. Moreover, the translocations of GLUT1 (Gibbs et al., 1991; Bosch et al., 2003) and GLUT4 in rat adipocytes (Standaert et al., 1997; Kotani et al., 1998; Watson and Pessin, 2001) and GLUT2 in rat intestinal IEC-6 cells (Zheng and Sarr, 2012) were observed in PKC-regulated glucose transport. Therefore, it is possible that PKC may also regulate protein translation, translocation, or transport kinetics of GLUT in BMEC to enhance glucose uptake. In summary, this study provided evidence that PKC may play a role in the regulation of glucose uptake and expression of GLUT1 and GLUT8 in BMEC.

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