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## Ultraviolet B (UVB) activates hypothalamic-pituitary-adrenal (HPA) axis in C57BL/6 mice

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### Abstract

To test the hypothesis that ultraviolet B (UVB) can activate the hypothalamic-pituitary-adrenal (HPA) axis, the shaved back skin of C57BL/6 mice was exposed to 400 mJ/cm<sup>2</sup> of UVB or was sham irradiated. After 12 and 24 h of exposure, plasma, skin, brain, and adrenals were collected and processed to measure corticotropin-releasing hormone (CRH), urocortin (Ucn),  $\beta$ -endorphin ( $\beta$ -END), ACTH and corticosterone (CORT) or brain was fixed for immunohistochemical detection of CRH. UVB stimulated plasma levels of CRH, Ucn,  $\beta$ -END, ACTH and CORT, and increased skin expression of Ucn,  $\beta$ -END and CORT at the gene and protein/peptide levels. UVB stimulated CRH gene and protein expression in the brain that was localized to the paraventricular nucleus of the hypothalamus. In adrenal glands it increased mRNAs of melanocortin receptor type 2, StAR and CYP11B1. Hypophysectomy abolished UVB stimulation of plasma but not of skin CORT levels, and had no effect on UVB stimulation of CRH and Ucn levels in the plasma, demonstrating the requirement of an intact pituitary for the systemic effect. In conclusion, we identify mechanism of the regulation of body homeostasis by UVB through activation of the HPA axis that originates in the skin and requires pituitary for the systemic effect.

### Keywords

skin; pituitary; hypothalamus; CRH; POMC; corticosterone

### Introduction

Skin with underlying subcutis is armed with neuroendocrine capabilities and represents the largest and one of the most complex organs in the human body (Slominski *et al.*, 2012). Strategically located at the interface with external environment, skin detects, integrates, and responds to stressors including ultraviolet radiation (UVR) (Fritsche *et al.*, 2007; Paus *et al.*,

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2006; Slominski and Wortsman, 2000; Slominski *et al.*, 2012; Slominski *et al.*, 2013b). UVB (290–320 nm) radiation has powerful biological actions not only on cutaneous biology but it also impacts many regulatory pathways involved in immune homeostasis that are both vitamin D-dependent and independent (Becklund *et al.*, 2010; Campbell *et al.*, 1993; Fritsche *et al.*, 2007; Holick, 2003; Krutmann *et al.*, 2012; Ndiaye *et al.*, 2013; Pawelek *et al.*, 1992; Schwarz *et al.*, 2011). The mechanisms of UVB-induced immune suppression (Vink *et al.*, 1997) are not completely understood. However, evidence is accumulating that DNA damage and other mechanisms such as the photoisomerization of urocanic acid, free-radical formation, and signal transduction-mediated activation of transcription factors and induction of neuroendocrine signaling may also contribute to the resulting pathological conditions as well (Fritsche *et al.*, 2007; Ndiaye *et al.*, 2013; Slominski and Pawelek, 1998; Slominski *et al.*, 2012; Vink *et al.*, 1997).

The main regulatory algorithm to maintain body homeostasis is the hypothalamic-pituitary-adrenal (HPA) axis, which requires activation of a complex range of responses involving the endocrine, nervous, and immune systems, collectively known as the stress responses (Chrousos, 2009; Selye, 1976; Smith and Vale, 2006; Vale *et al.*, 1981). Neural signals encoded by the limbic system as stressors trigger neurons of the paraventricular nucleus (PVN) of hypothalamus to produce and release corticotropin releasing hormone (CRH) into the hypophyseal portal circulation (Smith and Vale, 2006; Vale *et al.*, 1981). Next, CRH binds to CRH receptor type 1 (CRH-R1) on pituitary corticotropes, and induces the release of POMC-derived adrenocorticotrophic hormone (ACTH) into the systemic circulation. The melanocortin receptor type 2 (MC2R), expressed in the adrenal cortex, stimulates glucocorticoid (GC) synthesis and secretion after binding of ACTH (Chrousos, 2009; Miller and Auchus, 2011; Smith and Vale, 2006). GC (i.e., cortisol (COR) in humans and corticosterone (CORT) in rodents) maintain metabolic and stress-responses, suppress immune activity and are self-regulated, with negative feedback to the hypothalamus and pituitary to mute the HPA axis (Chrousos, 2009; Miller and Auchus, 2011).

The skin has neuroendocrine capabilities that also encompasses all elements of the “cutaneous HPA axis” that follow the organization of the central HPA axis [reviewed in (Slominski *et al.*, 2007)]. This concept was based on the evidence that vertebrate skin expresses CRH with functional CRH-R1 [reviewed in (Slominski *et al.*, 2013b)] and related POMC macromolecule, which is further processed to ACTH (Ito *et al.*, 2005; Schauer *et al.*, 1994; Slominski *et al.*, 1998; Slominski *et al.*, 1992; Slominski *et al.*, 2000c), which, after interaction with MC2R, induces steroidogenesis with the final production of highly adaptable COR or CORT (Cirillo and Prime, 2011; Ito *et al.*, 2005; Skobowiat *et al.*, 2011; Slominski *et al.*, 2000a; Slominski *et al.*, 2013a; Slominski *et al.*, 2005a; Slominski *et al.*, 2005b; Slominski *et al.*, 2006a; Vukelic *et al.*, 2011). Furthermore, Hiramoto *et al.* (Hiramoto *et al.*, 2003) have demonstrated that exposure of the eye to the UVB increases plasma  $\alpha$ -MSH levels with systemic stimulation of epidermal melanocytes.

Based on the above we have decided to test the hypothesis that UVB acting on the skin can regulate body homeostasis through activation of central HPA. Using mouse model we evidence that UVB activates the HPA axis both on the local (skin) and systemic (brain, adrenal and plasma) levels, with a latter requiring an intact pituitary. The UVB induced

increases in corticosterone production explain immunosuppressive effects of the UVB, while that of  $\beta$ -endorphin could explain a phenomenon of “UV addiction”.

## Results

### General design of UVB exposure

To prevent retinal or non-retinal eye signal transmission, the heads including eyes, were covered with aluminum foil, and the skin on the back was irradiated with UVB (Figure 1).

### Dose and Time Dependent Effects of UVB in the Skin

Previously, we have documented that UVB (290–320 nm), but not less energetic UVA (320–400 nm), is effective in stimulating HPA axis elements in human and mouse skin organ culture *in vitro* (Skobowiat *et al.*, 2011; Skobowiat *et al.*, 2013a; Skobowiat *et al.*, 2013b). In current experiments we first measured the CORT in the skin *in vivo* using different doses and time after UVB exposure and found that the dose of 400 mJ/cm<sup>2</sup> (2.1 minimal erythema doses (MED), Table S1) and 12 and 24 h after exposure were most optimal for enhancement of the CORT levels (Figure 2a,b). Lower dose (100 mJ/cm<sup>2</sup>) and shorter times of observation (3 and 6 h) showed significantly lower stimulatory effect. Similarly, markedly stronger stimulation of plasma CORT levels was observed at 400 in comparison to 100 mJ/cm<sup>2</sup> and at 12 in comparison to 3 h after UVB exposure (Figure S1). The histopathological analysis demonstrated that UVB at 400 mJ/cm<sup>2</sup> did not produce noticeable epidermal necrosis nor trigger marked/moderate inflammatory infiltrate (Figure 2c). However, small increase in infiltrating neutrophils and eosinophils was observed at 1–6 h after UVB exposure, which after 12 or 24 h returned to the control (Figure 2d). Therefore, based on the previous (Skobowiat *et al.*, 2013a) and current *in vivo* experiments we have chosen the dose of 400 mJ/cm<sup>2</sup> (2.1 MED) and a time of 12 and 24 h after UVB exposure for further experiments.

### UVB effects on cutaneous expression of CRH and Ucn

Since the CRH gene is not expressed in C57BL/6 skin (Slominski *et al.*, 2001; Slominski *et al.*, 2013b), and can be replaced by Ucn in activating HPA axis elements (Slominski *et al.*, 2000b; Slominski *et al.*, 2013b), we examined Ucn cutaneous expression as a potential triggering regulator. We also measured the concentration of CRH since this peptide can be released locally from cutaneous nerve fibers (Roloff *et al.*, 1998; Slominski *et al.*, 1996b). UVB radiation stimulated the Ucn expression at the gene (Figure 3a) and peptide levels (Figure 3l) with the similar patterns at 12 and 24 h post-radiation. *In situ* localization studies showed increased expression of Ucn in the main skin compartments including the epidermis, adnexal structures and stratum paniculosum (Figure 3g). There was also an increase in CRH peptide concentration in the skin after UVB exposure, as evaluated by ELISA (Figure 3k).

### UVB effects on cutaneous expression of POMC, ACTH and $\beta$ -END

Next, we checked cutaneous POMC expression, a “pituitary” element of the systemic HPA axis. UVB light stimulated a 2.5-fold increase in expression of POMC mRNA after 12 h, and which was still present after 24 h, although at lower levels (Figure 3b). Immunoblotting with antibodies directed against ACTH, which recognize the 33 kDA POMC precursor,

confirmed increased expression of this molecule (Figure 3e). UVB-induced increased ACTH production was also confirmed with quantitative IHC (Figure 3h) as was expression of POMC-derived  $\beta$ -END stimulated by UVB (Figure 3i).

### UVB effects on cutaneous expression of MC2R, CYP11A1, StAR, 3 $\beta$ -HSD and CORT

The expression of the next crucial element of the HPA axis, the MC2R (responsible for initiation of steroidogenesis upon ACTH activation), was upregulated 1.5 times after 12 h and almost two times after 24 h (Figure 3c). We also investigated StAR gene expression, which is required to transfer cholesterol from the outer to the inner mitochondrial membrane, and showed that its up-regulation occurred after 12 and 24 hrs post-UVB exposure (Figure 3d). Moreover, we evaluated the expression of the rate-limiting enzyme of steroidogenesis, the cytochrome P450<sub>scc</sub> (CYP11A1), which cleaves the cholesterol side chain to produce pregnenolone, a precursor of all steroids. Western blot analysis revealed high expression of P450<sub>scc</sub> at 12 h and 24 h post-UVB exposure, compared to appropriate controls (Figure 3f). Furthermore, immunohistochemistry for 3 $\beta$ -HSD, (the enzyme that transforms pregnenolone to progesterone), showed that this antigen is highly expressed in cutaneous adnexal structures and in the stratum paniculosum, but weakly expressed in the epidermis of untreated skin. UVB radiation enhanced 3 $\beta$ -HSD expression, especially in epidermal cells (Figure 3j). The final product of HPA axis activation, CORT was produced at significantly high level at 12 h and increased further at 24 h after UVB irradiation (Figure 3m).

### UVB effects on CRH expression in the Hypothalamus

The area corresponding to the hypothalamus (Bregma ~ -0.34 to -2.70 mm) was dissected out and processed either for CRH mRNA expression or peptide measurements. There was a significant increase of CRH gene expression (Figure 4a) and CRH peptide production (Figure 4b) both 12 and 24 h after UVB exposure. Furthermore, immunohistochemistry showed an increased number of CRH-immunopositive neurons and nerve fibers surrounding the PVN area. The highest immunopositive signal was observed at 12 h after UVB exposure, and its relative values are calculated and presented as an insert to Figure 4c.

### Changes in Adrenals after UVB exposure

QPCR analyses showed that MC2R mRNA was up-regulated after 12 and 24 h of UVB radiation (Figure 5a). Similarly, StAR mRNA was increased at the same time points (Figure 5b). Expression of the CYP11B1 gene was also up-regulated but only after 24 h of UVB exposure (Figure 5c).

### Changes in Plasma after UVB exposure

CRH peptide content was highly increased at 12 and 24 h after UVB irradiation, compared to that in sham-irradiated animals (Figure 5d). A similar effect was observed for Ucn (Figure 5e). ACTH concentrations were markedly stimulated at 12 and 24 h after UVB exposure (Figure 5f). Interestingly, UVB enhanced plasma concentrations of another peptide,  $\beta$ -END (Figure 5g), that results from POMC cleavage. The final component of the HPA stress-axis in rodents, CORT, was highly elevated after 12 and 24 h post-UVB exposure (Figure 5h).

## UVB effects in Hypox Animals

First, we tested the skin, and showed that both basal and UVB-stimulated cutaneous HPA activity was pronounced in mice with intact pituitary (sham-hypox controls) than in hypophysectomized (lacking the pituitary, hypox) mice (Figure 6a). Histological evaluation showed no significant change in the skin morphology between hypox and sham-hypox mice skin after UVB exposure (Figure 6b).

Second, changes evoked by UVB at the plasma level in mice with intact pituitary were markedly different from those of mice lacking the pituitary (hypox). Although, CRH and Ucn followed similar pattern of activation in both animal groups (Figure 6c,d), hypox mice had much lower plasma levels of ACTH and CORT (Figure 5e,f) and UVB failed to stimulate plasma levels of ACTH and CORT (Figure 6e,f).

## Discussion

This manuscript shows, that the exposure of the skin to UVB can activate the systemic HPA axis culminating in increased plasma levels of corticosterone, which requires a functional pituitary. These studies parallel an important discovery by Hiramoto's group that exposure of the eyes to UVB stimulates plasma levels of  $\alpha$ -MSH and increases number of epidermal melanocytes (Hiramoto *et al.*, 2003). Although the mechanism of this stimulation remains to be established (evidence for activation of any the hypothalamic nucleus awaits experimental demonstration), the requirement for an intact pituitary was demonstrated (Hiramoto *et al.*, 2003). However, the final product of UVB induced in the eye axis is  $\alpha$ -MSH, accompanied by an increased expression of pituitary prohormone convertase 2 (Hiramoto *et al.*, 2013; Hiramoto *et al.*, 2014), which contrasts the classical HPA axis. In the HPA axis, after pituitary stimulation by hypothalamic CRH, the main product of POMC process is ACTH that acts on the adrenal glands to stimulate glucocorticosteroidogenesis (Turnbull and Rivier, 1999). While Hiramato et al show a novel mechanism of UVB activity at the central level that is different from the HPA axis, our studies demonstrate activation of the classical HPA axis by skin derived factors acting at different entry points Figure S2, for detailed discussion see below).

We also show that UVB can induce in mouse skin *in vivo* expression of all the elements of the HPA axis including CRH, POMC and corticosteroidogenic pathway. Since UVB did not induce CRH gene expression in skin cells, CRH must be delivered from nerve endings supplying dermo-epidermal or follicular junctions. This is consistent with previous studies showing increases in skin CRH after trauma1-induced hair cycling in C57BL/6 mouse (O'Kane *et al.*, 2006; Roloff *et al.*, 1998; Slominski *et al.*, 1996b; Slominski *et al.*, 2004; Slominski *et al.*, 2001; Slominski *et al.*, 1999), and detection of CRH in skin nerve bundles (Roloff *et al.*, 1998). CRH, together with Ucn (of which both gene and protein expression are up-regulated by UVB), can interact with CRH-R1 and CRH-R2 to promote local POMC activity (Slominski *et al.*, 2013b) followed by the stimulation of local steroidogenesis (Slominski *et al.*, 2013a; Slominski *et al.*, 2014). Concomitantly, UVB enhanced ACTH and  $\beta$ -END production in the skin, with increased expression of crucial regulators of steroidogenesis such as MC2R (receptor for ACTH), StAR (transporter of cholesterol) and steroidogenic enzymes (CYP11A1/P450scc and 3 $\beta$ -HSD) culminating with the final

production of CORT. Thus, UVB activates all elements of the HPA axis in mouse skin *in vivo*, which is in agreement with previous studies on UVB induction of cutaneous POMC (Chakraborty *et al.*, 1999; Schiller *et al.*, 2004) and of all or selected elements of the HPA axis in skin cells (Slominski *et al.*, 1996a; Slominski *et al.*, 2006b; Zbytek *et al.*, 2006) or skin organ culture (Skobowiat *et al.*, 2011; Skobowiat *et al.*, 2013a).

The most striking and previously unreported observations were the stimulation of CRH in the brain localized to the PVN of the hypothalamus and plasma increases of CRH, Ucn, ACTH,  $\beta$ -END and CORT that were accompanied by up-regulation of adrenal MC2R, StAR and CYP11B1. Activation of all of these elements upon skin exposure to UVB clearly indicates that activation of the systemic stress response is centered in the HPA axis and is invoked by skin signals induced by UVB. The crucial role of skin factors in this activation is documented by our experimental design that shielded the head from UVB exposure, preventing retinal signal transmission. The possible mechanisms of this activation are outlined in Figure S1, which includes both the neural route and humoral signals sent from the skin to the central regulatory elements. UVB enhancement of the production of CRH mRNA and peptide in the hypothalamus with localization at the PVN supports the hypothesis that cutaneous signals are conveyed via DRG- spinal cord- dorsal column to the PVN, the center where the HPA axis begins (Slominski *et al.*, 2013b; Smith and Vale, 2006). However, the detailed mapping of this routing is beyond the confines of this project and represents a future challenge.

The nature of humoral signaling from the skin to increase systemic CORT is more complex because in addition to production of ACTH, Ucn and CRH (see above), UVB stimulates cytokine production (IL1, IL6 and TNF $\alpha$ ) and release into circulation (Kirnbauer *et al.*, 1991; Muthusamy and Piva, 2010; Schwarz and Luger, 1989), all of which can activate the pituitary POMC (Chrousos, 2009; McEwen, 2007; Slominski *et al.*, 2000c; Turnbull and Rivier, 1999). Although dissection of potential contributions of cytokines signals can be difficult or impossible to quantitate in this experimental setting, the most logical explanation would be the activation of CRH-R1 in anterior pituitary by the circulating natural ligands: CRH and Ucn, induced by UVB, to initiate the following cascade of HPA activity:



To substantiate this hypothesis, we employed hypophysectomized mice. After UVB exposure, there was significant enhancement of CRH and Ucn peptide levels in the skin and plasma of hypox and sham-hypox animals, with plasma increases of ACTH and CORT seen only in sham-hypox (control) mice. This documents that the pituitary is necessary for UVB activation of a final element of the HPA axis in rodents, CORT. Moreover, the elevated CRH, Ucn and ACTH in the skin after UVB exposure were insufficient for direct (omitting pituitary) humoral activation of adrenals. However, they were capable of stimulating cutaneous CORT but without the corresponding increases in the plasma. Thus, UVB is another trigger of the HPA axis that requires a functional pituitary to stimulate secretion of ACTH and plasma glucocorticoids, while activation of local (skin) HPA activity is restricted to this organ without systemic involvement, as previously predicted (Slominski *et al.*, 2000c).

The net phenotypic consequences of these UVB-induced processes will include homeostatic and immunosuppressive effects resulting from the action of glucocorticoids and POMC-derived peptides, consistent with the established role and function of the HPA axis in the regulation of body homeostasis (Chrousos, 2009; Selye, 1976; Slominski *et al.*, 2013b; Smith and Vale, 2006; Turnbull and Rivier, 1999). These activities would explain the powerful systemic UVB immunosuppressive effects (Kripke, 1994; Schwarz *et al.*, 2011) as well as systemic beneficial effects in attenuation of autoimmune processes that are independent of vitamin D3 (Becklund *et al.*, 2010; Juzeniene and Moan, 2012; Schwarz *et al.*, 2012). Thus, it may be possible to attenuate severity of autoimmune diseases such as rheumatoid arthritis, multiple sclerosis, inflammatory bowel disease or scleroderma, by stimulation of endogenous glucocorticoids in an organized fashion through UVB induction of the HPA axis.

The stimulation of  $\beta$ -END levels in the skin and plasma offers a mechanistic explanation for the recently described phenomenon of “UV addiction” (Kourosh *et al.*, 2010; Nolan *et al.*, 2009), as a secondary effect of UVB-induced production of POMC-derived s-END (Slominski *et al.*, 2012). The above data on  $\beta$ -END are supported by most recent report from Dr. Fisher group showing UVB mediated stimulation of  $\beta$ -END in the skin and serum (Fell *et al.*, 2014).

In summary, we provide the evidence that UVB can activate the central HPA axis and, based on its organization and function, we propose that this mode of action represents unique mechanism regulating body homeostasis in response to UVB spectrum of solar light.

## Materials and Methods

### Animals

All procedures involving mouse experiments and tissue handling were approved by the IACUC at the UTHSC, Memphis TN. Eighty seven week-old females C57BL/6 mice ( $n=6$ ) were purchased from the Taconic Farms (Hudson, NY). After arrival, animals were kept for 4 days to avoid transportation stress with free access to standard laboratory chow and water, maintained on a 12:12 light/dark cycle and room temperature (RT) ranged from 20 to 24 C. Eight week-old animals with all hairs at the telogen stage as judged by a lack of skin pigmentation (Slominski and Paus, 1993) were used for the experiments. This was further confirmed by histological examination of the hair cycle stage confirming that hairs were at the telogen (resting-quiescence) stage where the expression of neuroendocrine factors is expected to be the lowest (Slominski and Paus, 1993; Slominski *et al.*, 2000c) and, therefore, the predicted UVB stimulation would be the highest. To test the role of the pituitary in UVB-induced HPA-activation, hypophysectomized and control (sham hypophysectomized) ninety eight ( $n=6$ ) B6 females, 7 weeks old, were purchased from Charles River Laboratories (Wilmington, MA) and were kept on 5% sucrose-supplemented water according to the vendor's guidelines.

## General Experimental Design

Details of experimental design are in Figure 1 and its legend. Shortly, after irradiation, animals (3 animals per cage) were kept for 12 or 24 h, according to the group assignment. Finally, under deep anesthesia (isoflurane 4 %) blood was collected by retro-orbital phlebotomy into EDTA tubes, and animals were euthanized by cervical dislocation. Plasma was obtained by centrifugation (1,600 g, 5 min, 4 C), skin, brain and adrenals harvested and frozen immediately at -80 C or fixed with paraformaldehyde. We repeated experiments 3 times during the spring, summer and winter to avoid confounding effects caused by the seasonal fluctuation of the HPA axis activity (Cohen *et al.*, 2012). To test the role of the pituitary, we employed the hypophysectomized (hypox) and sham operated controls (sham-hypox) mice. The animals were treated as above. Blood and tissues were collected in the morning (6 – 7 am, 12 and 24 h after irradiation).

## Irradiation

UVB (290–320 nm) irradiation was performed with a Spectroline XX-15A lamp (Spectronics Corp., Westbury, NY) equipped with a UVB waveband bulb (USHIO G15T8E) from the distance of 2.5 inches (see Table S1. The UV dosimetry was described in (Skobowiat *et al.*, 2013b). During irradiation, the bulb was covered with a cellulose triacetate sheet (Kodacel filter, Kodacel™, Eastman Kodak, Rochester, NY) which cuts-off wavelengths shorter than 290 nm, as described in (Skobowiat *et al.*, 2013b). The time of UVB irradiation was calculated upon the formula  $\text{Time (s)} = \text{Dose (J/cm}^2\text{)}/\text{Intensity (W/cm}^2\text{)}$  and presented as a standard erythema dose (SED) and MED in Table S1. One SED is equivalent to an erythemal effective radiant exposures of 100 J/m<sup>2</sup> (MKS) or 0.01 J/cm<sup>2</sup> (CGS)(International Commission on Illumination (CIE), 1999).

## Quantitative real time RT-PCR (QRT-PCR)

A detailed description of QRT-PCR is given in Supplemental materials and methods. The list of primers used for amplification with SYBR Green polymerase (Kapa Biosystems, Inc., Woburn, MA) is in Table S2. Results are presented as fold change based on the  $2^{-\Delta\Delta C_t}$  method  $\pm$  SD.

## ELISA/EIA

A detailed description is given in Supplemental materials and methods. Commercially available kits used in this study are listed in Table S3. Results are presented as mean  $\pm$  SD in either pg/mL or ng/mL after recalculations by total protein concentration (brain, skin) or dilution (plasma).

## Immunohistochemistry

Detailed protocols are in supplementary file. Briefly, the hypothalamus was isolated at the level of anterior Bregma +1 mm up to posterior Bregma -2.70 mm, by the use of the Brain Slicer Matrix (Zivic Instrument, Pittsburgh, PA). Region resembling the whole hypothalamus (Bregma ~ -0.34 to -2.70) were characterized under light microscope based on the The Allen Reference Atlas (<http://mouse.brain-map.org/static/atlas>) and 10  $\mu$ m



coronal sections were mounted onto silanized slides (Dako, Carpinteria, CA), and subjected to immunofluorescence protocols described in supplementary file.

### Western Blot

A detailed description is in (Skobowiat *et al.*, 2011). Briefly, equal amounts of protein from a combination of 3 skins for each sample was denatured with Laemli buffer, subjected to SDS/PAGE, and proteins were transferred to a PVDF membrane and incubated with antibodies listed in Table S4. Next, the membrane was incubated with secondary IgG-HRP, and detection of immunocomplexes was performed with chemiluminescence.

### Statistics

Data are presented as means  $\pm$  SD and are analyzed using Prism 4.00 (GraphPad Software, San Diego, CA). Statistically significant differences are denoted by \* (Student's *t*-test for two groups) or # (one-way ANOVA Tuckey test for more than two groups), where  $p < 0.05$  is considered as statistically significant.

### Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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### Abbreviations

<b>ACTH</b>	adrenocorticotrophic hormone
<b><math>\alpha</math>-MSH</b>	$\alpha$ -melanocyte stimulating hormone
<b><math>\beta</math>-END</b>	$\beta$ -endorphin
<b>B6</b>	C57BL/6 mice
<b>CIE</b>	Commission Internationale de L'Éclairage (International Commission on Illumination)
<b>CORT</b>	corticosterone
<b>COR</b>	cortisol
<b>CRH</b>	corticotropin releasing hormone
<b>CRH-R1</b>	CRH receptor type 1
<b>CYP11B1</b>	gene coding of Steroid 11 $\beta$ -hydroxylase
<b>P450scc</b>	cytochrome P450 side-chain cleavage enzyme

<b>DRG</b>	dorsal root ganglia
<b>MC2R</b>	Melanocortin receptor 2
<b>POMC</b>	proopiomelanocortin
<b>PVN</b>	paraventricular nucleus
<b>SED</b>	standard erythematous dose
<b>StAR</b>	steroidogenic acute regulatory protein
<b>Ucn</b>	Urocortin
<b>3<math>\beta</math>-HSD</b>	3- $\beta$ -hydroxysteroid dehydrogenase

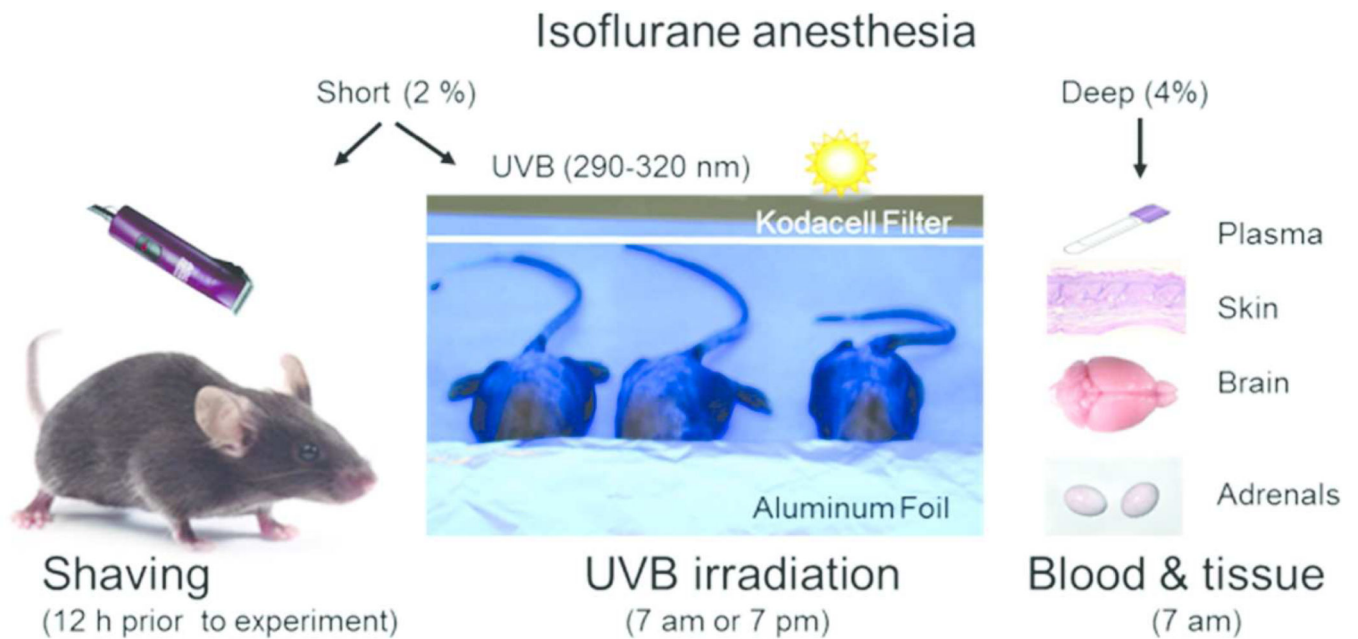
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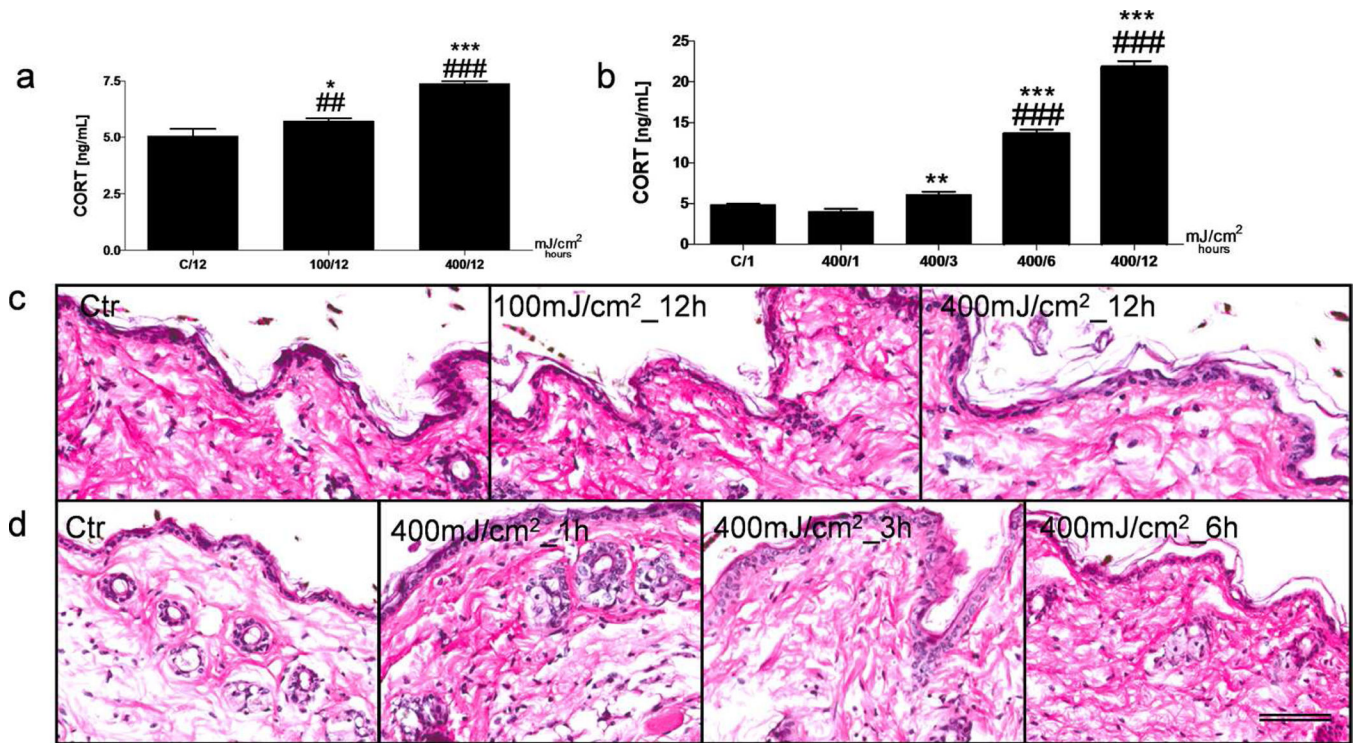
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**Figure 1.**

Outline of the experimental design. Under short (max. 1 min) vapor isoflurane (2 %) anesthesia, the back skin was shaved with animal clippers 12 h before the experimental procedures. The next day, animals were repeatedly anesthetized, eyes covered with aluminum foil (to prevent retinal signal transmission), and the skin on the back was irradiated with UVB ( $400 \text{ mJ/cm}^2$  for most experiments), either at 7 am (24 h group) or at 7 pm (12 h group). These optimal doses and times after exposure were based on initial testing of UVB at ranges of  $100 - 400 \text{ mJ/cm}^2$  vs control and time post-exposure of 3, 6, 12 and 24 hours.

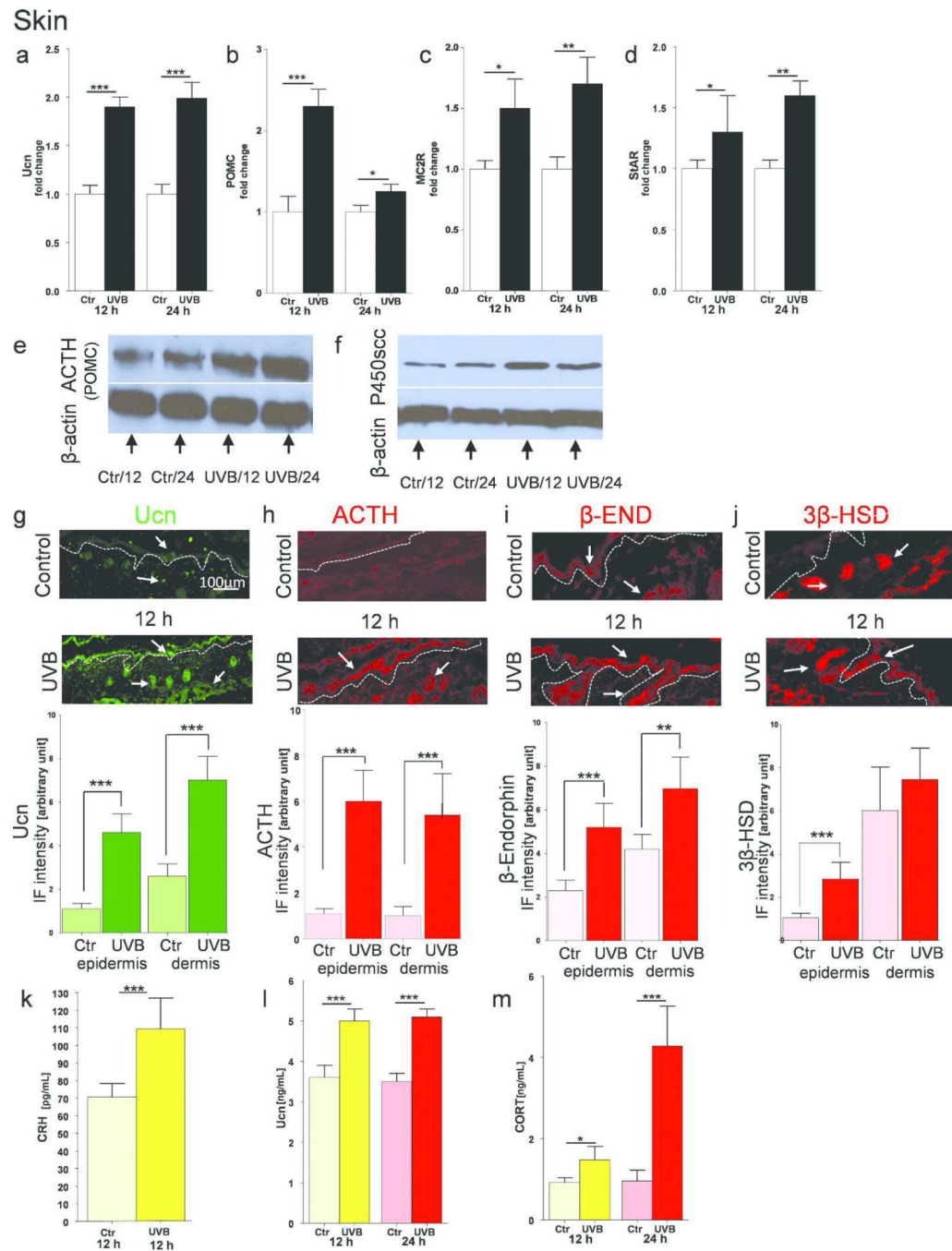


**Figure 2.**

Time and dose-dependent changes in CORT production and skin histological evaluation after UVB radiation.

Dose- (a) and time- (b) dependent increases in CORT production after UVB exposure in C57BL6 shaved mouse skin, evaluated with ELISA. A dose-dependent histological evaluation observed 12 h after UVB irradiation with a dose of 100 and 400 mJ/cm<sup>2</sup> (c).

Time-dependent histological changes evolved after 1, 3 and 6 h followed by UVB exposure of 400 mJ/cm<sup>2</sup> (d). H&E staining on formalin fixed and paraffin embedded skin representative sections. Data were analyzed using Student's *t*-test, \*  $p < 0.05$ , \*\*  $p < 0.01$ , and \*\*\*  $p < 0.001$  or one-way ANOVA, ##  $p < 0.01$ ; ###  $p < 0.001$ . Scale bar = 100  $\mu$ m.



**Figure 3.** Cutaneous equivalent of the HPA axis in C57BL6 mice is stimulated upon UVB radiation. Expression of genes coding Ucn (a), POMC (b), MC2R (c) and StAR (d) after UVB exposure compared to control (shame-treated) animals. Data presented as fold change  $\pm$  SD. Protein estimation with Western Blot for ACTH/POMC (e) and P450scc (f). *In situ* expression of Ucn (g), ACTH (h),  $\beta$ -END (i) and  $3\beta$ -HSD (j) antigens measured by immunofluorescence with corresponding quantification of immunopositive signal intensity (inserts to the subpanels). Arrows indicate examples of positive signals. ELISA evaluation



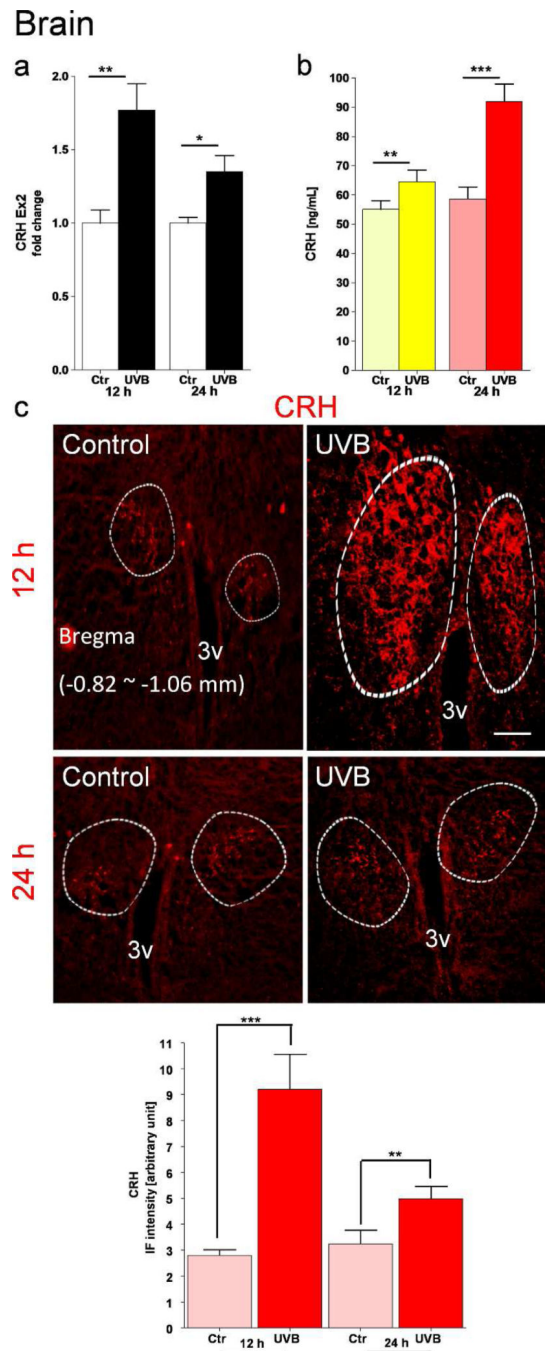
of peptide CRH (k), Ucn (l) and steroid CORT (m) concentrations. Data are presented in pg or ng/mL per 4  $\mu$ g of total proteins extracted, and analyzed using Student's *t*-test, \*  $p < 0.05$ , \*\*  $p < 0.01$ , and \*\*\*  $p < 0.001$ .

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**Figure 4.**

Up-regulation of CRH expression in the murine hypothalamus after exposure of shaved back skin to UVB. CRH gene expression is shown as fold change  $\pm$  SD (a). CRH peptide was measured by ELISA (b). Data are presented in pg/mL in relation to the same protein concentration (28  $\mu$ g/ $\mu$ L). CRH immunoreactivity *in situ* in the PVN was evaluated by immunofluorescence (c). Circled areas show *in situ* localization of CRH antigen visualized in perikarya and nerve fibers. Calculation of positive signals intensity is presented on the

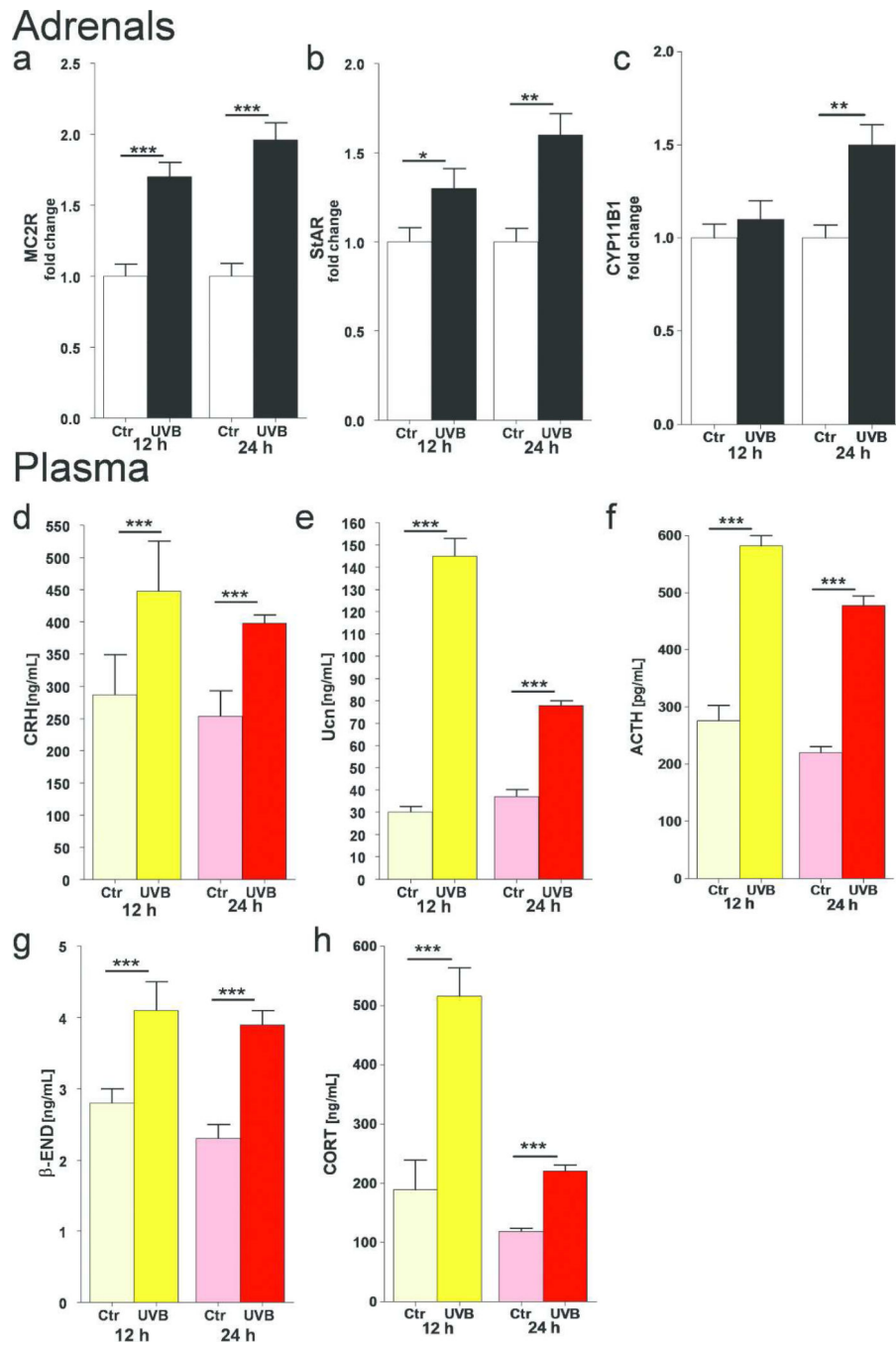
graph in the lower panel c. Data are analyzed using *t*-test, \*  $p < 0.05$ , \*\*  $p < 0.01$ , and \*\*\*  $p < 0.001$ . Scale bar = 100  $\mu\text{m}$ .

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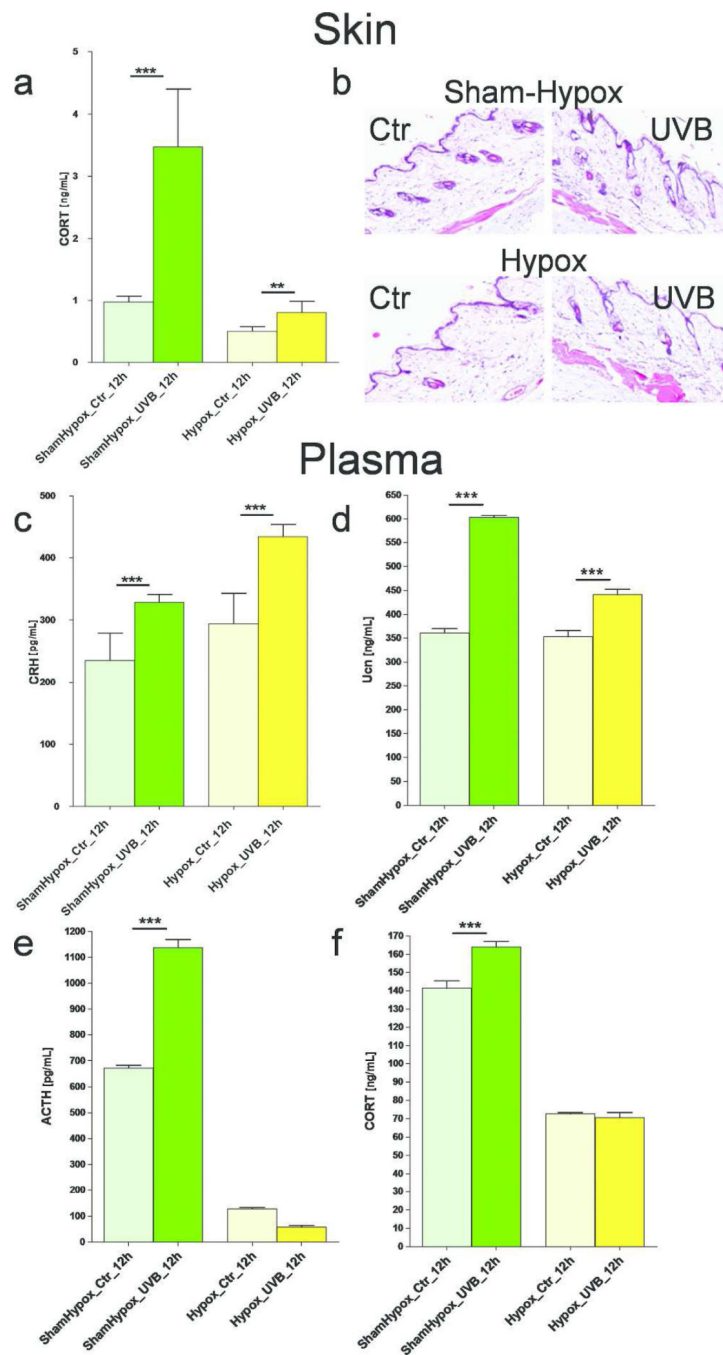
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**Figure 5.**

UVB stimulates the systemic HPA axis in C57BL/6 mice.

UVB stimulated expression of gene coding MC2R (a), StAR (b) and CYP11B1 (c) in C57BL/6 adrenals. Data are presented as fold changes  $\pm$  SD. UVB enhanced plasma concentrations of CRH (d), Ucn (e), ACTH (f),  $\beta$ -END (g), and CORT (h). Data are presented as pg or ng/mL after appropriate dilutions were performed separately for each assay and analyzed using *t*-test, \*  $p < 0.05$ , \*\*  $p < 0.01$ , and \*\*\*  $p < 0.001$ .



**Figure 6.** Differential effects of UVB on the HPA axis in hypox (pituitary removed) and sham-hypox (pituitary intact) C57BL6 mice. CORT levels were evaluated with ELISA and presented as ng/mL after adjustment of total protein content to 4  $\mu\text{g}/\mu\text{L}$  (a). Comparison of histological evaluation between hypox and sham-hypox mouse skin after UVB (400  $\text{mJ}/\text{cm}^2$ ) exposure (b). Plasma levels of peptide CRH (c), Ucn (d), ACTH (e) and CORT (f) after exposure of

shaved back skin to UVB. Data presented as pg or ng/mL after prior dilution performed separately for each assay, and analyzed using *t*-test, \*\*  $p < 0.01$ , and \*\*\*  $p < 0.001$ .

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