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2 **Fermented Colostrum Whey Upregulates Aquaporin-3 Expression in, and**  
3 **Proliferation of, Keratinocytes via p38/c-Jun N-terminal kinase Activation**

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19 **Running title:** Fermented colostrum whey upregulates AQP-3 levels

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78        **Fermented Colostrum Whey Upregulates Aquaporin-3 Expression in, and**  
79        **Proliferation of, Keratinocytes via p38/c-Jun N-terminal Kinase Activation**

80  
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82        **Abstract**

83        Colostrum, which contains various immune and growth factors, aids wound healing by  
84        promoting keratinocyte proliferation. Aquaporins (AQPs) are small, hydrophobic  
85        membrane proteins that regulate cellular water retention. However, few studies have  
86        examined the effect of processed colostrum whey on AQP-3 expression in human skin  
87        cells. Here, we investigated the effect of milk, colostrum, fermented milk, and fermented  
88        colostrum whey on *AQP3* expression in keratinocyte HaCaT cells. Concentrations of 100-  
89        400 µg/mL of fermented colostrum whey were found to induce HaCaT cell proliferation.  
90        AQP3 was found to be expressed exclusively in HaCaT cells. *AQP3* expression was  
91        significantly increased in 100 µg/mL fermented colostrum whey-treated cells compared  
92        with that in controls. Moreover, fermented colostrum increased p38 mitogen-activated  
93        protein kinase (MAPK) and c-Jun N-terminal kinase (JNK) phosphorylation, but not  
94        ERK1/2 phosphorylation. Thus, our results suggest that fermented colostrum whey  
95        increased AQP-3 expression in, and the proliferation of, keratinocytes via JNK and p38  
96        MAPK activation.

97  
98        **Keywords:** Colostrum, fermented colostrum, aquaporin-3, keratinocyte, skin disorder

99

## 100 **Introduction**

101 The skin is composed of a highly distinct, multilayered epithelium, which contains  
102 approximately 30% of the total water content of the body (Madison, 2013; Sougrat, et al.,  
103 2002). The stratum corneum, the outermost layer of the skin, serves as the primary line  
104 of defense against the external environment, and maintains homeostasis by regulating the  
105 evaporation of water (Jacobi, 1959). The moisture present in the stratum corneum  
106 contains various physiologically active substances that are necessary for the human body  
107 and help the skin remain smooth and moist; this is important for the healing of various  
108 types of skin damage (Greaves and Søndergaard, 1970; Verkman et al., 2008). A failure  
109 in the ability to maintain water homeostasis in the skin results in various skin diseases,  
110 such as atopic dermatitis (Sator et al., 2008; Segre, 2006), psoriasis (Ghadially, 1996),  
111 and wound-healing delay (Gallant-Behm and Mustoe, 2010).

112 Aquaporin (AQP) is a small, hydrophobic membrane protein that transports water  
113 molecules into cells, and thus regulates the process of water retention in cells (Agre et al.,  
114 2002; Hara-Chikuma and Verkman, 2008). Till date, 13 mammalian AQPs (AQP-0 to  
115 AQP-12) have been cloned; these can be classified into different categories, depending  
116 on their transport capacity. AQP-0, AQP-1, AQP-2, AQP-4, and AQP-5 selectively  
117 transport water; however, AQP-3, AQP-7, AQP-9, and AQP-10 transport water and small  
118 solutes (such as glycerol) (Agre et al., 2002; Verkman, 2011). Several types of AQPs are  
119 reportedly present in skin and skin appendages. Among these, AQP-3 forms the water and  
120 glycerol channels present in the basal layer keratinocytes of the skin epidermis (Agre et  
121 al., 2002; Hara-Chikuma and Verkman, 2005), and is crucial for the process of skin  
122 moisturization (Hara et al., 2002; Ma et al., 2002; Sougrat et al., 2002). In addition, AQP-  
123 3 is important for wound healing, as it promotes epithelial cell migration and proliferation

124 (Hara et al., 2002; Hara-Chikuma and Verkman, 2008b; Verkman et al., 2008). AQP-3  
125 knockout mice show severe skin disorders, such as impaired skin hydration, decreased  
126 wound-healing ability, and reduced skin elasticity (Hara et al., 2002; Ma et al., 2002).  
127 Therefore, the stimulation of AQP-3 expression in keratinocytes provides effective levels  
128 of epidermal hydration and could be used to effectively improve the procedures used for  
129 skin disease treatment. Whey is produced as a by-product of the rennet-type cheese  
130 manufacturing process and is collected after the coagulation of casein and fat in fermented  
131 milk. Whey accounts for 85–90% of the total milk volume and retains soluble whey  
132 proteins, growth factors, lactose, and minerals, along with microbial fermentation  
133 metabolites (Smithers, 2008; Yadav et al., 2015). Furthermore, the components of whey  
134 regulate the differentiation of cultured human epidermal keratinocytes, while whey  
135 peptides improve the wound-healing process in the skin (Baba et al, 2006; Wang et al.,  
136 2010).

137 Colostrum refers to the milk secreted within 72 h of delivery (Thapa, 2005). Colostrum  
138 is composed of three primary components: nutrients (Macy, 1949), immune substances  
139 (Ogra et al., 1997), and growth factors (Pakkanen and Aalto, 1997; Seo et al., 2018).  
140 Functionally, colostrum is known to enhance immune function and eliminate bacteria,  
141 toxins, and allergens (Zimecki and Kruzel, 2007). Notably, the growth factors in the  
142 colostrum play an important role in wound healing by promoting keratinocyte and  
143 fibroblast proliferation and migration (Chen et al, 2014; Hara and Verkman, 2002; Kovacs  
144 et al, 2009).

145 Despite the beneficial functions of colostrum, only a few studies have attempted to  
146 determine the effect of colostrum on the skin or identify the mechanism underlying  
147 colostrum-induced keratinocyte proliferation. In the present study, we used HaCaT cells

148 to investigate the relationship between colostrum and AQP-3 expression, which plays an  
149 essential role in keratinocyte proliferation.

150

## 151 **Materials and methods**

### 152 **Bacterial culture**

153 *Lactobacillus bulgaricus* (KCTC 3635) and *Streptococcus thermophilus* (KCTC 3658) were  
154 obtained from the Korean Collection for Type Cultures (KCTC), and *L. rhamnosus* GG (ATCC  
155 53103) was purchased from the American Type Culture Collection (ATCC, Manassas, VA, USA).  
156 The three bacterial strains were cultured for 24 h at 37 °C in MRS broth (Difco, Becton,  
157 Dickinson and Company, Sparks, MD, USA), transferred to fresh MRS broth, and incubated  
158 overnight at 37 °C. The activated cells were harvested via centrifugation at 850 ×g for 10 min at  
159 4 °C, washed twice, and then resuspended in phosphate-buffered saline (PBS). A starter culture  
160 was inoculated into the defatted colostrum and milk samples.

161

### 162 **Preparation of milk and colostrum samples**

163 The milk and colostrum used in this study were obtained from the Cheongwon pasture  
164 in Chungbuk, Korea. Colostrum was collected within 24 h after calf birth, whereas milk  
165 was collected randomly. Both colostrum and milk were pasteurized at 63 °C for 30 min  
166 and centrifuged at 3,000 g at 4°C for 30 min, to remove cellular debris and fat.  
167 Fermentations were performed in sealed bottles containing 500 mL of the fat-free milk  
168 and colostrum with three kinds of bacterial strains, which included *L. bulgaricus* (KCTC  
169 3635), *S. thermophilus* (KCTC 3658), and *L. rhamnosus* GG, for 1 h at 37 °C in an  
170 incubator. After fermented samples were incubated with rennet (0.2 µL/mL) at 37 °C for  
171 30 min, they were centrifuged at 10,000 g for 1 h at 4 °C for separation into a solid (curd)  
172 and a liquid (whey) layer. Finally, the whey samples were filtered using a 0.45 µm syringe

173 filter (Sartorius Stedim Biotech, Goettingen, Germany), lyophilized, and stored at -20 °C  
174 until further use. The final lyophilized product was dissolved in water to obtain a solution  
175 with an appropriate concentration. Colostrum whey, milk whey, fermented colostrum  
176 whey, and fermented milk whey samples were used for further experiments.

177

### 178 **Cell culture and reagents**

179 The human skin keratinocyte cell line HaCaT (KCLB, Seoul, Korea) was maintained  
180 in Dulbecco's Modified Eagle's Medium (Welgene, Gyeongsan, South Korea)  
181 supplemented with 10% fetal bovine serum (Welgene) and 1% penicillin-streptomycin  
182 (Lonza, Walkersville, MD, USA) at 37 °C under 5% CO<sub>2</sub> conditions. In addition, Du145  
183 cells were maintained in RPMI 1640 medium (Welgene) supplemented with L-glutamine  
184 (300 mg/L), 25 mM HEPES, 25 mM NaHCO<sub>3</sub>, and 10% fetal bovine serum (Welgene).  
185 The HaCaT and Du145 cells were sub-cultured once every three days. Dulbecco's PBS  
186 (DPBS; Welgene) and 0.25% trypsin-EDTA (Thermo Fisher Scientific, Rockford, IL,  
187 USA) were used also used during cell culture.

188

### 189 **Cell viability assay**

190 The cell viability assay was performed using an EZ-Cytox cell viability assay kit (EZ-  
191 3000, DoGen, Seoul, Korea). HaCaT cells were seeded into 12-well plates (approximately  
192  $3 \times 10^4$  cells/well) and treated with 0, 1, 10, 100, 200, and 400 µg/mL of the colostrum  
193 whey, milk whey, fermented colostrum whey, and fermented milk whey samples for 24  
194 h. Subsequently, 100 µL of EZ-Cytox reagent was added to each well and incubation was  
195 performed for 1 h. Two-hundred microliters of the supernatant were transferred into each  
196 well of a 96-well plate, and the absorbance of the samples was measured at 450 nm using

197 a microplate reader (Tecan Sunrise, Salzburg, Austria). Cell viability values were  
198 normalized to those of the cells from the untreated control groups.

199

### 200 **Real-time PCR and quantitative RT-PCR (QPCR)**

201 Total RNA was isolated from HaCaT cells using the RNeasy Mini Kit (Qiagen, Hilden,  
202 Germany), along with on-column DNase treatment (Qiagen), as per the manufacturer's  
203 instructions. A cDNA synthesis kit (MGmed, Seoul, Korea) was used to synthesize cDNA  
204 from 1.5 µg of total RNA, using the Oligo (dT) 30 primer. Target gene expression was  
205 analyzed using the gene-specific primers listed in Table 1. The amplified products were  
206 subjected to electrophoresis on a 1.5% agarose gel and visualized with ultraviolet  
207 illumination.

208 RT-PCR was performed at 94°C for 2 min, followed by 35 cycles at 94°C for 10 s, 60°C  
209 for 30 s, and 72°C for 20 s. For QPCR, cDNA strands synthesized from the isolated total  
210 RNA were used as templates. PCR was performed on a StepOnePlus Real-Time PCR  
211 System (Qiagen), using the SYBR Green PCR Mastermix (BioRad, Hercules, CA). The  
212 cycle threshold values were normalized against the GAPDH gene expression levels. PCR  
213 was performed at 94°C for 1 min; subsequently, 40 cycles were performed at 94°C for 10  
214 s, 57°C for 10 s, and 72°C for 20 s.

215

### 216 **Western blotting analysis**

217 Treated and untreated cultured HaCaT cells were harvested with  
218 radioimmunoprecipitation assay (RIPA) buffer supplemented with a protease inhibitor.

219 The total protein content was quantified using the BCA Protein Assay (Thermo Scientific,  
220 Seoul, Korea). Equal quantities of proteins (40 µg each) were separated using 12%

221 polyacrylamide gel electrophoresis, and the resultant bands were transferred onto  
222 polyvinylidene difluoride membranes. Non-specific binding was blocked using TBST  
223 containing 1% bovine serum albumin (BSA) at 26°C for 1 h. Subsequently, the membrane  
224 was incubated overnight at 4 °C with primary antibodies diluted in TBST and a solution  
225 containing 1% BSA: AQP-3 ( 1:2,000 dilution; Merck Millipore, MA, USA), phospho-  
226 ERK (1:2,000 dilution; Cell Signaling Technology, Beverly, MA, USA), ERK (1:1,000  
227 dilution; Cell Signaling Technology), phospho-JNK (1:1000 dilution; Cell Signaling  
228 Technology), JNK (1:1,000 dilution; Cell Signaling Technology), phospho-p38 (1:1,000  
229 dilution; Cell Signaling Technology), p38 (1:1000 dilution; Cell Signaling Technology),  
230 and  $\beta$ -actin (1:1,000 dilution; Santa Cruz Biotechnology, Santa Cruz, CA). The  
231 membranes were then washed in TBST and incubated at room temperature for 1 h with  
232 horseradish peroxidase-linked anti-rabbit or anti-mouse IgGs (1:10,000 dilution; Santa  
233 Cruz Biotechnology, Santa Cruz, CA). Proteins were visualized using an enhanced  
234 chemiluminescence detection system (Thermo Scientific) and autoradiography X-ray  
235 film (AGFA, Mortsels, Belgium).  $\beta$ -Actin was used as a control to determine if each  
236 sample contained the same amount of protein.

237

### 238 **Immunocytochemistry**

239 To detect cellular AQP-3 membrane protein, HaCaT cells were seeded (approximately 1  
240  $\times 10^5$  cells/well) into a 6-well plate and treated with varying concentrations of fermented  
241 colostrum whey for 3 h (0, 1, 10, and 100  $\mu$ g/mL). HaCaT cells were rinsed with DPBS,  
242 fixed with 4% paraformaldehyde for 15 min; then, membrane permeabilization was  
243 performed for 10 min with PBS containing 0.2% Triton X-100. Non-specific protein  
244 binding was blocked using 1% BSA in PBS for 30 min at room temperature (RT), and the

245 cells were incubated overnight at 4 °C with AQP-3 primary antibodies (Millipore, diluted  
246 1:200) in 1% BSA in DPBS. The cells were then washed three times with DPBS and  
247 incubated with secondary goat anti-rabbit IgG antibodies (1:1,000 dilution; Santa Cruz  
248 Biotechnology, Texas, USA) for 1 h at RT in 1% BSA in DPBS. Samples were incubated  
249 with 3'-diaminobenzidine (DAB) solution (Burlingame, CA, USA) prepared in DPBS for  
250 5 min for DAB staining. Finally, the samples were mounted with a mounting solution  
251 (DAKO, Carpinteria, CA, USA).

252

### 253 **Statistical analysis**

254 The SPSS software (version 24.0) for Windows (SPSS, Chicago, IL, USA) was used for  
255 the statistical analysis. Differences among three or more groups or time points were  
256 determined using one-way analysis of variance, followed by Tukey's honest significant  
257 difference test. Different significance levels have been indicated using different  
258 lowercase letters.

259

## 260 **Results and Discussion**

### 261 **Effect of samples on HaCaT Cell Viability**

262 The cell viability assay was performed to investigate the cytotoxicity of colostrum  
263 whey, milk, fermented colostrum, and fermented milk in HaCaT cells. HaCaT cells were  
264 treated with increasing concentrations of colostrum whey, milk, fermented colostrum, and  
265 fermented milk (1, 10, 100, 200, and 400 µg/mL) for 24 h; the corresponding viabilities  
266 of HaCaT cells treated with these varied concentrations of colostrum whey were 102.19  
267 ± 1.06%, 101.13 ± 1.08%, 104.16 ± 0.87%, 104.41 ± 1.68%, and 107.48 ± 1.09%,  
268 respectively. Values were expressed as a percent of the control group values (regarded as

269 100%). The viabilities of milk whey (1, 10, 100, 200, and 400  $\mu\text{g/mL}$ )-treated cells were  
270  $96.28 \pm 1.07\%$ ,  $98.27 \pm 1.25\%$ ,  $98.98 \pm 2.23\%$ ,  $103.70 \pm 1.08\%$ , and  $107.51 \pm 0.59\%$ ,  
271 respectively; the viabilities of the fermented colostrum whey (1, 10, 100, 200, and 400  
272  $\mu\text{g/mL}$ )-treated cells were  $101.72 \pm 1.27\%$ ,  $104.31 \pm 0.97\%$ ,  $108.97 \pm 1.34\%$ ,  $123.17 \pm$   
273  $1.08\%$ , and  $124.49 \pm 1.94\%$ , respectively; and those of the fermented milk whey (1, 10,  
274 100, 200, and 400  $\mu\text{g/mL}$ )-treated cells were  $104.95 \pm 1.39\%$ ,  $105.52 \pm 2.71\%$ ,  $106.55 \pm$   
275  $1.21\%$ ,  $107.23 \pm 0.57\%$ ,  $112.92 \pm 0.37\%$ , respectively (Fig. 1A). No notable cytotoxicity  
276 was observed in all treated HaCaT cell samples at concentrations of up to 400  $\mu\text{g/mL}$ .  
277 The cell viability increased significantly at a concentration of 400  $\mu\text{g/mL}$  in case of all  
278 samples; especially, the viability increased significantly from 100  $\mu\text{g/mL}$  to 400  $\mu\text{g/mL}$   
279 in case of the cells treated with fermented colostrum whey (Fig. 1A). These results  
280 indicated that fermented colostrum whey significantly increased cellular proliferation.  
281 Yoon et al. (2002) reported that treatment of keratinocytes with bovine colostrum  
282 promoted keratinocyte proliferation. In addition, bovine colostrum promoted  
283 proliferation of HaCaT cell, although no significant changes were seen in HaCaT cell  
284 growth after 48 h of colostrum treatment, which suggests that the enriched bioactive  
285 substances in the colostrum plays an essential role in HaCaT cell proliferation (Kovacs et  
286 al. 2009). These previous studies and the current data together reveal the ability of  
287 colostrum to promote HaCaT cell proliferation.

288

### 289 **Expression of AQP-3 in human keratinocyte HaCaT cells**

290 To identify the subtypes of AQPs expressed in HaCaT, we performed RT-PCR using  
291 AQP-specific primers (Table 1). Among the tested AQPs, the expression levels of AQP-  
292 3 and AQP-10 mRNAs were identified; AQP-3 expression was abundant in HaCaT cells

293 and was observed to occur in DU145 cells, which were used as positive controls (Fig.  
294 1B). DU145 cells are prostate cancer cell line, and it is well known that AQP3 is expressed  
295 in these cells (Almeida et al., 2021). In the correlation of keratinocyte proliferation with  
296 *AQP-3* expression, two complex mechanisms were found to be defined and undefined,  
297 respectively. AQP-3-mediated cell proliferation is one of the well-defined mechanisms.  
298 Human keratinocytes with AQP-3 knockdown and keratinocytes from AQP-3-knockout  
299 mice showed reduced levels of proliferation (Hara-Chikuma and Verkman, 2008). The  
300 upregulation of AQP-3 by the transfection of AQP-3 plasmid DNA has been shown to  
301 promote the proliferation of human keratinocytes and increased cellular glycerol and ATP  
302 levels (Nakahigashi et al., 2011).

303

#### 304 **Fermented colostrum upregulates *AQP-3* expression in HaCaT cells**

305 Next, we investigated whether the whey obtained from colostrum, milk, fermented  
306 colostrum, and fermented milk altered the expression of *AQP-3*. HaCaT cells were treated  
307 with 1, 10, or 100  $\mu\text{g}/\text{mL}$  of all four whey samples, and the AQP-3 mRNA level was  
308 measured using RT-PCR and QPCR (Fig. 2). AQP-3 mRNA expression was not altered  
309 in cells treated with colostrum, milk, and fermented milk (Fig. 2A, B and D, respectively),  
310 but was markedly increased in cells treated with fermented colostrum whey at a  
311 concentration of 100  $\mu\text{g}/\text{mL}$  (Fig. 2C). Consistently, the QPCR analysis data showed that  
312 treatment with colostrum and milk did not alter the AQP-3 expression at any  
313 concentration, but treatment with fermented colostrum significantly increased the AQP-  
314 3 expression at 10 and 100  $\mu\text{g}/\text{mL}$  (Fig. 2E); the increase in the AQP-3 expression level  
315 was  $3.06 \pm 0.05$ -fold higher than that observed for cells treated with milk or colostrum  
316 (at a concentration of 100  $\mu\text{g}/\text{mL}$ , Fig. 2E). In addition, treatment with fermented milk

317 resulted in a significant  $1.23 \pm 0.05$ -fold increase in the AQP-3 expression level at a  
318 concentration of 100  $\mu\text{g}/\text{mL}$  compared with the level observed with milk and colostrum  
319 (Fig. 2E). The duration of AQP-3 expression after treatment with fermented colostrum  
320 was determined by treating HaCaT cells with 100  $\mu\text{g}/\text{mL}$  of fermented colostrum. The  
321 expression of AQP-3 mRNA increased after treating cells for 2 h, peaked at 3 h, and  
322 decreased after 6 h (Fig. 2F).

323 To identify AQP-3 expression at the protein level (Fig. 3), HaCaT cells were treated  
324 with 0, 1, 10, and 100  $\mu\text{g}/\text{mL}$  of fermented colostrum for 3 h, and subsequently analyzed  
325 via western blotting and immunocytochemistry. AQP-3 protein expression levels  
326 increased following fermented colostrum treatment at concentrations of 10 and 100  
327  $\mu\text{g}/\text{mL}$  (Fig. 3A). Densitometric analyses of AQP-3 and beta-actin immunoblots showed  
328 that AQP-3 protein expression levels were 10.5 and 10.9-fold higher than that of the  
329 negative control (Fig. 3B). Consistently, the immunocytochemistry data also showed that  
330 AQP-3 was expressed in the cell membrane, and strong AQP-3 staining signal was  
331 visually observed in fermented colostrum-treated cells compared with that in the control,  
332 in a dose-dependent manner (Fig. 3C). These data demonstrate that the upregulation of  
333 AQP-3 expression in keratinocyte occurred because of fermented colostrum treatment.

334 Currently, many studies are investigating the active compounds that increase AQP-3  
335 expression levels, to enable the treatment of skin disorders. Many studies have reported  
336 plant-derived substances such as green *Coffea arabica* seed oil extract (Del Carmen  
337 Velazquez Pereda et al., 2011), trans-zeatin purified from *Zea mays* (Yang et al., 2009),  
338 *Piptadenia colubrina* extract (Del Carmen Velazquez Pereda et al., 2011), and asiaticoside,  
339 isolated from *Centella asiatica* (Wijayadi and Darmawan, 2017).

340 Although there is no direct evidence in this study, it is possible that fermented  
341 colostrum regulates HaCaT keratinocyte proliferation via the upregulation of AQP-3.  
342 EGF, which is abundantly present in colostrum, is a known upstream regulator of AQP-3  
343 expression in cultured colorectal cancer (Li et al., 2013), ovarian cancer (Ji et al., 2008),  
344 and pancreatic cancer (Liu et al., 2012) cells. As expected, the upregulation of AQP-3 in  
345 cancer cells resulted in their enhanced proliferation. Milk and fermented milk whey  
346 contain various bioactive substances that regulate skin cells. For instance, lactoferrin  
347 stimulates proliferation and wound healing in epidermal keratinocytes (Tang et al., 2010),  
348 and exopolysaccharides from *Streptococcus thermophilus* prevent the occurrence of  
349 ultraviolet-induced skin damage (Morifuji et al., 2017). Additionally, bioactive peptides  
350 with low molecular mass are also produced during milk fermentation and the rennet  
351 enzyme reaction (Shinagawa et al., 2018). Shinagawa et al. reported the increase in  
352 keratinocyte proliferation with increase in AQP-3 expression after treatment with  
353 compounds with molecular weight < 3 kDa that were produced during the cheese  
354 fermentation process (Shinagawa et al., 2018). Interestingly, lactose, calcium, and other  
355 abundant molecules with low molecular mass did not affect AQP-3 expression and  
356 keratinocyte proliferation (Shinagawa et al., 2018), suggesting that low-molecular-weight  
357 peptides produced during fermentation activated the keratinocytes. A previous study  
358 found that milk whey was abundantly present in phospholipids, which supports this  
359 hypothesis (Vaghela and Kilara, 1995). The topical treatment of murine dorsal skin with  
360 phospholipids promotes keratinocyte proliferation and differentiation (Kumura et al.,  
361 2012). Fermented colostrum whey might contain abundant amounts of both bioactive  
362 peptides and phospholipids that were generated by fermentation. Because we found that  
363 the use of fermented colostrum whey resulted in 2-fold higher *AQP-3* expression levels

364 than those following treatment with fermented milk whey, it is possible that fermented  
365 colostrum whey has a greater abundance of bioactive peptides associated with *AQP-3*  
366 expression than fermented milk whey. However, the specific whey components that  
367 regulate *AQP-3* expression in HaCaT keratinocytes remain unclear. Further experiments  
368 are required to understand the regulatory effects of fermented colostrum whey  
369 components on HaCaT keratinocytes.

370 Interestingly, *AQP-3* expression levels were maximal at 3 h post treatment with  
371 fermented colostrum whey and decreased to levels equivalent to those of the control  
372 within 6 h. Other studies also showed that the change in *AQP-3* expression caused by  
373 external stimuli occurs rapidly. For examples, when cells were treated with 10–20 mg/mL  
374 of *Piptadenia colubrina* extract, *AQP-3* expression increased after 2 h, peaked at 6 h, and  
375 decreased after 24 h (Del Carmen Velazquez Pereda et al., 2010). Treatment with a green  
376 *Coffea arabica* seed extract resulted in increased *AQP-3* expression after 3 to 6 h (Del  
377 Carmen Velazquez Pereda et al., 2009). These results support our observation that the  
378 *AQP-3* expression level began to increase after 2 h and peaked at 3 h.

379

### 380 **Fermented colostrum whey regulates the MAPK signaling pathway in HaCaT cells**

381 To understand the regulatory mechanism by which fermented colostrum whey  
382 effectively induced HaCaT cell proliferation, the present study investigated whether  
383 MAPK pathways were involved in the proliferation of HaCaT cells, using fermented  
384 colostrum whey. The phosphorylation levels of three mitogen-activated protein kinases  
385 (MAPKs), including ERK 1/2, JNK1/2, and p38, were assessed. HaCaT keratinocytes  
386 were treated with 100 µg/mL of fermented colostrum whey for varying durations. The  
387 results showed that treatment with 100 µg/mL of fermented colostrum whey rapidly

388 induced the phosphorylation of JNK1/2 and p38 MAPK within 20 min (Fig 4A, B),  
389 whereas the levels of phosphorylated ERK 1/2 were not increased (Fig. 4C). In summary,  
390 exposure to fermented colostrum whey increased the expression level of AQP-3 mRNA  
391 in, and induced the proliferation of, HaCaT keratinocytes via the activation of p38 and  
392 pJNK signaling (Fig. 4D). In the present study, the phosphorylation of p38MAPK and  
393 JNK were observed in HaCaT keratinocytes after their treatment with fermented  
394 colostrum whey. Although studies identifying the underlying mechanism of keratinocyte  
395 proliferation and effects of colostrum or milk have not been undertaken thus far, similar  
396 results were seen with human osteoblasts. The treatment of human osteoblasts with  
397 concentrated low-molecular-weight bovine milk whey proteins (1–30 kDa) increased  
398 alkaline phosphatase activity and mineralization, by upregulating JNK-activating  
399 transcription factor 4 (ATF4) via the phosphorylation of JNK (Tsuji-Naito and Jack, 2012).  
400 Interestingly, they used low-molecular-weight (1–30 kDa) fractions of whey proteins, and  
401 this fraction typically exhibited an abundance of growth factors and cytokines (Hwang et  
402 al., 2012).

403 These prior reports and our data provide strong evidence that certain factors in the  
404 colostrum are important for keratinocyte activation. MAPK families play a crucial role in  
405 the regulation of different processes, including cell proliferation, development, differentiation,  
406 transformation, and apoptosis. Five distinct groups of MAPKs have been characterized in  
407 mammals, including ERK1/2, JNKs, p38, and ERK3, 4, and 5. In addition, the most  
408 extensively studied groups of vertebrate MAPKs include ERK1/2, JNKs, and p38 kinases  
409 (Kyriakis and Avruch, 2001). Evidence has shown that the JNK signal pathway was involved  
410 in keratinocyte proliferation. Gazel et al. has reported about JNK-induced epidermal  
411 keratinocyte proliferation and the inhibition of differentiation (Gazel et al., 2006). In addition,

412 p38 MAPKs were defined as cellular protein kinases that were activated in response to stresses,  
413 such as osmotic stress, inflammatory cytokines, ultraviolet light, and changes in the oxygen  
414 content (Brewster et al, 1993; Han et al., 1994). In contrast, a recent study has found that  
415 p38-MAPKs were also activated in response to a variety of extracellular growth and  
416 differentiation cues (Nebreda and Porras, 2000). Although ERK1/2 was not activated,  
417 fermented colostrum whey induced HaCaT cell proliferation and also the activation of  
418 the p38 MAPK and JNK signaling pathways. In contrast with our results, Li et al (2015).  
419 reported the downregulation of AQP-3 expression by lipopolysaccharides via the p38/c-  
420 Jun signaling pathway in HT-29 human colon epithelial cells. Taken together, our results  
421 suggest that fermented colostrum whey has potential for effectively improving the  
422 efficacy of treatment procedures for various skin disorders, as it activates AQP-3  
423 expression and proliferation with activation of p38/c-Jun N-terminal kinase in  
424 keratinocytes.

425

#### 426 **Competing interests**

427 The authors state that they have no competing interests to declare.

428

#### 429 **Acknowledgements**

430 This study was funded by a National Research Foundation of Korea (NRF) grant provided by the

431 Korean government (MSIT) (grant number NRF2019R1A2C 1008310).

432 This paper was supported by Konkuk University Researcher Fund in 2020.

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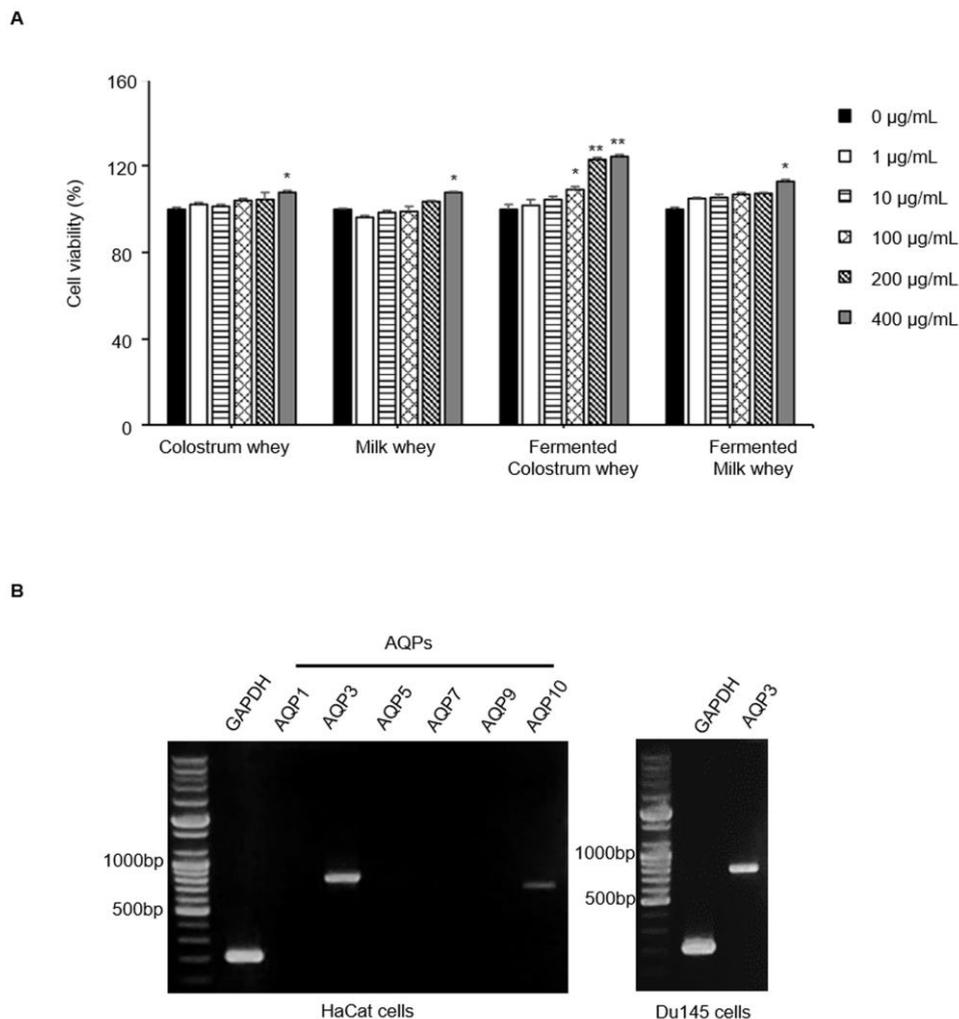
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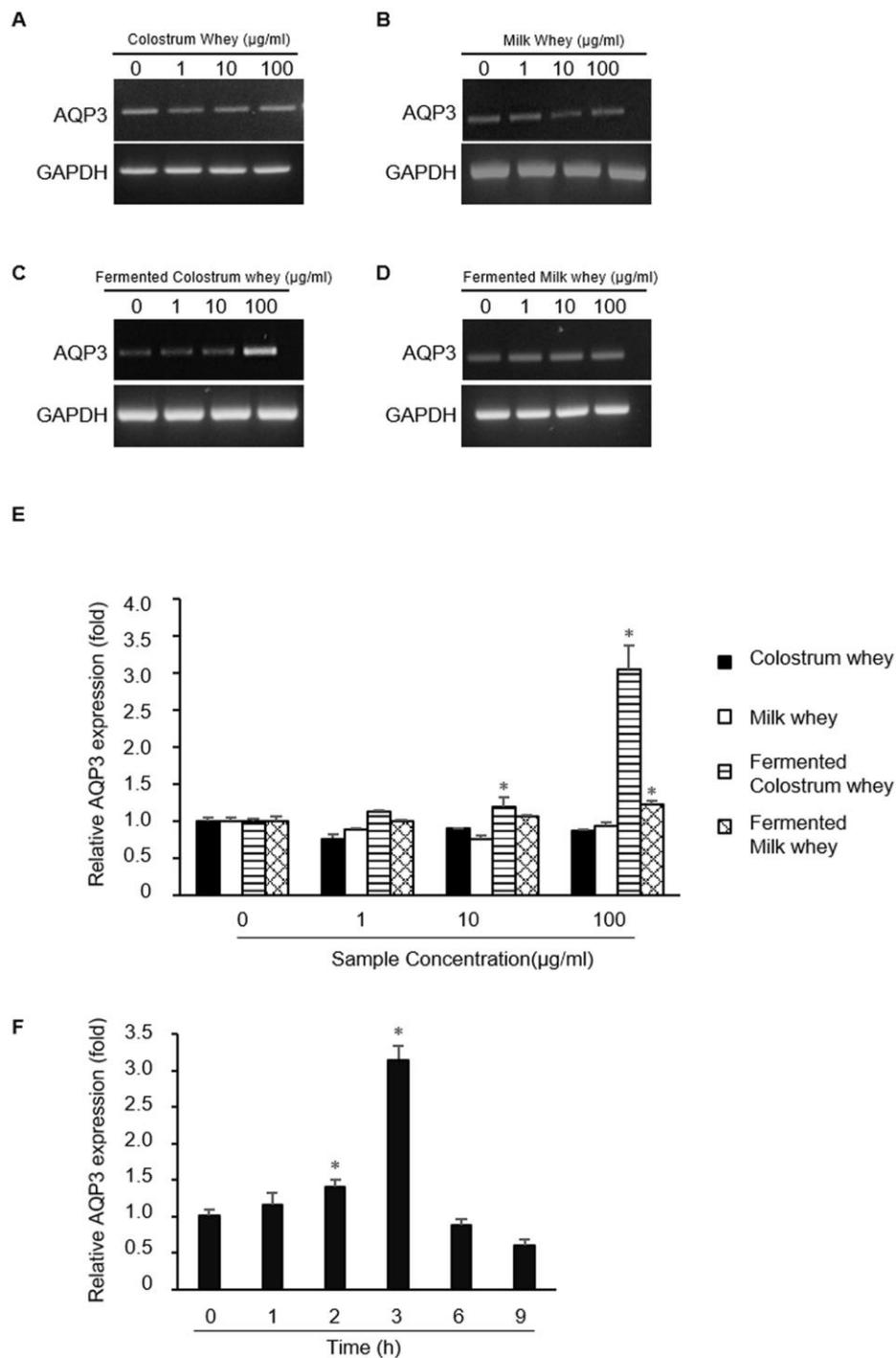
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625

626 **Figure legends**

627

628 **Figure 1.** (A) Effects of 0-400 µg/mL colostrum whey, milk whey, fermented colostrum  
 629 whey, and fermented milk whey on the viability of HaCaT cells. The cell viability was  
 630 measured by the MTT assay. Results expressed as a % of control absorbance. Data were  
 631 presented as the means  $\pm$  SE values of three independent experiments (n=3). (B) PCR  
 632 analysis of AQP expression in cultured HaCaT cells. RNA (1.5 µg) was reverse-  
 633 transcribed into first-strand cDNA and amplified by PCR, using AQP (1, 3, 5, 7, 9, 10)-  
 634 specific primers. AQP-3 is the most abundant AQP in HaCaT cells. *AQP-3* expression  
 635 was detected in DU145 cells, which served as the positive controls.

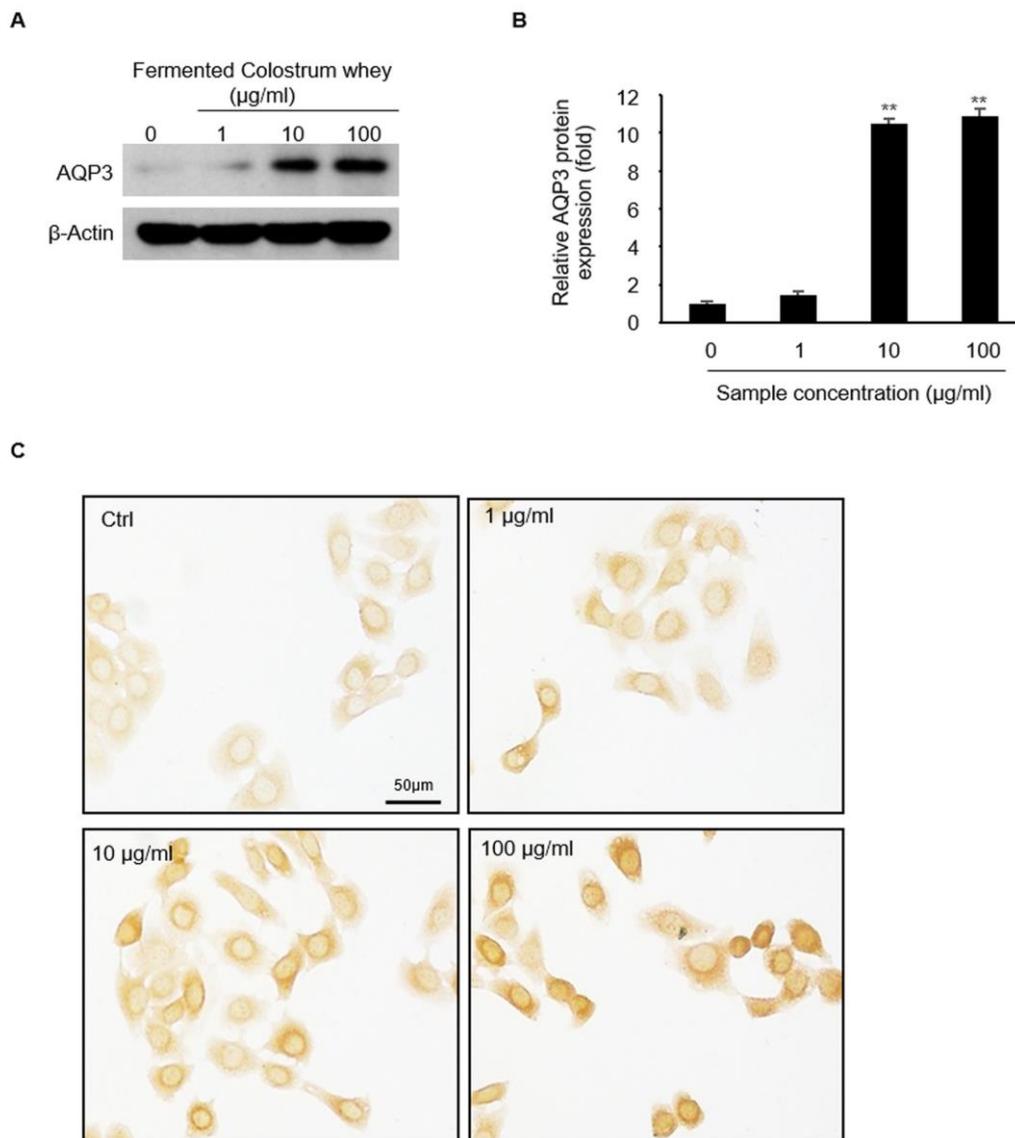


636

637 **Figure 2.** Effect of fermented colostrum whey on AQP-3 expression. HaCaT cells were  
 638 treated with different concentrations (1, 10, and 100  $\mu\text{g/mL}$ ) of (A) colostrum whey, (B)  
 639 milk whey, (C) fermented colostrum whey, and (D) fermented milk whey, for 3 h. AQP-  
 640 3 mRNA expression was analyzed by RT-PCR. GAPDH was used as a control. (E)  
 641 AQP-3 mRNA levels were quantified by real-time PCR (QPCR). GAPDH mRNA level

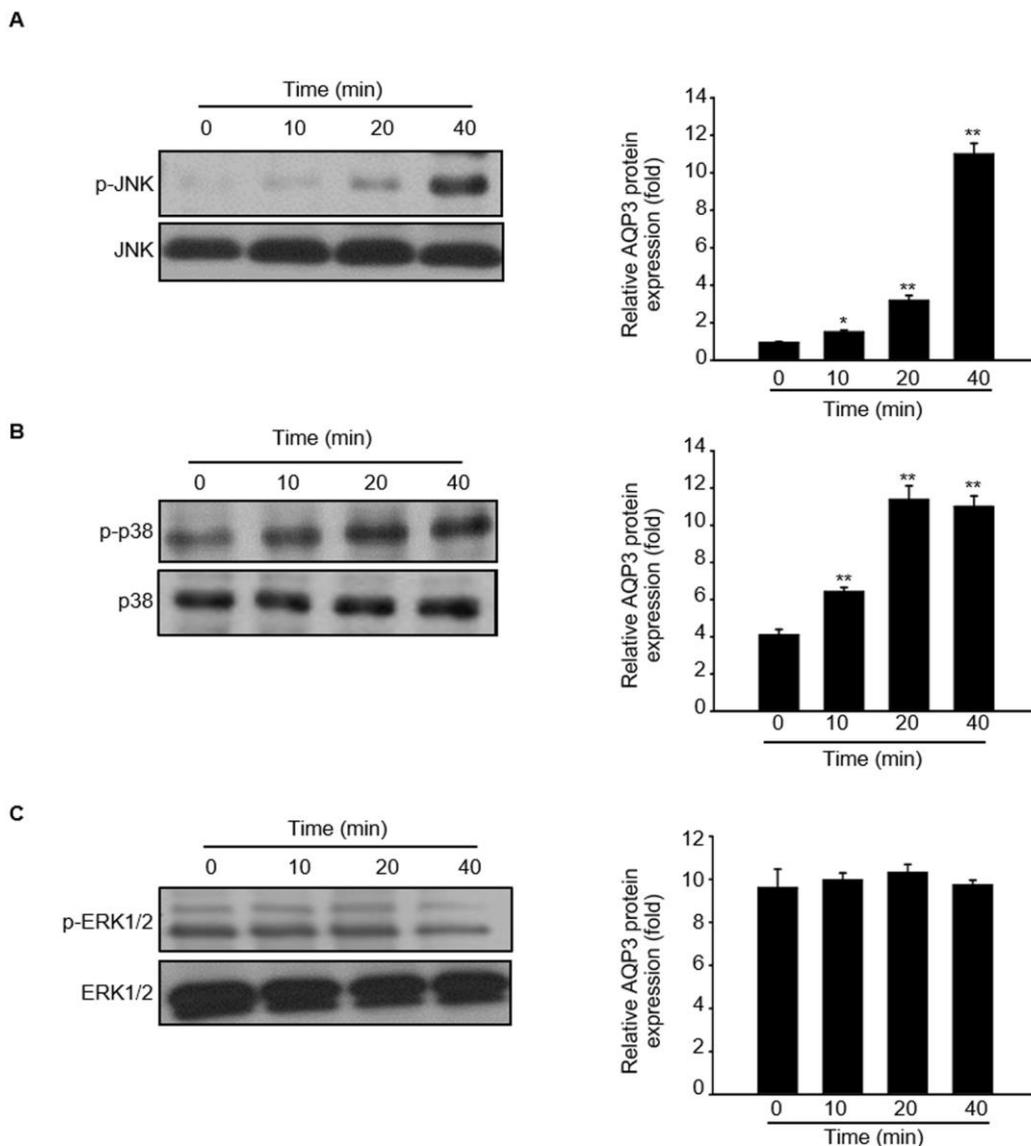
642 was used for the normalization of the mRNA levels of AQP-3. (F) HaCaT cells were  
643 treated with a concentration of 100  $\mu\text{g/mL}$  of fermented colostrum whey, and the AQP-3  
644 mRNA levels at different durations (1, 2, 3, 6, and 9 h) were detected by QPCR.  
645 GAPDH mRNA levels were used for normalization. Data were presented as the means  $\pm$   
646 SE values of three independent experiments (n=3).  
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649 **Figure 3.** Effect of fermented colostrum whey on AQP-3 protein expression. (A)  
 650 HaCaT cells were treated with different amounts (0, 1, 10, and 100 µg) of fermented  
 651 colostrum whey for 3 h. AQP-3 protein expression was analyzed by immunoblotting.  
 652 The β-actin antibody was used as a control to confirm that equal amounts of each  
 653 protein were loaded. (B) In the bar graph, the relative intensity of AQP-3 is expressed as  
 654 a ratio of AQP-3/β-actin and compared with that for the untreated group. Data were  
 655 presented as the means ± SE values of three independent experiments (n=3). (C) AQP-3  
 656 protein expression was analyzed by immunocytochemistry at a 400× magnification  
 657 (Scale bars = 50 µm; all images were taken at the same magnification).  
 658 Immunocytochemical staining was performed using the DAB solution.



659

660 **Figure 4.** Role of MAPK signaling in fermented colostrum whey-induced proliferation  
 661 of HaCaT keratinocytes. HaCaT cells were treated with 100  $\mu\text{g}/\text{mL}$  of fermented  
 662 colostrum whey for varying time periods. Phosphorylation levels of MAPKs such as (A)  
 663 p-JNK, (B) p-p38, and (C) pERK1/2 were analyzed by western blotting. The  $\beta$ -actin  
 664 antibody was used as an internal control to confirm that equal amounts of each protein  
 665 were loaded. The bar graph represents the normalized values of the densities of each  
 666 band relative to the densities of the bands of the non-active form of each protein. The  
 667 values represent the mean  $\pm$  SE values of three independent experiments. (n=3).

668

669 **Table1.** Primer sequences for real-time PCR

670

Gene	Sequence	Length (b.p.)
AQP-1	F: 5'-CTTTGTCTTCATCAGCATCGGTTC-3' R: 5'- ATGTCGTCGGCATCCAGGTCATAC-3'	711
AQP-3	F: 5'-ACCCTCATCCTGGTGATGTTTG-3' R: 5'-TCTGCTCCTTGTGCTTCACAT-3'	781
AQP-5	F: 5'- CTCTTGGTGGGCAACCAGATC-3' R: 5'-TCACTCAGGCTCAGGGAGTTGG-3'	839
AQP-7	F: 5'- GGGAGCTACCTTGGTGTCAACTT-3' R: 5'- CATCTTGGGCAATACGGTTATCC-3'	720
AQP-9	F: 5'- ACGTTTTGGAGGGGTCATCAC-3' R: 5'- CAGGCTCTGGATGGTGGATTTC-3'	664
AQP-10	F: 5'- ATAGCCATCTACGTGGGTGGTAAC-3' R: 5'- TTTGTGTTGAGCAGACACCAGATC-3'	651
GAPDH	F: 5'- GGATTTGGTCGTATTGGG-3' R: 5'- GGAAGATGGTGATGGGATT-3'	205

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