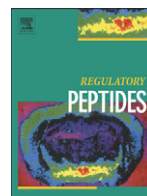




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# Corticotropin-releasing hormone attenuates vascular endothelial growth factor release from human HaCaT keratinocytes

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## ABSTRACT

**Objectives:** Corticotropin-releasing hormone (CRH) is a central component of the local hypothalamic-pituitary-adrenal (HPA) axis, which has a functional equivalent in the skin. To determine whether CRH and its receptor, CRH-R1, modulate the expression of vascular endothelial growth factor (VEGF), which is overexpressed in psoriatic epidermis and plays a causal role in the pathogenesis of psoriasis, we investigated the effect of CRH on the expression of VEGF in a human keratinocyte cell line (HaCaT) and whether this effect is via the mitogen-activated protein kinase (MAPK) signal transduction pathways.

**Methods:** Real-time RT-PCR, ELISA assay and western blot were used in the present study to investigate the expression of VEGF in CRH-treated HaCaT cells.

**Results:** The mRNA and protein levels of VEGF in CRH-treated HaCaT cells were significantly attenuated. However, this downregulation was abrogated by pretreatment with antalarmin, SB203580 and SP600125; pretreatment with PD98059 did not attenuate the effects of CRH on the expression of VEGF. In addition, CRH treatment induced rapid phosphorylation of p38 MAPK and JNK1/2, and antalarmin, SB203580 and SP600125 inhibited CRH-induced phosphorylation of p38 MAPK and JNK1/2.

**Conclusions:** CRH might downregulate the expression of VEGF through the CRH-R1 and MAPK (p38 MAPK and JNK1/2) signaling pathways in human HaCaT cells.

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## 1. Introduction

Corticotropin-releasing hormone (CRH), a 41-amino acid neuropeptide, is produced mainly in the hypothalamus and regulates endocrine and behavioral responses to stress through the activation of the hypothalamic-pituitary-adrenal (HPA) axis [1]. CRH exerts its actions via interaction with specific CRH receptors (CRH-Rs) [2]. Three subtypes of CRH-Rs – CRH-R1, CRH-R2 and CRH-R3 – belong to the G protein-coupled seven-transmembrane receptors [2–5]. Recent research has indicated that CRH and CRH-Rs are expressed and have functions in the skin [6–9]. In human skin, CRH-R1 is the major receptor in epidermis and dermis [7].

The human skin is an independent peripheral endocrine organ [10]. It is a prominent target organ for numerous neurotransmitters and neuropeptide signals that have a profound impact on skin biology in health and disease [11]. Skin has its own functional peripheral equivalent of the HPA axis; CRH produced peripherally and other HPA axis components comprise the cutaneous HPA systems [2,6,12]. Cutaneous CRH is synthesized by cutaneous cells and immune cells in human skin [1,13,14] and regulates various functions of the skin, especially maintaining local homeostasis [2,15]. Compelling evidence has sug-

gested that CRH inhibits the proliferation of both human primary keratinocytes and immortalized keratinocytes [15,16] and stimulates differentiation of keratinocytes [17,18]. Slominski et al. demonstrated a CRH-based homeostatic response system in the skin, and CRH acts as a pleiotropic cytokine [19]. The abnormal differentiation of keratinocytes resulting in a suboptimal barrier function of the skin may be evidence of a protective function for CRH and CRH-R1 in the skin [20].

Psoriasis is a chronic inflammatory disease characterized by erythematous plaques with silvery scales. Psoriatic lesions exhibit proliferation of epidermal keratinocytes, inflammatory cell infiltration, and increased angiogenesis of the superficial dermal vessels [21]. The prominence of dermal microvascular expansion in the psoriatic lesion demonstrates that psoriasis is an angiogenesis-dependent disease [22]. Vascular endothelial growth factor (VEGF) is a crucial regulator of angiogenesis and vascular permeability in both physiological and pathological conditions such as tumor growth and chronic inflammation [23–25]. It was originally identified as an endothelial cell-specific growth factor that can stimulate endothelial cells to undergo angiogenesis and induce vascular permeability, thus facilitating the development of some diseases [26]. VEGF is expressed and secreted by epidermal keratinocytes in normal human skin [27]. Keratinocytes overexpress VEGF in clinically involved and uninvolved skin of patients with chronic plaque psoriasis [28]. In transgenic mice with epidermis-specific overexpression of VEGF, enhanced skin vascularity and vascular

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permeability [29], chronic transgenic delivery of VEGF to the skin induced inflammation and all characteristics of psoriasis spontaneously, and the VEGF antagonist reversed the phenotype. These findings suggested a causative role of VEGF in the pathogenesis of psoriasis [30].

However, little is known about the exact role of CRH in skin. We hypothesized that CRH may modulate VEGF expression and investigated the effect of CRH and its receptor CRH-R1 on the expression of VEGF in a human keratinocyte cell line, HaCaT. We examined whether this effect functioned via the mitogen-activated protein kinase (MAPK) signal transduction pathway, particularly p38 mitogen-activated protein kinase (p38 MAPK), extracellular signal-regulated protein kinase 1/2 (ERK1/2) and c-Jun N-terminal kinase (JNK) [31,32].

## 2. Materials and methods

### 2.1. Antibodies and reagents

Human CRH and the CRH-R1 antagonist antalarmin were from Sigma (St. Louis, MO, USA). SB203580 (the inhibitor of p38 MAPK), PD98059 (the inhibitor of ERK1/2) and SP600125 (the inhibitor of JNK1/2) were from Biosource (Camarillo, CA). Antibody against  $\beta$ -actin was from Santa Cruz Biotechnology (Santa Cruz, CA). Antibodies against phospho-p38 MAPK, p38 MAPK, phospho-JNK1/2 and JNK1/2 were from Cell Signaling Technology (Beverly, MA, USA); horseradish peroxidase-conjugated anti- $\beta$ -actin and anti-rabbit IgG antibodies were from Santa Cruz Biotechnology. The human VEGF ELISA kit was from R&D Systems (Minneapolis, MN, USA).

### 2.2. Cell culture

Immortalized human HaCaT keratinocytes were maintained at 37 °C and 5% carbon dioxide (CO<sub>2</sub>) in Dulbecco's modified Eagle's medium supplemented with 10% heat-inactivated fetal bovine serum, 100 U/ml penicillin and 100  $\mu$ g/ml streptomycin. HaCaT cells were digested for 5 min at 37 °C with phosphate-buffered saline (PBS) containing 0.25% trypsin and 0.02% EDTA. Then, the cells were collected by centrifugation at 800 rpm for 10 min at 4 °C, and cultured at a density of  $1 \times 10^6$  cells/plate at 37 °C in a humid atmosphere of 5% CO<sub>2</sub> and 95% air. The culture medium was changed twice a week.

### 2.3. Cell pretreatment

HaCaT cells were seeded at a density of  $1 \times 10^6$  cells/plate, grown for 48 h until 70% confluence, and then cells were washed with serum-free media and maintained with media without FBS at least 12 h prior to the experiments. HaCaT cells were pretreated with 10  $\mu$ M antalarmin, SB203580, PD98059 or SP600125 respectively for 1 h, and then different concentrations of CRH (1, 10 and 100 nM) were added to the cells.

### 2.4. Real-time RT-PCR

After treatment, total RNA was extracted from HaCaT cells by use of TRIzol (Invitrogen, Carlsbad, CA) according to the manufacturer's instructions. Reverse transcription was performed at 37 °C for 1 h by use of an MLV Kit (Promega, Madison, WI), and 1  $\mu$ g total RNA was used. Real-time RT-PCR involved Light Cycler 2.0 (Roche Applied Science, USA). cDNA was amplified with the use of Light-Cycler-FastStart DNA Master SYBR Green I (Roche, Indianapolis, IN). The primers for VEGF (GenBank NM\_001033756.1) were 5'-AAGTGGTCCCAGGCTGCA-3' (forward), and 5'-ACTCCAGGCCCTCGTCA-3' (reverse). The primers for human  $\beta$ -actin (GenBank NM\_001101.2) were 5'-TGGACATCCGAAAGAC-3' (forward) and 5'-GAAAGGGTGAACGCAACTA-3' (reverse). The cycling conditions were denaturation for 10 s at 95 °C, amplification for 35 cycles, with denaturation for 5 s at 95 °C, annealing for 5 s at 55 °C for  $\beta$ -actin and 60 °C for VEGF, and extension for 15 s at 72 °C. At the end of each cycle, the fluorescence emitted by the SYBR Green I dye was measured.

After amplification, a melting curve was generated by holding the reaction at 65 °C for 30 s and then heating slowly to 95 °C with a ramp rate of 0.1 °C/s. The data were analyzed by use of Light Cycler v4.0 (Roche Applied Science). VEGF mRNA expression was normalized to that of the housekeeping gene human  $\beta$ -actin. Relative VEGF mRNA levels were calculated using the 2<sup>(-Delta Delta C(T))</sup> Method [33]. Three independent experiments were performed in triplicate.

### 2.5. VEGF ELISA

After stimulation for 24 h, culture supernatants of cells were collected, centrifuged (15,000 rpm, 5 min) and stored at -80 °C until analysis. The concentration of VEGF in the culture supernatant was measured by an immunoassay kit according to the manufacturer's instructions. The human VEGF ELISA kit was from R&D Systems (Minneapolis, MN, USA). Three independent experiments were performed in triplicate.

### 2.6. Western blot analysis

Stimulated HaCaT cells were lysed in ice-cold RIPA buffer containing 1% phenyl methyl sulfonyl fluoride and centrifuged at 15,000 rpm for 10 min at 4 °C. Supernatants were collected, and the protein concentrations were measured with the use of a BCA protein assay kit (Beyotime, Jiangsu, China). Equal amounts of protein were boiled for 5 min, and 30  $\mu$ g of total protein was separated on 12% SDS-PAGE and transferred to a 0.22- $\mu$ m nitrocellulose membrane (Bio-Rad, Hercules, CA). After being blocked with 5% non-fat milk, the blots were washed with PBS containing 0.1% Tween 20 and incubated with an appropriate primary antibody at 4 °C overnight. Antibodies against phospho-p38 MAPK, p38 MAPK, phospho-JNK1/2 and JNK1/2 were used at 1:500 dilution; antibody against  $\beta$ -actin was used at 1:5000 dilution. After several washes, the membranes were incubated with horseradish peroxidase-conjugated secondary antibody (1:5000) for 1 h at 37 °C and then washed again. The blots were visualized with use of an enhanced chemiluminescence kit (Millipore, Billerica, MA, USA). The images were recorded by use of FluorChem9900 (Alpha Innotech, CA, USA) and analyzed with use of Quantity One software (Bio-Rad Laboratories, Hercules, CA). Three independent experiments were performed.

### 2.7. Statistical analysis

For each condition, data from at least three independent experiments were quantified and analyzed by one-way ANOVA with post-hoc LSD *t* test. A *P* < 0.05 was considered statistically significant. Analysis involved use of SPSS v16.0 (SPSS Inc., Chicago, IL, USA).

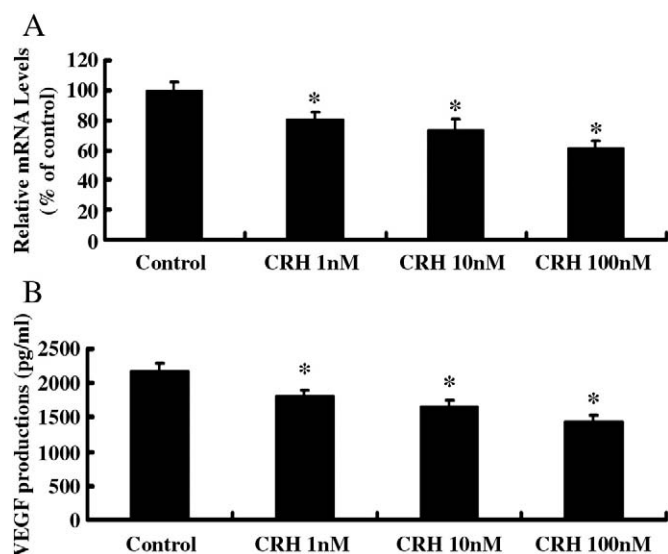
## 3. Results

### 3.1. Effect of CRH on VEGF production in HaCaT cells

To study the effect of CRH on VEGF mRNA expression in HaCaT cells, the cells were treated with different concentrations of CRH (1, 10 and 100 nM). Total RNA was extracted after 4 h and real-time RT-PCR for VEGF was performed. VEGF production was measured by ELISA kit from supernatants of CRH-treated cultured HaCaT cells collected 24 h after the exposure. Real-time RT-PCR and ELISA revealed VEGF mRNA expression and production, respectively, significantly decreased by CRH (1, 10 and 100 nM) in HaCaT cells in a dose-dependent manner, with the maximal effect at 100 nM (Fig. 1A, B).

### 3.2. CRH-R1 and MAPK signaling pathways are involved in VEGF production

Since CRH treatment led to decreased VEGF expression, we examined whether CRH-R1 and MAPK signaling pathways participated in the regulation of VEGF expression. HaCaT cells were pretreated with



**Fig. 1.** Effect of CRH on the mRNA expression and production of vascular endothelial growth factor (VEGF) in HaCaT cells. (A) Real-time RT-PCR analysis of VEGF mRNA level in HaCaT cells treated with different concentrations of CRH (1, 10 and 100 nM) for 4 h. (B) ELISA of VEGF production in cells treated with the indicated concentrations of CRH for 24 h. Data are mean  $\pm$  SD of experiments performed in triplicate. \*  $P < 0.05$  compared with control cells.

regulated VEGF expression in HaCaT cells by CRH-R1 through p38 MAPK and JNK1/2 but not ERK1/2 pathways. 206 207

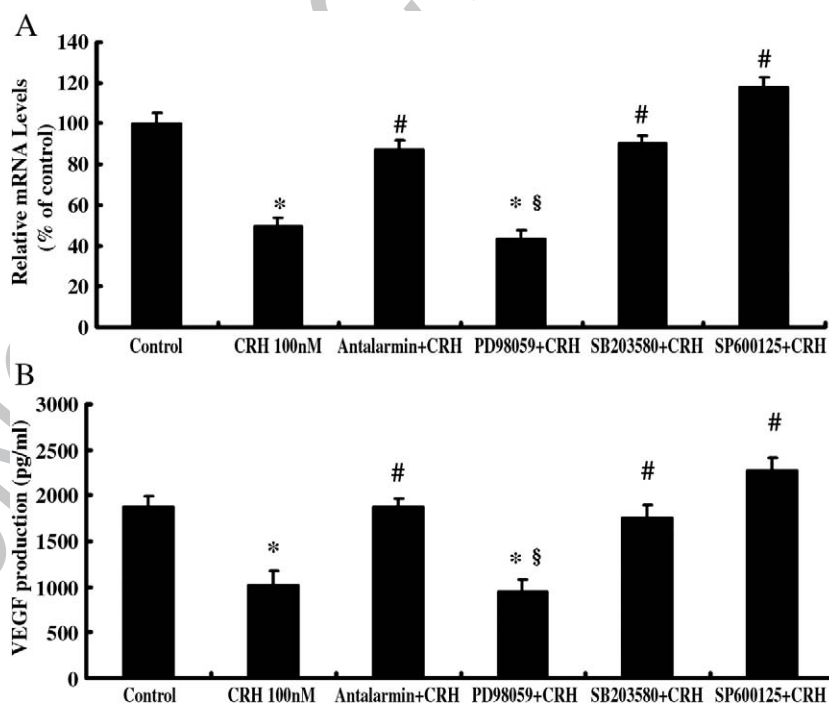
### 3.3. CRH activates p38 MAPK and JNK1/2 phosphorylation in HaCaT cells 208

To give evidence of the activation of p38 MAPK and JNK1/2 pathways, we used western blot analysis to evaluate the phosphorylation of p38 MAPK and JNK1/2 in CRH-treated HaCaT cells. CRH induced a rapid phosphorylation of p38 MAPK and JNK1/2, with a peak at 5 min (Fig. 3A). Normalization to total p38 MAPK and JNK1/2 confirmed that CRH specifically altered p38 MAPK and JNK1/2 phosphorylation but not the expression levels. Pretreating HaCaT cells respectively with SB203580, SP600125 or antalarmin for 1 h significantly inhibited the CRH-induced phosphorylation of p38 MAPK and JNK1/2 (Fig. 3B). These data indicate that CRH activated p38 MAPK and JNK1/2 phosphorylation in HaCaT cells, and CRH-R1 was involved in the CRH-induced phosphorylation of p38 MAPK and JNK1/2. 209 210 211 212 213 214 215 216 217 218 219 220

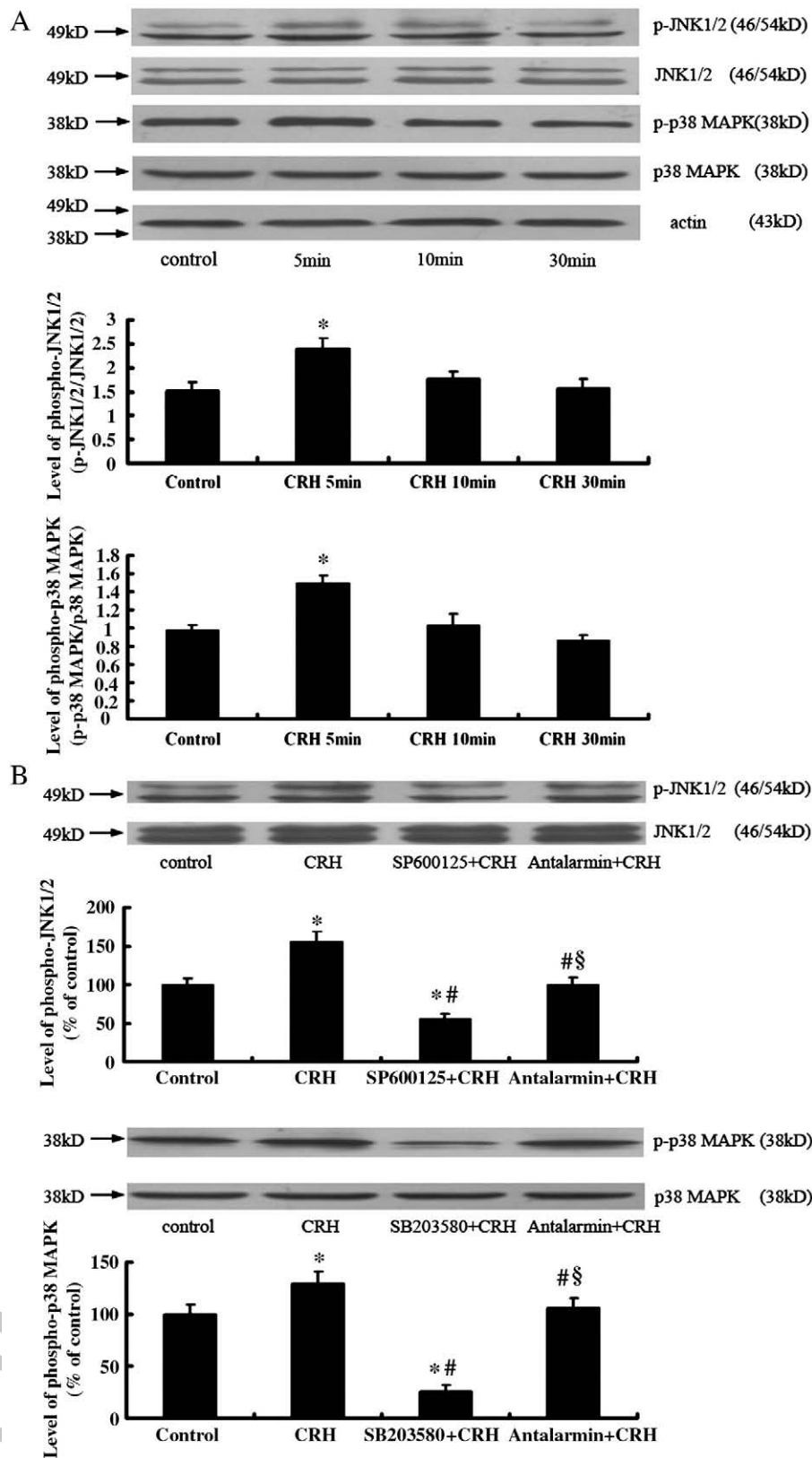
## 4. Discussion 221

In the present study, we investigated the regulatory effect of CRH on the expression of VEGF in HaCaT cells and the MAPK pathway involved. CRH attenuated VEGF expression in human HaCaT cells through CRH-R1. Furthermore, the effect of CRH on VEGF expression was associated with activation of p38 MAPK and JNK1/2 signal transduction pathways through CRH-R1. 222 223 224 225 226 227

VEGF is a major epidermis-derived vessel-specific growth factor and regarded as a potent angiogenic factor in many cutaneous diseases [34,35]. The level of VEGF is an important parameter in maintaining balanced skin angiogenesis, and it has been identified as a major keratinocyte-derived vessel-specific growth factor [36]. Keratinocytes in the lesional skin are a major source of pro-angiogenic cytokines in psoriasis; studies have identified several angiogenic factors from psoriatic epidermis, including tumor necrosis factor- $\alpha$  and VEGF [37,38]. VEGF was reported to be strongly up-regulated in psoriatic skin lesions 228 229 230 231 232 233 234 235 236



**Fig. 2.** Effect of the antagonist of CRH-R1 and MAPK inhibitors on VEGF expression induced by CRH in HaCaT cells. HaCaT cells were treated respectively with antalarmin (the CRH-R1 antagonist) and SB203580 (the inhibitor of p38 MAPK), PD98059 (the inhibitor of ERK1/2), or SP600125 (the inhibitor of JNK1/2) at 10  $\mu$ M for 1 h before CRH (100 nM). (A) Real-time RT-PCR analysis of VEGF mRNA expression after 100 nM CRH treatment for 4 h. (B) ELISA of VEGF production in cells treated with 100 nM CRH for 24 h. Data are mean  $\pm$  SD of three independent experiments performed in triplicate. \*  $P < 0.01$  vs control; #  $P < 0.01$ , §  $P > 0.05$  vs CRH-treated group.



**Fig. 3.** CRH activates p38 MAPK and JNK1/2 phosphorylation in HaCaT cells. (A) Western blot analysis of the activation of p38 MAPK and JNK1/2 pathways and phosphorylation of p38 MAPK and JNK1/2 in HaCaT cells treated with CRH. CRH induced a rapid activation of p38 MAPK and JNK1/2 as determined by phosphorylation levels. Phosphorylation of p38 MAPK and JNK1/2 peaked at 5 min. Data are mean  $\pm$  SD of three independent experiments. \*  $P < 0.05$  vs control. (B) Effect of pretreating HaCaT cells with SB203580 (the p38 MAPK inhibitor), SP600125 (the JNK1/2 inhibitor) or antalarmin (the CRH-R1 antagonist) for 1 h on the CRH-induced phosphorylation of p38 MAPK and JNK1/2. Data are mean  $\pm$  SD of three independent experiments. \*  $P < 0.05$ , §  $P > 0.05$  vs control; #  $P < 0.05$  vs CRH-treated group.

237 and may play a key role in mechanisms underlying the development of  
238 psoriasis [28,30].

239 Recently, CRH has been shown to modulate cytokine production  
240 and induce changes in growth and differentiation [15,18]; it may  
241 induce a shift away from proliferation activity and towards immu-  
242 noreactivity [16]. Previously, the pro-inflammation role of CRH was  
243 reported with CRH-induced activation of mast cells in stress-related  
244 exacerbation of cutaneous inflammatory diseases, such as psoriasis  
245 [39,40]. Meanwhile, exogenously added CRH was found to stimulate  
246 production of interleukin (IL)-6 and IL-11 and inhibit production of  
247 IL-1 $\beta$  in HaCaT cells, and CRH inhibited lipopolysaccharide (LPS)-  
248 stimulated IL-6 production [41]. CRH also could inhibit VEGF mRNA  
249 expression in cultured extravillous trophoblasts, suggesting that CRH  
250 may inhibit angiogenesis during early placentation [42]. In another  
251 study, CRH downregulated IL-18 expression, a pro-inflammatory factor  
252 [43]. Thus, CRH also has an anti-inflammatory role.

253 HaCaT cells are a unique spontaneously immortalized keratinocyte  
254 cell line derived from normal adult human skin [44]. HaCaT cells exhibit  
255 most of the characteristics of basal keratinocytes, especially hyperpro-  
256 liferation, which is typical of epidermal keratinocytes in psoriatic lesions  
257 [45]. Therefore, the line has been widely used for *in vitro* testing of anti-  
258 psoriatic compounds [46,47]. We used HaCaT cells to examine the effect  
259 of CRH on VEGF production. VEGF mRNA expression and production  
260 were significantly downregulated by CRH in a dose-dependent manner,  
261 with the maximal effect at 100 nM CRH. CRH at 100 nM was used in  
262 comparable models, such as regulation of IL-1 $\beta$  production in mono-  
263 cytes [48], suppression of the stress-related NF- $\kappa$ B pathway in human  
264 HaCaT keratinocytes [49], and in human trophoblast cells, in which  
265 CRH augmented LPS induced cytokine secretion [50]. Although CRH  
266 produced locally plays an active and pivotal role in peripheral organs,  
267 especially in regulating local homeostasis [2,15], the biological actions  
268 of CRH are mediated through interaction with CRH-Rs, which may be a  
269 central element [2]. We found the effect of CRH on VEGF expression  
270 antagonized by the CRH-R1 antagonist, antalarmin, which indicates that  
271 CRH-R1 was involved in the role of CRH in HaCaT cells.

272 To investigate the precise mechanism underlying the role of the  
273 CRH/CRH-R1 system in HaCaT cells, we studied the activation of MAPKs,  
274 the most extensively analyzed cytoplasmic signal transduction path-  
275 ways. Three main MAPKs have been well characterized: ERK1/2, p38  
276 MAPK and JNK [31,32]. In our study, pretreating HaCaT cells with  
277 SB203580 (the inhibitor of p38 MAPK) or SP600125 (the inhibitor of  
278 JNK1/2) completely abrogated the effects of CRH on the downregulation  
279 of VEGF, whereas pretreatment with PD98059 (the inhibitor of ERK1/2)  
280 did not attenuate the effects of CRH on the expression of VEGF. These  
281 data indicate that p38 MAPK and JNK1/2 but not ERK1/2 play an im-  
282 portant role in the downregulation of VEGF under the CRH/CRH-R1  
283 signaling pathway.

284 The roles of MAPKs have previously been demonstrated in psoriasis,  
285 with contradictory results. The activation of p38 MAPK and JNK1/2 is  
286 mainly related to stress and inflammatory cytokines, and the kinases  
287 therefore modulate various inflammatory responses [51]. The p38  
288 MAPK is activated in keratinocytes in psoriasis, in wound healing and in  
289 response to different stimuli such as inflammatory cytokines, UV ra-  
290 diation and oxidative stress [52,53]. JNK1/2 is activated in skin in  
291 response to UV light and other stress signals [53]. Psoriatic epidermis  
292 showed selective activation of ERK and JNK but not p38 MAPK, which  
293 might be related to hyperproliferation and abnormal differentiation of  
294 psoriatic epidermis [54]. In our previous study, we found the levels of  
295 phosphorylated ERK1/2 and p38 MAPK enhanced in lesional psoriatic  
296 skin [55]. In another study, the activity of p38 MAPK and ERK1/2 was  
297 increased in lesional psoriatic skin compared with nonlesional psoriatic  
298 skin, and clearance of psoriasis normalized the p38 MAPK and ERK1/2  
299 activity [56]. MAPK signal pathways have multiple roles in the path-  
300 ogenesis of psoriasis. In our present study, the activation of p38  
301 MAPK and JNK1/2 was involved in the downregulation of VEGF by  
302 CRH in HaCaT cells *in vitro*, which suggests an anti-angiogenic role of

CRH/CRH-R1. Although the effects of ERK activation in skin to a large  
extent parallel those of p38 MAPK [57], ERK was not involved in the  
downregulation of VEGF expression in HaCaT cells by CRH/CRH-R1.

## 5. Conclusion

In summary, we demonstrate that CRH attenuates the expression  
of VEGF through CRH-R1 and p38 MAPK–JNK1/2 signaling pathways  
in human HaCaT cells. Overall, our study provides further evidence for  
the active and pivotal role of CRH and CRH-R1 in the skin, and may  
provide an insight into the pathophysiology of neuroinflammatory  
skin diseases such as psoriasis.

## 6. Uncited reference

[58]

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