AG and UAG induce β -casein expression via activation of ERK1/2 and AKT pathways

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Abstract

The ghrelin peptides were found to circulate in two major forms: acylated ghrelin (AG) and unacylated ghrelin (UAG). Previous studies showed that AG regulates β -casein (CSN2) expression in mammary epithelial cells. However, little is known about the mechanisms by which AG regulates CSN2 gene and protein expression. Evidence suggests that UAG has biological activity through GHSR1a-independent mechanisms. Here, we investigated the possible GHSR1a-mediated effect of UAG on the expression of CSN2 in primary bovine mammary epithelial cells (pbMECs) isolated from lactating cow. We found that both AG and UAG increase the expression of CSN2 in a dose-dependent manner in pbMECs in comparison with the control group. Increased expression of CSN2 was blocked by [D-Lys3]-GHRP-6 (an antagonist of the GHSR1a) and NF449 (a Gs- α subunit inhibitor) in pbMECs. In addition, both AG and UAG activated AKT/protein kinase B (AKT) and extracellular signal-regulated kinase 1/2 (ERK1/2) pathways, whereas [D-Lys3]-GHRP-6 and NF449 inhibited the phosphorylation of AKT and ERK1/2 in pbMECs respectively. Blockade of ERK1/2 and AKT signaling pathways prevented the expression of CSN2 induced by AG or UAG. Finally, we found that both AG and UAG cause cell proliferation through identical signaling pathways. Taken together, these results demonstrate that both AG and UAG act on ERK1/2 and AKT signaling pathways to facilitate the expression of CSN2 in a GHSR1a-dependent manner.

Key Words

- acylated ghrelin
- unacylated ghrelin
- β-casein
- bovine mammary epithelial cells

Journal of Molecular Endocrinology (2016) **56**, 213–225

Acylated (AG) and unacylated ghrelin (UAG) are circulating peptides encoded by the preproghrelin gene peptides and are mainly secreted by the X/A-like enteroendocrine cells of the stomach (Kojima *et al.* 1999, Muller *et al.* 2015). AG stimulates growth hormone (GH) release and positive energy balance by binding in the hypothalamus and pituitary gland to the only functional ghrelin receptor that has been characterized thus far, GH

secretagogue receptor 1a (GHSR1a) (Howard *et al.* 1996, Barnett *et al.* 2010, Muller *et al.* 2015). The presence of an acyl side chain (mainly *n*-octanoic acid) attached to the AG peptide is required for full agonism of GHSR1a (Barnett *et al.* 2010, Heppner *et al.* 2014, Kern *et al.* 2014).

In addition to the gut, AG and GHSR1a were expressed in mammary glands and cultured primary mammary epithelial cells (MECs) (Zhang *et al.* 2013,

Jedrzejczak & Szatkowska 2014, Ogorevc & Dovc 2015). Therefore, understanding the regulating actions of these peptides is of interest. AG has a specific action during lactation. In addition to increasing plasma insulin, food intake, and glucose concentrations (Itoh et al. 2006, Fukumori et al. 2013), AG also affects milk production and milk components in rats and dairy cows (Nakahara et al. 2003) and affects milk fat and protein synthesis in water buffaloes (Gil et al. 2013). β-casein (CSN2) is the most abundant protein in cow milk and is often used as a marker for mammary differentiation. CSN2 was previously shown to be expressed under in vitro conditions in different species, such as in several bovine mammary cell lines (Jedrzejczak & Szatkowska 2014), goat primary mammary cells (Zhang et al. 2013, Ogorevc & Dovc 2015), and mammary gland explants from cows. Northern blot analysis revealed that daily injection of AG significantly increased mammary gland CSN2 mRNA expression in rats (Nakahara et al. 2003). Zhang and colleagues showed that levels of CSN2 protein increased in MECs of lactating goat after treated with AG compared with control group (Zhang et al. 2013). However, the effects and mechanisms of AG in milk protein synthesis in primary bovine mammary epithelial cells (pbMECs) have not been examined.

A number of reports have been published on the regulation of AG secretion in the context of energy metabolism that did not distinguish between acyl-modified and des-acyl AG (ThidarMyint & Kuwayama 2008, Delhanty *et al.* 2014, Pinkney 2014). In contrast to AG, the regulating action of UAG in the expression of CSN2 remains poorly understood. In several studies, the effects of AG are mimicked by UAG (Granata *et al.* 2007, Reano *et al.* 2014). Later studies revealed a weak but full agonism of binding by UAG to GHSR1a (Gauna *et al.* 2007, Heppner *et al.* 2014). However, the potential role of UAG in the expression of CSN2 in pbMECs has yet to be identified. Moreover, whether UAG can activate a similar pathway to AG to regulate the expression of CSN2 is not clear.

G-protein-coupled receptors (GPCRs) are one of the major classes of cell surface receptors and are associated with a group of G-proteins consisting of three subunits termed alpha, beta, and gamma. G-proteins are classified into four families according to their α subunit: Gi alpha subunit (G α i), Gs alpha subunit (G α s), G12/G13 alpha subunits (G α 12/13), and Gq alpha subunit (G α q). Attempts to elucidate the peripheral effects of AG have identified several signaling mechanisms involving G-protein-dependent pathways, highlighting the complexity of GHSR1a activation (Evron et al. 2014, Kern et al. 2014, Yin et al. 2014). Depending on the tissues and cells, activation of GHSR1a may trigger a diverse signaling mechanism and subsequently distinct physiological responses. Both AG and UAG stimulate cell proliferation and prevent apoptosis via activation of the ERK1/2 and AKT pathways in different cells, such as pancreatic β-cells, skeletal muscle cells (Reano et al. 2014, Yu et al. 2014), and human endothelial cells (Xiang et al. 2011). In addition, both pbMEC number and activity can be affected by IGF-I signaling by activating the AKT and ERK1/2 pathways. Moreover, the AKT pathway is activated in response to milk yield (Murney et al. 2015). However, whether AG and UAG activate the ERK1/2 and AKT pathways and induce the expression of CSN2 via GHSR1a in pbMECs of lactating cows is not known.

The aim of this study was to evaluate the effects of AG and UAG on the expression of CSN2 in pbMECs of lactating cows. Furthermore, the signal pathways that are involved in the expression of CSN2 induced by both AG and UAG were also investigated.

Materials and methods

Reagents

Human AG (Ser3-Decanoyl), Human UAG (Ser3-Des-Octanoyl), and [D-Lys3]-GHRP-6 were obtained from Phoenix Pharmaceuticals, Inc. (Belmont, CA, USA). Antibodies to P-ERK 1/2 (Thr202/Tyr204), ERK1/2 (Thr202/ Tyr204), AKT(Ser473), and P-AKT (Ser473) were purchased from Cell Signaling Technology Inc. Antibodies to AG, GHSR1a, and CSN2 were purchased from Abcam. Bovine insulin, prolactin, hydrocortisone, phosphatase inhibitor cocktail 2, protease inhibitor cocktail, and DAPI were obtained from Sigma. The antibodies for goat anti-rabbit IgG-HRP, rabbit anti-cytokeratin 18 (CK-18), and rabbit anti-β-tubulin (1-19) were purchased from Santa Cruz Biotechnology. The selective inhibitors of NF449 (Gas), pertussis toxin (PTX, Gai), and U0126 (ERK 1/2) were obtained from TOCRIS (Ellisville, MO, USA). The selective inhibitor of YM-254890 (Gq11) was provided by Astellas Pharma Inc. (Tokyo, Japan). The selective inhibitor of MK2206 (AKT) was from Selleckchem (Houston, DE, USA).

PbMECs cultures and treatments

PbMECs from the alveolar tissue of four mid-lactation (d 150–200 after parturition) from healthy Holstein cows were isolated using a procedure described previously

(Sorg *et al.* 2012, Jedrzejczak & Szatkowska 2014). Cells were maintained in DMEM-F12, supplemented with 10% fetal bovine serum, penicillin ($100 \mu/mL$), streptomycin (100 mg/mL), bovine insulin ($5 \mu g/mL$), hydrocortisone ($1 \mu g/mL$), and prolactin ($1 \mu g/mL$). The expression of CK-18 was identified as a marker of pbMECs by both immunofluorescence and flow cytometry. The gene expression of *CSN2*, ghrelin (*GHRL*), and *GHSR1a* were detected by quantitative RT-PCR and western blot in pbMECs from passages 1–4.

The cells were treated with increasing concentrations of AG or UAG (0, 0.01, 0.1, 1, 10, and 100 ng/mL) for 12 or 24 h. The cells were then collected and frozen at -80° C until mRNA or protein extraction. For some experiments, the cells were pre-incubated with inhibitors for 1 h, and then treated for different time periods with AG or UAG.

Quantitative RT-PCR

After treatment with AG or UAG for different time periods, the pbMECs were harvested. Total RNA was extracted from pbMECs and reversed transcribed into cDNA from 4 µg total RNA, as described previously (Li *et al.* 2013). The mRNA levels of various genes were evaluated using the SYBR Green QuantiTect RT-PCR Kit. The comparative $2^{-\Delta\Delta ct}$ method was used to calculate the relative gene expression levels of *CSN2*, *GHRL*, and *GHSR1a*, with β-actin serving as the housekeeping gene. The PCR products were visualized via 1.5% agarose gel electrophoresis. The following primers were used: GHSR1a, 5'-CCGACAATGACTCGCT GACT-3' and 5'-CAAGTTCCAGGGCCGGTAAT-3'; GHRL, 5'-ATGCCCGCCCGTGGACCAT-3' and 5'-AACCGGAT TTCCAGCTCGTC-3'; CSN2, 5'-GGCTATGGCTCCTAAGC



Figure 1

Cultured and detection of pbMECs. (A) Morphology of cultured pbMECs. Islands of epithelial cells are visible (bar = 1000 µm). (B) The expression of CK-18 in pbMECs was studied by flow cytometry. (C) Immunofluorescence labeling of CK-18 in pbMECs (200x magnification). (D) Gene expression of *CNS2*, *GHSR1a*, and *GHRL* in cultured pbMECs (passages 1–4). (E) Protein expression of CSN2, AG, and GHSR1a in pbMECs (passages 1–4). A full colour version of this figure is available at http://dx.doi.org/10.1530/ JME-15-0287.

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ACA-3' and 5'-GAGAAAGGGACAGCACGGAC-3'; and β -actin, 5'-TCACCAACTGGGACGACA-3' and 5'-GCATA CAGGGACAGCACA-3'.

Western blot analysis

After treatment with AG or UAG for different time periods, whole-cell proteins were extracted in accordance with our published protocol (Li et al. 2013). The cells were washed with ice-cold PBS and harvested using RIPA lysis buffer containing 10mM HEPES, pH 7.5; 10mM KCl; 0.1 mM EGTA; 0.1 mM EDTA; phosphatase inhibitor cocktail 2, and protease inhibitor cocktail. Protein samples (40µg) were subjected to 12% SDS-PAGE and transferred to PVDF membrane (Chemicon International, Millipore, Billerica, MA, USA). After blocking with 5% milk, the membranes were incubated overnight at 4°C with the specific antibodies to P-ERK1/2 (1:1000), ERK1/2 (1:2000), AKT (1:2000), P-AKT (1:1000), GHSR1a (1:4000), CSN2 (1:1000), AG (1:500), or β-tubulin (1:10,000). The membranes were washed four times with 0.1% TBS-T and incubated with goat anti-rabbit IgG-HRP for 1 h at room temperature (RT). The immunoreactive proteins were visualized using the ECL detection system (Millipore).

Cell viability assay

Cells were seeded in 96-well plates at a density of 5×10^4 cells/well and treated with AG or UAG (0, 0.01, 0.1, 1, 10, and 100 ng/mL) for 24, 48, and 72 h. After treatment, the cells were incubated with 1 mg/mL MTT for approximately 4 h and then incubated with 150 µL of dimethylsulfoxide for 10 min. The optical density of the samples was quantified by measuring the absorbance at 570 nm and expressed as the percentage relative to the untreated control. Each experiment included five replicates per treatment, and the experiments were repeated a minimum of three times. For some experiments, the cells were pre-incubated with inhibitors for 1 h and then treated for 24 h with AG or UAG.

Immunofluorescence assay

To evaluate the expression of CK-18 in pbMECs, 1×10^4 cells/well were seeded in a slide in 24-well plates for 24 h. Slides were fixed with 4% paraformaldehyde in PBS for 10 min at RT. After blocking with 3% normal goat serum for 2h at RT, the slides were incubated with primary

http://jme.endocrinology-journals.org DOI: 10.1530/JME-15-0287 antibodies to CK-18 (1:200) overnight at 4°C. After washing, the slides were incubated with PE-conjugated goat anti-rabbit IgG (1:500) for 1 h at RT. DAPI was used to stain the cell nuclei at a concentration of $1.43 \,\mu$ M for 10 min. After staining, slides were viewed with an inverted fluorescence microscope.

Flow cytometry assay

To evaluate the population of CK-18-expressing pbMECs, the cells were seeded into 6-well plates at a density of 1×10^6 cells/well. After 24h, the cells were aspirated and



Figure 2

AG and UAG induce CSN2 gene and protein expression in pbMECs. (A) Cells were incubated for 12 h with increasing concentrations of AG or UAG. Gene expression of CSN2 was determined by quantitative RT-PCR. (B) Cells were incubated for 24 h with increasing concentrations of AG or UAG. Protein expression of CSN2 was examined by western blot. (C and D) The data are presented as the mean \pm s.b. (n > 3 independent experiments; *P<0.05, **P<0.01 vs non-treated (NT) control group).

washed with PBS twice. After blocking, the cells were stained with CK-18 (1:500) for 16h at 4°C. The cells were incubated with donkey anti-rabbit IgG-FITC for 1 h at RT after washing. Flow cytometry (BD Biosciences, USA) was used to determine the population of CK-18⁺ cells.

Statistical analysis

The data were analyzed using GraphPad Prism 5 (GraphPad InStat Software, San Diego, CA, USA). The comparison among groups was carried with an ANOVA followed by Dunnett's test. In each panel, experiments were repeated with similar results at least three times. Data are presented as the mean \pm s.D. A *P*-value of 0.05 or 0.01 was considered statistically significant.

Results

AG and UAG promote the expression of CSN2 in pbMECs

We first detected the morphology (Fig. 1A) and the expression of CK-18 before each experiment to ensure the purity of pbMECs. Cells from a population with 99% CK-18 expression were used in the subsequent

experiments (Fig. 1B and C). We next detected the expression of CSN2, AG, and GHSR1a in cultured pbMECs from passage 1 to passage 4. The results showed that genes of *CSN2*, *GHRL*, and *GHSR1a* were expressed at passages 1–4, although the gene level was lower at passage 1 and passage 2 (Fig. 1D). Moreover, the proteins of CSN2, AG, and GHSR1a were also stably expressed in pbMECs at passages 1–4 (Fig. 1E).

Previous studies showed that AG increases CSN2 mRNA expression in MECs from dairy goats (Zhang et al. 2013). However, whether AG influences the expression of CSN2 in pbMECs is not known. In addition, UAG is a weaker agonist of GHSR1a compared with AG (Gauna et al. 2007). Therefore, we hypothesized that treatment with UAG has similar but maybe less potent effects than AG on the expression of CSN2. To test this hypothesis, the gene and protein expression of CSN2 were detected after treatment for 12 or 24h with increasing concentrations of either AG or UAG in pbMECs. The results showed that compared with the non-treated (NT) control group, only 10 ng/mL AG or UAG significantly increased CSN2 gene expression (Fig. 2A, B and C). Western blot results showed that the expression of CSN2 protein was significantly higher in these pbMECs treated with AG (1-100 ng/mL)



Figure 3

AG and UAG increased the expression of CSN2 in a GHSR1a-dependent manner in pbMECs. The cells were pre-treated for 1 h with [D-Lys3]-GHRP-6 (100 µg/mL) and then incubated for 12 or 24 h with AG or UAG. The gene and protein expression of CSN2 was examined by (A and B) quantitative RT-PCR and (C and D) western blot. (E, F, G, and H) The data are presented as the mean \pm s.D. (*n*>3 independent experiments; **P*<0.01 vs NT; **P*<0.05, ***P*<0.01 vs AG or UAG treatment).

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and peaked at 10 ng/mL and then decreased at 100 ng/mL (P<0.01; Fig. 2B, C and D). For UAG treatment at 10 ng/mL and 100 ng/mL, significantly higher values of protein expression of CSN2 were observed (P<0.01; Fig. 2B, C and D). In addition, we also investigated whether AG and UAG influences the expression of GHSR1a in pbMECs. Our results showed that both AG and UAG have no significant effect on GHSR1a gene and protein expression (Supplementary Figure 1, see section on supplementary data given at the end of this article). Considering the above results, 10 ng/mL was selected as the best concentration for AG and UAG in subsequent experiments.

GHSR1a and $G\alpha s$ are required for AG- and UAG-induced expression of CSN2 in pbMECs

Previous research reported that both AG and UAG regulate peripheral glucose metabolism through a GHSR1a-dependent mechanism (Heppner *et al.* 2014).

[D-Lys3]-GHRP-6 is regarded as a highly selective GHSR1a antagonist and has been widely used to investigate the dependency of GHSR1a signaling (Granata et al. 2007, Zhang et al. 2013). To determine whether the effect of AG and UAG on the expression of CSN2 depends on GHSR1a activation, pbMECs were pre-incubated with [D-Lys3]-GHRP-6 (100µg/mL) for 1h, followed with AG or UAG for 24h. The results showed that both AG and UAG significantly increased gene and protein expression of CSN2 in pbMECs compared with the NT control group (Fig. 3A, B, C, D, E, F, G, and H). However, the increased gene level of CSN2 was inhibited by [D-Lys3]-GHRP-6 (*P*<0.01; Fig. 3A, B, E, and F). Moreover, increased protein level of CSN2 was also suppressed by [D-Lys3]-GHRP-6 (*P*<0.01; Fig. 3C, D, G, and H). However, stimulation with [D-Lys3]-GHRP-6 alone had no significant effects on the expression of CSN2 compared with the NT control group.

AG binding to GHSR1a has been shown to activate intracellular second messengers coupled to $G\alpha s$, $G\alpha i/o$,



Figure 4

Gas is coupled to GHSR and is required for AG- and UAG-induced expression of CSN2 in pbMECs. The cells were pre-treated for 1 h with NF449 (10 μ M) and then incubated for 12 or 24 h with AG or UAG. The gene and protein expression of CSN2 was examined by (A and B) quantitative RT-PCR and (C and D) western blot. (E, F, G, and H) The data are presented as the mean ±s.p. (n > 3 independent experiments; #P<0.01 vs NT; *P<0.05, **P<0.01 vs AG or UAG treatment).

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Figure 5

AG and UAG induce ERK1/2 and AKT phosphorylation in GHSR-dependent manner in pbMECs. Cultured pbMECs were incubated with either AG or UAG in the presence or absence of [D-Lys3]-GHRP-6 (100 µg/mL) for 5, 30, and 60 min. At the indicated time points, cell lysates were prepared and subjected to western blotting using ERK1/2, P-ERK1/2, AKT, and P-AKT antibodies. (A and B) AG- and UAG-induced phosphorylation of ERK1/2 and AKT were abolished upon treatment with [D-Lys3]-GHRP-6. (C, D, E, and F) Bar graph showing mean of P-ERK1/2 and P-AKT levels. (n > 3 independent experiments; #P<0.01 vs NT; *P<0.05, **P<0.01 vs AG or UAG treatment).

and $G\alpha q/11$ and enhanced GH release, appetite, and proliferation (Granata et al. 2007, Evron et al. 2014, Reano et al. 2014). GHSR1a couples to different intracellular signaling pathways and mediates different effects depending on the cell type. However, which G-protein subunits couple to GHSR1a to mediate the effect of AG or UAG in pbMECs remains unknown. To investigate

this mechanism, the cells were pre-incubated for 1h with NF449 (10µM), YM254890 (10µM), and PTX (10µM), which are selective G α s, Gq/11, and Gi/o protein-coupled receptor antagonists respectively. Next, the cells were incubated for 12 or 24h with AG or UAG. Pre-incubation with NF449 resulted in a complete blockade of both AG- and UAG-induced gene and protein expression of



Figure 6

Gas is required for ERK1/2 and AKT phosphorylation induced by AG or UAG in pbMECs. Cells were incubated with either AG or UAG in the presence or absence of NF449 (10 µM) for 5, 30, and 60 min. At the indicated time points, cell lysates were prepared and subjected to western blotting using ERK1/2, P-ERK1/2, AKT, and P-AKT antibodies. (A and B) Phosphorylation of ERK1/2 and AKT activated by AG or UAG was inhibited by NF449, a Gas subunit-selective G-protein antagonist. (C, D, E, and F) Bar graph showing mean of P-ERK1/2 and P-AKT levels. (n>3 independent experiments; #P<0.01 vs NT; *P<0.05, **P<0.01 vs AG or UAG treatment).

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CSN2 (P<0.01; Fig. 4A, B, C, D, E, F, G, and H). However, treatment with PTX and YM-254890 had no effect on both AG- and UAG-induced mRNA and protein expression of CSN2 in pbMECs (Supplementary Figure 2). These results implied that activation of GHSR1a by AG and UAG is likely coupled to G α s, not Gq/11 and Gi/o, and promotes the expression of CSN2 in pbMECs.

AG and UAG induce ERK1/2 and AKT phosphorylation in pbMECs via activation of GHSR1a and G α s

Recent studies have demonstrated that under defined conditions, GHSR1a couples to $G\alpha$ s, resulting in activation of phospho-ERK1/2 and phospho-AKT (Granata *et al.* 2007, Evron *et al.* 2014, Kern *et al.* 2014). Thus, we next assessed the potential role of GHSR1a in regulating the activation of ERK1/2 and AKT signaling in pbMECs. The cells were pre-incubated with [D-Lys3]-GHRP-6 for 1h and followed with AG or UAG for 5, 30, and 60 min. After treatment with AG, the phosphorylation of ERK1/2 and AKT was significantly increased compared with the NT

control group but suppressed after pre-incubation with [D-Lys3]-GHRP-6 at 60 min for phospho-ERK1/2, or 5 and 60 min for phospho-AKT, respectively (P < 0.01; Fig. 5A, C, and D). In addition, the increased phosphorylation of ERK1/2 and AKT induced by UAG was also blocked by [D-Lys3]-GHRP-6 at 5 min for phospho-ERK1/2 and phospho-AKT compared to with the NT control group, respectively (P < 0.01; Fig. 5B, E, and F).

Our previous results showed that AG- and UAGinduced expression of CSN2 involves a G α s protein-coupled receptor, which, in turn, has been shown to activate ERK1/2 and AKT signaling (Granata *et al.* 2007, Yu *et al.* 2014). Therefore, we investigated whether these pathways mediate the increased expression of CSN2 induced by AG or UAG. In agreement, NF449 also inhibited AG-induced ERK1/2 phosphorylation at 30 and 60 min (P < 0.01; Fig. 6A and C), and increased phospho-AKT was also suppressed by NF449 at 5, 30, and 60 min (P < 0.01; Fig. 6A and D). In addition, stimulation of pbMECs with UAG leads to significantly increased phosphorylation of ERK1/2, which is reduced by NF449 at 5, 30, and 60 min (P < 0.01;



Figure 7

AG and UAG induce the expression of CSN2 through the ERK1/2 signaling pathway in pbMECs. The cells were pre-treated for 1 h with U0126 (10 μ M), followed by AG or UAG for 12 or 24 h. The gene and protein expression of CSN2 was examined by (A and B) quantitative RT-PCR and (C and D) western blot. (E, F, G and H) The data are presented as the mean ±s.D. (*n*>3 independent experiments; **P*<0.01 vs NT; **P*<0.05, ***P*<0.01 vs AG or UAG treatment).

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Fig. 6B and E). Moreover, induced phospho-AKT was also decreased at 30 min (P<0.01; Fig. 6B and F); however, no effect was found using inhibitors alone. These results suggested that GHSR1a and G α s are required for AG- and UAG-induced phosphorylation of ERK1/2 and AKT in pbMECs.

AG and UAG stimulate the expression of CSN2 by activating ERK1/2 and AKT signaling

To further determine the potential role of ERK1/2 and AKT pathways in AG- and UAG-induced expression of CSN2 in pbMECs, the cells were pre-incubated for 1h with U0126 and MK2206, which are specific inhibitors of ERK1/2 and AKT respectively. Our results showed that treatment with U0126 resulted in a significant blockade of ERK1/2 phosphorylation (data not shown). The gene and protein levels of CSN2 induced by AG and UAG were also inhibited by U0126 in pbMECs (P<0.01; Fig. 7). In addition, stimulated with MK2206 for 1h completely blockaded the phosphorylation of AKT in pbMECs

(data not shown). Increased gene and protein expression of CSN2 was also suppressed by MK2206 in pbMECs (P<0.01; Fig. 8). However, stimulation with U0126 and MK2206 alone had no significant effects on the expression of CSN2 compared with the NT control group. Together, these results demonstrated that AG- and UAG-induced expression of CSN2 might depend on the phosphorylation of ERK1/2 and AKT in pbMECs.

AG and UAG stimulate cell proliferation in pbMECs

Both AG and UAG not only stimulated cell proliferation but also suppressed apoptosis in pancreatic β -cells, skeletal muscle, and macrophages (Gauna *et al.* 2007, Evron *et al.* 2014, Li *et al.* 2015). Thus, finally, we investigated the effects of increasing the concentration of AG and UAG treatment on the growth of pbMECs. As shown in Fig. 9, compared with the NT control group, pbMECs proliferation was enhanced following increased AG stimulation for 24, 48, and 72h, with the maximum



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Figure 8

AG and UAG induce the expression of CSN2

through the AKT signaling pathway in pbMECs.

The cells were pre-treated for 1 h with MK2206

(10 μ M), followed by AG or UAG for 12 or 24 h. The gene and protein expression of CSN2 were examined by (E and F) quantitative RT-PCR and (A and B) western blot (C and D). (E, F, G, and H) The data are presented as the mean ±s.D. (n>3 independent experiments; *P<0.01 vs NT; *P<0.05, **P<0.01 vs AG or UAG treatment).



Figure 9

AG and UAG induce pbMEC proliferation. PbMECs were cultured with increasing concentrations of AG or UAG for 24, 48, and 72 h with or without AG or UAG. Then, cell proliferation was measured in a MTT assay. (A and B) AG and UAG markedly enhanced cell proliferation at 24, 48, or 72 h. The cells were pre-treated for 1 h with D-Lys, NF449, U0126, or MK2206, followed by AG or UAG for 24 and 48 h. (C and D) AG- and UAG-induced cell proliferation were inhibited by D-Lys, NF449, U0126, or MK2206. Values are presented as the mean \pm s.p. (n > 3 independent experiments; #P<0.01 vs NT; *P<0.05, **P<0.01 vs AG or UAG treatment).

proliferative response at 100 ng/mL (P<0.05; Fig. 9A). In addition, UAG significantly induced cell proliferation in a dose-dependent manner at 24 h, and was lower but still above basal level after 48 and 72 h (P < 0.05; Fig. 9B). These results suggested that cell proliferation was markedly enhanced as the concentration of AG and UAG. However, there were no cytotoxic effects on the growth of pbMECs.

To examine whether GHSR1a and Gas mediate AGand UAG-induced proliferation in pbMECs, the cells

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were pre-incubated with [D-Lys3]-GHRP-6 and NF449 for 1h followed by incubation with AG or UAG. The results showed that pre-incubation with [D-Lys3]-GHRP-6 or NF449 remarkably reduced AG- or UAG-induced proliferation in pbMECs compared with AG or UAG group (P < 0.05; Fig. 9C and D). To elucidate whether activation of the ERK1/2 and AKT pathways are required to AG- and UAG-induced proliferation, we assessed the effect of U0126 and MK2206. Both AG- and UAG-induced cell proliferations were blocked by U0126 or MK2206 in pbMECs compared with AG group (P < 0.05; Fig. 9C and D). In addition, stimulation with [D-Lys3]-GHRP-6, NF449, U0126, and MK2206 alone had no significant effect on cell proliferation compared with the NT control group. These results showed that cell proliferation induced by AG or UAG was likely mediated through a GHSR1a and Gas signal cascade.

Discussion

Our findings show that both AG and UAG increase the gene and protein expression of CSN2 in pbMECs of lactating cows. The ERK1/2 and AKT signal pathway that is activated by GHSR1a and $G\alpha$ s likely mediates these effects.

The initiation of lactation dramatically alters the metabolism of many organs so that the mammary glands can be supplied with the nutrients necessary for milk protein synthesis (Yu et al. 2015). Protein production is controlled by the complex interplay of peptide and steroid hormones, predominantly the lactogenic hormones: prolactin, insulin, and hydrocortisone. More recently, other hormones have been discovered in breast milk such as AG (Savino et al. 2012). The third amino acid residue of AG, serine (Ser3), is modified by an acyl group; this modification is essential for AG biological activity. In addition to n-octanovl AG, other forms of AG peptide exist, including UAG, which lacks an acyl modification, and other acylated AG species, such as n-decanoyl AG, whose Ser3 residue is modified by n-decanoic acid (Yoh et al. 2011). It has been shown that n-decanoyl AG was one of the active forms of the AG peptide and that it could interact with the receptor in a manner similar to that of *n*-octanoyl AG. A number of reports on the regulation of AG secretion in the context of energy metabolism and food intake in cows that did not distinguish between acyl-modified and des-acyl AG have been published (Itoh et al. 2006, ThidarMyint et al. 2006, Bradford & Allen 2008, Roche et al. 2008, ThidarMyint & Kuwayama 2008, Fukumori et al. 2013). In addition, the effects of n-octanoyl

AG on plasma concentrations of glucose and milk yield and quality in lactation cows have been described in detail (Itoh *et al.* 2006, Roche *et al.* 2008, Gil *et al.* 2013). However, the effects and mechanisms of *n*-octanoyl AG and UAG in milk protein synthesis *in vitro* have not been examined. Thus, *n*-octanoyl AG and UAG were used in our experiment.

We first evaluated the effects of AG on the expression of CSN2 in pbMECs of lactation cows. With the exception of the epithelial cells of alveoli and ducts throughout the lactation in dairy goats (Zhang et al. 2013), our results demonstrated that AG and GHSR1a were expressed in pbMECs. We also find that only 10 ng/mL AG significantly increased the gene expression of CSN2. Western blot results showed that the expression of CSN2 protein was significantly higher in these pbMECs treated with a concentration range of 1-100 ng/mL compared with NT control group. Previous studies suggested that unlike AG, UAG is not acylated at Ser3 (Delhanty et al. 2013, 2014), and its effect on the expression of CSN2 has not been elucidated. However, here we show for the first time that UAG, similar to AG, significantly stimulated the gene expression of CSN2. Although the expression of CSN2 mRNA after treatment with UAG exhibited a similar tendency with AG-treated group, there are some different effects between AG and UAG. We observed significantly increased protein expression of CSN2 when treated with 1 ng/mL of AG compared with NT control group; however, there is no significantly increased when treated with UAG. Furthermore, highest protein expression of CSN2 was achieved in 10 ng/mL of AG, whereas 100 ng/mL for UAG. The cause of this results maybe that UAG is a full agonist of the GHSR1a, with activity in the high nanomolar range (Gauna et al. 2007). In a word, our results suggested that AG and UAG could directly modulate the expression of CSN2 in the cultured pbMECs, this is in agreement with findings in goat (Zhang et al. 2013).

The specific receptor mediating the signaling responses to AG or UAG in pbMECs is unknown. Cell-based assays have demonstrated that AG requires acylation at the Ser3 residue to activate GHSR1a at concentrations within the low-nanomolar range (Heppner *et al.* 2014). This evidence led to the assumption that the biological effects of UAG are independent of GHSR1a activation. However, a more recent study reported that UAG interacts with GHSR1a at concentrations within the high-nanomolar or micromolar range. Although some data indicate that UAG is a weak GHSR1a agonist, other data suggest that UAG is actually a full agonist of GHSR1a and induces similar maximal receptor stimulation as AG, such as the regulation of cell proliferation and glucose metabolism (Chung *et al.* 2013, Pinkney 2014). Thus, we investigated whether the effects of AG or UAG on the expression of CSN2 were dependent on GHSR1a. Our results showed that both AG- and UAG-induced expression of CSN2 were significantly blocked by [D-Lys3]-GHRP-6, an inhibitor of GHSR1a. Altogether, our *in vitro* results are consistent with the literature and suggest that UAG is an agonist of GHSR1a and regulates the expression of CSN2 in pbMECs.

GHSR1a belongs to the class A G-protein-coupled receptor (GPCR) family. When GHSR1a binds to the GPCR, a conformational change occurs in the GPCR, which allows it to activate an associated G-protein such as G α s, G α i/o, or G α q/11 (Evron *et al.* 2014). Here, we have investigated which G-protein subunit couples to GHSR1a in AG- and UAG-induced expression of CSN2 in pbMECs. We find that only NF449, the specific inhibitor of G α s, significantly reduced both AG and UAG-induced expression of CSN2 at both the gene and protein levels in pbMECs, whereas PTX and YM-254890 had no effect (data not shown). This result revealed the selective coupling of activated GHSR1a to G-protein subtypes.

Previous studies demonstrated that AG is correlated with the expression of CSN2 in vivo or in vitro in different species (Gil et al. 2013, Nakahara et al. 2003, Zhang et al. 2013). However, the mechanisms of this correlation are undefined. Downstream signaling of GHSR1a is mediated by several signaling pathways including the receptors ERK1/2 and AKT (Gauna et al. 2007, Xiang et al. 2011, Evron et al. 2014, Reano et al. 2014). It is well known that ERK1/2 and AKT play important roles in the regulation of cell survival (Xiang et al. 2011). AG- or UAG-induced activation of ERK1/2 and AKT is observed in many cell types, such as pancreatic β -cell line HIT-T15 (Evron *et al.* 2014), pancreatic β -cells, and macrophages (Li *et al.* 2015). To uncover the molecular mechanism underlying the tissue-specific response in GHSR1a-mediated intracellular signaling, we investigated the phosphorylation of ERK1/2 and AKT. In agreement with previous findings, stimulation by AG or UAG induces significant activation of ERK1/2 and AKT compared with NT control group (Granata et al. 2007). However, increased phosphorylation of ERK1/2 or AKT induced by AG or UAG was suppressed after pre-incubated with the inhibitor of GHSR1a, [D-Lys3]-GHRP-6. Moreover, previous research demonstrated that G-protein signaling cascades play an important role in prolactin-induced synthesis of milk products (Howard et al. 1996, Larrea et al. 1999). AG binding to GHSR1a has

http://jme.endocrinology-journals.org DOI: 10.1530/JME-15-0287 © 2016 Society for Endocrinology Printed in Great Britain

been shown to activate intracellular second messengers coupled to $G\alpha$ s, $G\alpha$ /o, and $G\alpha$ q/11, leading to activation of AKT, ERK1/2, AMPK, or mTOR signal pathways (Xiang *et al.* 2011). GHSR1a couples to different intracellular signaling pathways depending on the cell type. Here, our results showed that only treatment with NF449 for 1h inhibited the phosphorylation of ERK1/2 and AKT induced by AG or UAG in pbMECs. This observation is consistent with the report of Granata that G\alphas is required for AG- or UAG-induced phosphorylation of ERK1/2 and AKT (Granata *et al.* 2007).

To further determine whether the ERK1/2 and AKT pathway are required for the expression of CSN2 induced by AG or UAG, we examined the effect of U0126 and MK2206, which are specific inhibitors of ERK1/2 and AKT respectively. The chemical inhibition of ERK1/2 and AKT resulted in a significant suppression of AG- and UAG-induced expression of CSN2 in pbMECs, indicating that both AG and UAG stimulated the expression of CSN2 via activation of the ERK1/2 and AKT signaling pathways. Although previous studies indicate that mTOR and STAT5 signaling pathways play an important role in milk protein synthesis in the lactating cows (Kojima et al. 1999), here we are the first to demonstrate that activation of GHSR1a by AG or UAG may couple to G α s and signals to ERK1/2 and AKT to increase CSN2 expression in pbMECs.

It has shown that both AG and UAG promote cell proliferation and inhibit apoptosis in pancreatic β -cells (Granata et al. 2007), endothelial cells, and skeletal muscle cell (Yu et al. 2014). Milk proteins are synthesized and excreted by MECs during lactation. We presume that the milk protein synthesis within the MECs may be regulated through cell proliferation. In agreement, our results also proved that both AG and UAG stimulate pbMECs proliferation in a dose-dependent manner at 24h, even for 48h. Pre-treatment with [D-Lys3]-GHRP-6 and NF449 abolishes AG- or UAG-induced cell proliferation in pbMECs. In addition, the ERK1/2 and AKT inhibitors U0126 and MK2206, respectively, also suppress cell proliferation induced by AG or UAG. These results are consistent with previous reports that activation of the ERK1/2 and AKT pathways regulate pbMEC number and activity (Murney et al. 2015).

Collectively, our results demonstrate that GHSR1a is a G α s-coupled receptor involved in the AG- and UAGinduced expression of CSN2 in pbMECs and that increased the expression of CSN2 induced by either AG or UAG are likely mediated by the phosphorylation of ERK1/2 and AKT signaling pathways.

Supplementary data

This is linked to the online version of the paper at http://dx.doi.org/10.1530/ JME-15-0287

Declaration of interest

The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

Funding

This work was funded by the National Natural Science Foundation of China (project number 31372396 and 31572479), the Jilin Scientific and Technological Development Program (project no. 20130206036NY), and the Graduation Innovation Fund of Jilin University (project number 2015006), "985 Project" of Jilin University, Graduate Innovation Fund of Jilin University (project no. 2015861220).

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Received in final form 23 January 2016 Accepted 10 February 2016 Accepted Preprint published online 12 February 2016