Topical hypochlorite ameliorates NF-κB–mediated skin diseases in mice

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Nuclear factor-κB (NF-κB) regulates cellular responses to inflammation and aging, and alterations in NF-κB signaling underlie the pathogenesis of multiple human diseases. Effective clinical therapeutics targeting this pathway remain unavailable. In primary human keratinocytes, we found that hypochlorite (HOCl) reversibly inhibited the expression of CCL2 and SOD2, two NF-κB–dependent genes. In cultured cells, HOCl inhibited the activity of inhibitor of NF-κB kinase (IKK), a key regulator of NF-κB activation, by oxidizing cysteine residues Cys114 and Cys115. In NF-κB reporter mice, topical HOCl reduced LPS-induced NF-κB signaling in skin. We further evaluated topical HOCl use in two mouse models of NF-κB–driven epidermal disease. For mice with acute radiation dermatitis, topical HOCl inhibited the expression of NF-κB–dependent genes, decreased disease severity, and prevented skin ulceration. In aged mice, topical HOCl attenuated age-dependent production of p16INK4a and expression of the DNA repair gene Rad50. Additionally, skin of aged HOCl-treated mice acquired enhanced epidermal thickness and proliferation, comparable to skin in juvenile animals. These data suggest that topical HOCl reduces NF-κB–mediated epidermal pathology in radiation dermatitis and skin aging through IKK modulation and motivate the exploration of HOCl use for clinical aims.

Introduction

Hypochlorite (HOCl) is used worldwide in homes, health care offices, and swimming pools in concentrations ranging between 0.005% and 6%. Dilute HOCl baths (0.005% v/v) are used for treating human atopic dermatitis (1), but it remains controversial whether disease improvement reflects anti-inflammatory or antimicrobial effects (2). HOCl functions as an oxidant and may interfere with cellular signaling processes by oxidizing key mediator(s). We and others have previously shown that nuclear factor-κB (NF-κB) activates proinflammatory gene programs (3, 4) and that NF-κB transcriptional activity is sensitive to intracellular reduction-oxidation (redox) conditions (5). Thus, HOCl may modulate NF-κB signaling and inflammatory responses in the skin.

NF-κB is activated by a wide variety of stimuli including TNF, LPS, and ionizing irradiation (6). NF-κB plays an integral role in important and diverse processes, including human inflammatory responses, immune system development, apoptosis, and aging. In resting cells, NF-κB is a dimeric protein sequestered in the cytoplasm via its association with inhibitory proteins called inhibitors of NF-κB (IκB). The IκB kinase (IKK) complex mediates IκB protein phosphorylation and is composed of two catalytic subunits, IKKα and IKKβ. Upon cell stimulation, IκB proteins are phosphorylated, ubiquitinated, and degraded. NF-κB then enters the nucleus and activates appropriate target genes.

Previous work has revealed a role for NF-κB in skin development, aging, and radiation response. Mice with a skin-restricted genetic NF-κB deletion (7) or with overexpression of an inhibitor of NF-κB exhibited epidermal hyperplasia, increased cell proliferation, and normal keratinocyte maturation (8, 9). Moreover, mice with a skin-restricted genetic deletion of IKKα or IKKβ also demonstrated epidermal hyperplasia and disrupted skin homeostasis (10). With age, NF-κB activity increases in many tissues. Conditional genetic inhibition of NF-κB in aged mice restored molecular and cellular features of juvenile skin, and in two different mouse models of accelerated aging, the inhibition of NF-κB delayed age-related phenotypes (11–13). Finally, ionizing irradiation initiates cellular responses driven by classical activation of NF-κB and can induce radiation dermatitis, a common adverse side effect in patients undergoing radiation therapy that has no standardized clinical treatment (10, 14, 15). Therefore, direct inhibitors of the NF-κB pathway should be useful in clinical medicine.

Here, we demonstrate that dilute HOCl oxidizes and inhibits IKK function in vitro and in vivo. This inhibitory effect was lost in cells where endogenous IKKβ was replaced with a Cys114→Cys115 to Ala114→Ala115 mutant. Strikingly, dilute HOCl baths helped treat or ameliorate two NF-κB–driven conditions in mice: acute radiation dermatitis and aging. Our findings provide a mechanistic basis for understanding the efficacy of HOCl in human diseases like atopic dermatitis and show that HOCl could be used in other NF-κB–driven disease processes.

Results

HOCl reversibly inhibits TNF-α–stimulated NF-κB–dependent gene expression in human keratinocytes. To test whether HOCl modulates NF-κB signaling, we measured the induction of established NF-κB–dependent inflammatory genes CCL2 (chemokine ligand-2 or MCP-1) and SOD2 (superoxide dismutase 2) in primary human keratinocytes stimulated with TNF-α. Exposure of human keratinocytes to 0.005% HOCl for 1 hour prior to TNF-α stimulation eliminated the induction of both genes (Figure 1A). Cell death was not increased in HOCl-treated cells, and HOCl did not alter the media pH, which can affect TNF-α responsiveness (Figure 1B). By contrast, CCL2 and SOD2 mRNA induction were unaffected in keratinocytes exposed to HOCl for 1 hour and then treated with TNF-α 24 hours later, suggesting that HOCl effects are reversible (Figure 1C). Thus,

HOCI inhibits NF-κB signaling independently of IκB. We extended these results using cultured mouse fibroblasts. HOCI blocked NF-κB–dependent gene expression induced either by TNF-α or LPS stimulation (Figure 2, A and B, and Supplemental Figure 1A; supplemental material available online with this article; doi:10.1172/JCI70895DS1), ligands that activate the NF-κB pathway through independent receptors, suggesting that HOCI inhibits intracellular NF-κB signaling. Using mouse genetics and a cell line that permits in vivo monitoring of NF-κB cellular location, we showed that HOCI blocks nuclear translocation of all NF-κB dimers, suggesting that HOCI inhibits cytoplasmic functions that regulate NF-κB activation (ref. 3 and Figure 2, C–E). HOCI blocked NF-κB–dependent gene expression in a dose-dependent manner (Supplemental Figure 1B). By contrast, HOCI induced the expression of Hmox1 (heme oxygenase 1 or HO-1), a known oxidation-sensitive enzyme, and increased p38 MAPK phosphorylation (Figure 2, F and G). Thus, while HOCI inhibited intracellular NF-κB signaling, it did not impair gene transcription and protein signaling in all pathways.

Previous work demonstrated that chloramines like HOCI can block NF-κB activation by oxidizing methionine 45 in IκBα, generating IκBβ forms resistant to TNF-α–induced degradation (16–18). We tested whether HOCI inhibited NF-κB signaling in IκBβ-deficient cells, which maintain NF-κB signaling through compensatory IκBβ activity (19). In the absence of IκBα, HOCI pretreatment still blocked TNF-α–stimulated Cxcl10 mRNA induction (Figure 2H). Western blot analysis showed that HOCI exposure inhibited TNF-α–induced degradation of both IκBα and IκBβ, suggesting that HOCI inhibits signaling steps preceding IκB protein degradation (Figure 2I). Consistent with this view, HOCI pretreatment in wild-type cells prevented TNF-α–stimulated IκBα Ser32 phosphorylation, a modification mediated by the IKK complex that precedes IκBα degradation (Supplemental Figure 1C). Thus, HOCI impeded TNF-α–stimulated phosphorylation and degradation of IκB proteins.

To test whether other oxidants also block NF-κB activation, we pretreated cultured mouse fibroblasts with hydrogen peroxide (H2O2; 90 μM) and measured TNF-α–stimulated induction of Ccl2 and Hmox1 mRNAs. H2O2 exposure induced Hmox1 transcript levels, indicating an appropriate cellular oxidative stress response. However, H2O2 pretreatment did not block TNF-α–stimulated Ccl2 expression (Figure 2J). Thus, HOCI, but not H2O2, can inhibit NF-κB activation.

HOCI oxidizes and inactivates IKK. Since IKK proteins phosphorylate and target IKK proteins for degradation, we assessed the effects of HOCI on IKK function. IKK is composed of the catalytic subunits IKKα and IKKβ. IKKα regulates noncanonical NF-κB signaling by a group of TNF-like receptors, including lymphotoxin β receptor (LTβR) (19). LTβR activates NF-κB by proteasomal processing of the NF-κB precursor protein p100 into the mature p52 subunit, thereby stimulating expression of target genes including Ccl2 (20). As expected, in LTβR-exposed wild-type cells, we observed an accumulation of p52 protein and induced Ccl2 expression. Both responses were eliminated by HOCI pretreatment (Figure 3, A and B). TNF-α–stimulated IKKβ activation requires phosphorylation at Ser177/Ser181. HOCI pretreatment blocked this phosphorylation, suggesting that HOCI inhibits IKKβ function (Figure 3C).

Collectively, HOCI inhibits both IKKα and IKKβ.

Previous studies suggest that IKK is oxidation sensitive, but the underlying mechanism remains unexplored (21, 22). Moreover, depending on the context, oxidants can activate or inhibit NF-κB (14). Thus we performed a kinase assay using activated recombinant human IKKβ and showed that HOCI inhibited IKKβ kinase activity (ref. 23 and Figure 3D, left panel). Concomitant addition of dithiothreitol (DTT), a strong reducing agent, blocked HOCI inhibition of IKKβ kinase activity (Figure 3D, middle panel), consistent with the hypothesis that HOCI inhibits IKK by oxidation. In contrast, sequential exposure of recombinant IKKβ protein to HOCI followed by DTT resulted in reduced IKKβ kinase activity (Figure 3D, right panel), suggesting that HOCI irreversibly oxidizes IKKβ. We observed earlier that HOCI-mediated inhibition of NF-κB signaling is reversible (Figure 1C), and
Figure 2
HOCl blocks intracellular NF-κB signaling independent of IkBα. (A) Relative mRNA levels of Cxcl10 in TNF-α–stimulated mouse wild-type cells with (dotted line) and without (solid line) HOCl pretreatment. (B) Relative fold changes of luciferase activity in LPS-stimulated transgenic wild-type cells carrying an integrated NF-κB reporter gene driving luciferase with (dotted line) and without (solid line) HOCl pretreatment. (C) Relative mRNA transcript levels of Ccl2 mRNA in TNF-α–stimulated p65-deficient cells with (dotted line) and without (solid line) HOCl pretreatment. (D) Visualization of TNF-α–stimulated p65-dsRed nuclear translocation (TNF-α panels; red arrows). HOCl pretreatment blocks TNF-α–stimulated p65-dsRed fusion protein nuclear translocation (HOCl and HOCl + TNF-α panels). (E) Quantification of p65-dsRed nuclear translocation per 100 cells. (F) Relative mRNA transcript levels of Hmox1 in TNF-α–stimulated mouse wild-type cells with (dotted line) and without (solid line) HOCl pretreatment. (G) Western blot analysis of phospho-p38 (P-p38) and total p38 protein in HOCl-treated cells. Black lines indicate a grouping of images from different parts of the same gel. (H) Relative mRNA levels of Cxcl10 in TNF-α–stimulated IkBα-deficient cells with (dotted line) and without (solid line) HOCl pretreatment. (I) Western blot analysis of IkBα and IkBβ protein levels in TNF-α–stimulated wild-type cells with and without HOCl pretreatment. (J) Relative Ccl2 and Hmox1 mRNA transcript levels in TNF-α–stimulated wild-type cells with (gray bars) and without (black bars) H2O2 pretreatment. Data are presented as the average ± SEM.
now we have identified an irreversible modification by HOCl. This difference is likely reconciled by ongoing IKKβ protein turnover. Consistent with this possibility, cycloheximide treatment, which inhibits protein synthesis, significantly prolonged the duration of HOCl-mediated NF-κB inhibition (Figure 3E).

**HOCl inhibits IKKβ by oxidizing cysteines 114-115.** Methionine and cysteine residues are amino acids commonly modified by oxidation (24). While methionine oxidation is reversible, oxidation of cysteine to cysteic acid is irreversible. Mass spectrometry (MS) of untreated and HOCl-treated IKKβ protein identified five cysteines that were oxidized to cysteic acid at residues 99, 114, 115, 412, and 525. Cysteines 99, 114, and 115 were conserved between IKKβ and IKKα-deficient cells (25). Each mutant cell line had similar TNF-α-stimulated Ccl2 transcript levels (26, 27). Although not identified in our analysis, Cys179 of IKKβ is shown. Alleles encoding indicated cysteine-to-alanine missense mutations are denoted as C99A, C114A/C115A, C412A, and C525A. C114A/C115A is a double mutant. (B) Relative Ccl2 mRNA transcript levels in TNF-α-stimulated wild-type cells with (gray bars) and without (black bars) cycloheximide treatment. *P = 0.004. (F) Percentage inhibition of Ccl2 transcript levels in TNF-α-stimulated IKKβ-deficient cells reconstituted with the ikkβ transgene; wild-type and four mutant transgenes of IKKβ are shown. Alleles encoding indicated cysteine-to-alanine missense mutations are shown. Alleles encoding ikkβ transgenes were denoted as C99A, C114A/C115A, C412A, and C525A. C114A/C115A is a double mutant. (B) Relative Ccl2 mRNA transcript levels in TNF-α-stimulated wild-type cells with (gray bar) or without (black bar) HOCl pretreatment in the presence of SnPP. *P < 0.001. Data are presented as the average ± SEM.

**Figure 3**

HOCl inhibits IKK function by oxidizing cysteines 114-115. (A) Western blot analysis of p52 protein levels and (B) relative Ccl2 mRNA transcript levels in LTβR-stimulated mouse wild-type cells with (black bars) and without (gray bars) HOCl pretreatment. (C) Western Blot analysis of phospho-IKKβ-Ser176/180 in TNF-α-stimulated wild-type cells with and without HOCl pretreatment. (D) Relative kinase activity of recombinant IKKβ protein with decreasing amounts of HOCl (left panel), simultaneous addition of DTT and HOCl (middle panel), and sequential addition of HOCl then DTT (right panel). (E) Percentage of induction of Ccl2 transcript in HOCl-pretreated TNF-α-stimulated wild-type cells with (gray bars) and without (black bars) cycloheximide treatment. *P = 0.004. (F) Percentage inhibition of Ccl2 transcript levels in TNF-α–stimulated IKKβ-deficient cells reconstituted with the ikkβ transgene; wild-type and four mutant transgenes of IKKβ are shown. Alleles encoding indicated cysteine-to-alanine missense mutations are denoted as C99A, C114A/C115A, C412A, and C525A. C114A/C115A is a double mutant. (G) Relative Ccl2 mRNA transcript levels in TNF-α–stimulated wild-type cells with (gray bar) or without (black bar) HOCl pretreatment in the presence of SnPP. *P < 0.001. Data are presented as the average ± SEM.
induction (Figure 3G). Thus, inhibition of NF-κB signaling by HOCl likely does not require NRF2-dependent responses.

**HOCl blocks in vivo NF-κB activation.** To assess whether HOCl could inhibit NF-κB signaling in vivo, we used an established NF-κB reporter mouse line that expresses a luciferase transgene from NF-κB-responsive cis-regulatory elements (26). Reporter mouse abdominal skin was exposed to HOCl or water (control) for 30 minutes, followed by LPS injection to activate NF-κB systemically. Two hours after LPS injection, the isolated epidermis of water-treated control mice had a 6-fold induction of reporter activity, an effect that was reduced by 50% in HOCl-treated animals (Figure 4A). Thus, a single, brief exposure to topical HOCl markedly blocked in vivo NF-κB-dependent transcriptional activation.

**HOCl ameliorates acute radiation dermatitis.** We used an established model of radiation dermatitis to test whether HOCl affects the development of acute radiation dermatitis in vivo (32). Radiation dermatitis was reproducibly induced on the back skin of 4-week-old C57BL/6 female mice with 6 Gy irradiation for 10 days. Skin was examined daily and assessed using Radiation Therapy Oncology Group (RTOG) criteria, which grades skin ulceration as the highest score (grade 4). Skin ulceration is defined as round excavations that result in the complete loss of epidermis and some portion of the dermis. To test the effect of HOCl, before each dose of ionizing radiation, we immersed the back skin of 5–7 animals in HOCl (or water for control mice) for 30 minutes. Irradiated skin in control mice initially developed erythema (days 10–12) followed by patchy moist desquamation (days 13–15) (Figure 4B, top panels, and Supplemental Figure 4). On day 20, irradiated skin began to develop skin ulcerations that continued to worsen through day 30 (Figure 4B, top panels, red arrowheads). HOCl-treated irradiated skin demonstrated mild-to-moderate erythema with intact epidermis. While the irradiated skin of all control animals eventually developed ulcerations, none of the HOCl-treated irradiated skin ulcerated, even by day 30 (Figure 4B, bottom panels). While a small area of HOCl-treated irradiated skin developed an intact scar, most of the HOCl-treated irradiated skin showed hair regrowth, indicating that intact hair follicles were maintained in the irradiated skin area (Supplemental Figure 4). This is consistent with the phenomenon of anagen effluvium, which is described in patients following radiation or chemotherapy treatment (33, 34). Overall, HOCl-treated animals had significantly reduced RTOG scores across the time course, with scores of 1.8 ± 0.24 and 4.0 ± 0 with and without HOCl treatment, respectively, by day 30 (Figure 4C).

Histological analysis of skin biopsies taken on day 14 revealed that the skin of irradiated control animals exhibited classical signs of radiation dermatitis, including a lichenoid infiltrate, loss of skin appendages, and epidermal/dermal swelling (Figure 4D, top panel). In contrast, the skin of HOCl-treated animals was largely normal, with retention of skin appendages and a mild increase in dermal cellularity (Figure 4D, lower panel; skin appendages are denoted by red arrowheads). Finally, irradiation of mouse skin induced the expression of five well-established NF-κB–dependent genes, and HOCl exposure significantly diminished this response (Figure 4E). Taken together, HOCl treatment markedly attenuated radiation-induced expression of NF-κB–dependent genes, ameliorated the severity of acute radiation-induced dermatitis and prevented skin ulceration.

**HOCl reverses the hallmark features of epidermal aging.** Aged skin is characterized by epidermal thinning and decreased proliferative capacity. In newborn mice, the epidermis comprises 3–4 cell layers and thins to 1 to 2 cell layers with age (Figure 5A, top and middle panels). To test whether HOCl inhibition of NF-κB might rejuvenate skin in vivo, we immersed the abdominal skin of 18-month-old C57BL/6 female mice in HOCl (or water for control mice) for 30 minutes daily for 2 weeks. After each treatment, both HOCl-treated and control animals were rinsed in water. HOCl exposure induced epidermal hyperplasia (Figure 5A, bottom panel). The average epidermal thickness increased more than 2-fold compared with water-treated control mice and approached the same thickness found in young mice (Figure 5B). Histology did not reveal any evidence of inflammation in the epidermis or dermis of mice exposed to HOCl. After treatment cessation, HOCl-induced epidermal thickening rapidly diminished within 1 week and was largely lost by 2 to 3 weeks (Figure 5B), suggesting that the beneficial effect is reversible and requires sustained HOCl exposure.

During normal skin development, keratinocytes in the basal layer adjacent to the basement membrane proliferate, then undergo cell-cycle arrest and migrate to outward layers, where they terminally differentiate. Consistent with the increase in epidermal thickness, HOCl treatment increased the expression of Ki-67, a cell proliferation marker, from 8% to 28% (Figure 5, C and D). Cell proliferation in HOCl-exposed animals only occurred in the basal layer of keratinocytes, the normal proliferative compartment. Similar to a study of primary fibroblasts, primary human keratinocytes treated with HOCl had significantly increased rates of cell proliferation (ref. 35 and Supplemental Figure 5). Rejuvenated skin also exhibited proper spatial organization of mature stratified epidermis, including expression of keratin 14 in the basal layer, keratin 10 in the suprabasal layer, and loricrin in the granular layer (Figure 5E). Keratin 17 is a marker of hyperproliferation and truncated epidermal differentiation (36); in HOCl-rejuvenated skin, we did not detect inappropriate keratin 17 expression in the epidermis. Rather, we found appropriate expression of keratin 17 in hair follicles and infundibulum (the opening of the hair follicle into the epidermis) (Figure 5F, white arrowheads). In skin diseases

### Table 1

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of hyperproliferation, keratin 10 expression was reduced or eliminated, but in HOCl-rejuvenated skin, keratin 10 expression was maintained (Figure 5E and ref. 37).

Skin aging induces the expression of RAD50, a DNA repair protein, and p16INK4A, a cell-cycle regulator (11, 12, 38). The expression of these two markers was significantly reduced in the epidermis of aged mice exposed to HOCl treatment to a level comparable to that found in 2-day-old mice (Figure 6, A–C). Taken together, HOCl induced epidermal hyperplasia, with increased cell proliferation and reversal of two aging-associated genes, while maintaining normal tissue homeostasis and differentiation, effects that require sustained exposure to HOCl (Figure 5B). These results are consistent with the previous findings that mice lacking NF-kB activity in the epidermis exhibit epidermal hyperplasia and increased cell proliferation, while maintaining normal keratinocyte maturation (7, 9).
sections of 18-month-old mouse skin with and without HOCl treatment. (A) H&E staining of epidermis from indicated mice. (B) Average epidermal thickness (number of cell layers) for each treatment group. Skin thickness from the 18-month-old plus HOCl group was determined 1 week and 2 weeks after cessation of HOCl (n = 3 for the HOCl treatment groups 1 week and 2 weeks after cessation of treatment). *P = 0.001. (C) Ki-67 immunostaining (green) on paraffin-embedded sections of 18-month-old mouse skin with and without HOCl treatment. (D) Percentage of basal keratinocytes expressing Ki-67. (E) Keratin 14 (K14), keratin 10 (K10), and loricrin immunostaining (green), and (F) keratin 17 (K17) immunostaining (green) on paraffin-embedded sections of skin from the 18-month-old plus HOCl treatment group. Expected staining in the lower portion of the hair follicle and hair infundibulum is denoted with white arrowheads. White dashed line in E indicates basement membrane. Nuclei in panels C, E, and F were counterstained with DAPI (blue). Original magnification, ×20 (A, C, E and F). Representative images were selected from each group. (B and D) Conditions with and without HOCl are denoted by black and gray bars, respectively. Data are presented as the average ± SEM.

Discussion

Previous studies have shown that oxidants can either induce or repress NF-kB signaling in a context- and cell type–specific manner, but little is known about the involved mechanism(s) or their potential effect in vivo (5). Our studies reveal that HOCl oxidizes IKK irreversibly at Cys114 and Cys115 to cysteic acid in order to inhibit NF-kB signaling in human keratinocytes; Cys114 and Cys115 are conserved between IKKα and IKKβ, and HOCl likely inhibits both subunits. This inhibitory effect is likely mediated by steric hindrance due to cysteic acid, and not by the disruption of disulfide bonds, since the IKKβ Cys114-115 to Ala mutants retain their inducible function. The possible effects of HOCl on targets other than IKK are not ruled out by our work; HOCl may oxidize regulatory cysteines in other proteins to affect their function. After HOCl removal, NF-kB signaling is likely restored by new synthesis of IKK protein.

Our data also suggest that oxidants can affect signaling pathways in a relatively specific manner. Although HOCl and H2O2 are in the same oxidant class and induced cellular antioxidant responses to a similar degree, only HOCl blocked NF-kB signaling. These distinct responses likely reflect intrinsic chemical differences between H2O2 and HOCl. H2O2 biological effects are mainly due to secondary products, and H2O2 is thought to covalently modify biological macromolecules only rarely. Compared with H2O2, HOCl has a relative reaction rate with glutathione that is 105-fold greater (39), indicating that HOCl is a more potent oxidant. HOCl also preferentially reacts with thiol residues, including cysteines (35). HOCl is naturally produced by neutrophils in the phagosome, and similar oxidative modifications can be found under biologically relevant conditions (40). Taken together, the ability of HOCl to modify IKK directly may explain why only some oxidants can block NF-kB signaling.

We also show that HOCl functions in vivo to attenuate two distinct and different NF-kB–driven disease processes: acute radiation dermatitis and aging. In our in vivo mouse models, we infer that HOCl is predominantly inhibiting NF-kB signaling in keratinocytes. In our radiation dermatitis experiments, many of the assessed NF-kB–dependent genes are cytokines produced in epithelial cells and known to recruit inflammatory cells; by inhibiting cytokine production, fewer immune cells can be recruited to the skin, thereby decreasing inflammation and subsequent tissue damage. In our aging model, our observations are consistent with prior studies that used conditional genetics to reduce NF-kB signaling specifically in keratinocytes for the alteration of skin homeostasis (10). Moreover, histologic analysis of rejuvenated skin did not reveal any gross changes in dermal cell populations. Thus, our data support the hypothesis that the primary effect of HOCl is on keratinocytes, but other studies are needed to assess the possibility that dermal cells are affected.

NF-kB signaling increases with age in diverse tissues, likely through a combination of genetic and epigenetic mechanisms. Heterochronic parabiosis experiments have demonstrated that
youthful features can be restored in aged tissues, which suggests that some features of aging are reversible (41, 42). We speculate that NF-κB blockade may reverse the epigenetic programs underlying skin aging. By rejuvenating aged skin to a more youthful state, HOCl may promote the treatment of skin pathologies exacerbated by aging. It is known that cutaneous wounds in the elderly heal much slower than in the young (43). HOCl may help treat chronic nonhealing wounds, including sacral decubitus pressure ulcers, trauma-induced wounds, and diabetic foot ulcers. These are common clinical problems that lack effective treatments and represent an important unmet clinical need (44). Radiation dermatitis is a common side effect in cancer patients undergoing radiation therapy that also lacks effective treatments or prevention regimens.

Bortezomib is the sole compound in clinical formularies used to target NF-κB signaling in human cancers, albeit in an indirect manner. Bortezomib inhibits proteasome function, thereby blocking IκB degradation, and treatment is associated with significant systemic toxicities. By contrast, HOCl is a safe, well-characterized, inexpensive, and widely available topical agent. Dilute concentrations of HOCl are used in swimming pools and medical clinics with minimal toxicities in humans. For example, HOCl has proven clinical efficacy in human atopic dermatitis. To our knowledge, there have been no reports of HOCl-induced premalignant skin changes. Thus, although our current study is limited to in vivo findings in mouse models, our observations should motivate clinical testing of HOCl in radiation dermatitis and aging skin. Future clinical trials will be required to assess the responses of aged skin to HOCl and to define optimal treatment strategies. Finally, our studies suggest that topical HOCl treatment may be readily extended to other NF-κB–driven inflammatory disease processes.

**Methods**

**Materials.** HOCl (Clorox) was diluted in H2O or DMEM media for a final concentration of 0.005%. We use “HOCl” to describe this mixture.

**Figure 6**

HOCl reverses markers of epidermal aging. Data from control 2-day-old mice and 18-month-old mice treated topically with H2O or HOCl for 14 days (n = 7–11 for each 18-month-old treatment group; n = 3 for 2-day-old mice). (A) p16INK4A immunostaining. (B) Quantification of p16INK4A immunostaining. *P = 0.03. (C) Relative mRNA levels of Rad50. *P = 0.001. Black bars represent HOCl treatment. Original magnification, ×20. Representative immunohistochemistry images for each group are shown. Data are presented as the average ± SEM.

**Cell culture.** Primary human keratinocytes were a gift from Paul Khavari (Stanford University). They were grown and passaged as previously described (45). IKKα, IKKβ, and IKKα/IKKβ doubly deficient MEFs were provided by Inder Verma (Salk Institute, San Diego, California, USA). Transgenic mouse fibroblasts harboring an NF-κB reporter gene driving luciferase have been previously described (4). Cells were maintained as previously described (46). Cells at 100% confluence were treated for 48 hours with medium containing 0.5% serum and then stimulated with the appropriate concentration of TNF-α (10 ng/ml; Roche), LPS (0.5 μg/ml; Sigma-Aldrich), or LTβR (2 μg/ml; Enzo Life Sciences) over a period of 0 to 6 hours. Cycloheximide (25 μg/ml; Sigma-Aldrich) or tin protoporphyrin (10 μM, Santa Cruz Biotechnology Inc.) was added to the media as indicated in Figure 2 and Figure 3E. The luciferase assay was performed according to the manufacturer’s instructions (Promega). Cell proliferation quantification was performed using CellTiter-Blue per the manufacturer’s instructions (Promega).

**Animals.** Transgenic NF-κB reporter mice harboring a luciferase reporter gene driven by NF-κB–binding sites were purchased from Taconic. C57BL/6 female mice (18–20 months of age) were purchased from The Jackson Laboratory. All mice were housed in the animal facility of Stanford University on a 12-hour light/12-hour dark cycle with ad libitum access to water and normal chow.

**Dilute HOCl baths.** Prior to the experiment, the hair of the lower abdomen or the back of each mouse was shaved closely with an electric razor. Five-liter buckets containing 500 ml of H2O or dilute HOCl (0.005%) at 37°C were prepared. One mouse was placed into each bucket; the water level allowed each mouse to stand comfortably on four feet. After 30 minutes, each mouse was removed and placed for 5 minutes into a second 5-liter bucket containing H2O at 37°C. Afterward, each mouse was dried and returned to its cage.

**In vivo NF-κB reporter activity.** NF-κB reporter mice were treated with a control or dilute HOCl bath as described above. LPS (2 mg/kg; Sigma-Aldrich) was injected via the tail vein. After 2 to 3 hours, the mice were injected with d-luciferin (~130 mg/kg; Sigma-Aldrich). The mice were euthanized 5 minutes later, and the abdominal skin was isolated. Bioluminescence imaging of the abdominal skin was performed using a Xenogen IVIS 200 system (PerkinElmer). Luminescence was measured regularly for 20 minutes, and multiple readings were collected and averaged for each sample.

**IKK kinase assay and MS.** The IKK assay was performed according to the manufacturer’s instructions (Cell Signaling Technology). MS was performed at the Stanford University Mass Spectrometry Core Facility. Protein solution samples were prepared by performing ~80°C acetone precipitation, followed by reconstitution in 8 M urea, 50 mM ammonium bicarbonate, and 0.2% ProteaseMAX solution (Promega). DTT reduction and propionamide alkylation were performed, followed by trypsin (Promega) digestion overnight at 37°C. Liquid chromatographic-mass spectrometric (LC-MS) analysis was performed, using an LTQ Orbitrap Velos mass spectrometer (Thermo Scientific) set in high-low data-depend-
dent fashion to perform tandem MS (MS/MS) on the top twelve most intense precursor ions. A Proxeon EASY-nLC (Thermo Scientific) was set at a flow rate of 300 nl/minute using a self-packed nano C18 LC column. Data were searched against the International Protein Index (IPI) Human v. 3.66 Fasta database (Matrix Science) using SEQUEST with 20 ppm precursor mass tolerance on a SORCERER platform, where data visualization and analysis were performed with Scaffold proteome software (Proteome Software).

**Real-time RT-PCR.** Cells for RT-PCR analysis were collected in TRIzol (Invitrogen). Total RNA was isolated using Direct-zol RNA MiniPrep (Zymo Research). The RNA concentration was measured by Nanodrop 1000 (Thermo Scientific). cDNA synthesis was performed with Maxima Reverse Transcriptase (Thermo Scientific) per the manufacturer's instructions. One-step quantitative RT-PCR was performed and analyzed using an ABI Prism 7300 detection system (Applied Biosystems) with TaqMan one-step RT-PCR Master Mix Reagents or SYBR Green one-step RT-PCR Master Mix Reagents (Applied Biosystems). TaqMan probes were purchased for mouse (RAD50, CCL2, CXCL10, H6, MNSOD, NFκBIA, HMOX1, and Nrf2) and human (SOD2, CCL2) genes (Life Technologies). Primer sequences for mouse genes assessed by SYBR Green chemistry are described in the Supplemental Methods.

**Western blot analysis.** Total protein was prepared from cultured cells by direct application of SDS sample buffer. Equal amounts of protein were resolved on SDS-PAGE and transferred to polyvinylidine fluoride membranes (Amersham Pharmacia) for immunoblotting with specific antibodies, including rabbit polyclonal anti–IkBα (1:1,000); rabbit monoclonal anti–phospho-IκBα (Ser176)/IKK (1:1,000); rabbit polyclonal anti–IKKβ (1:1,000); rabbit polyclonal anti–IKKα (Ser177) (1:1,000); rabbit polyclonal anti-p100/p52; rabbit monoclonal phospho-p38 MAPK (1:1,000); rabbit monoclonal p38 MAPK (1:1,000) (all from Cell Signaling Technology); and rabbit polyclonal anti–β-actin–HRP (1:5,000; Santa Cruz Biotechnology Inc.). Signal was visualized using ECL detection (Amersham Pharmacia) on Kodak film after further incubation with HRP-conjugated secondary antibodies.

**Mutant plasmids.** To generate mutant plasmids, a human IKKβ cDNA expression vector (gift from D. Goeddel, Column Group, San Francisco, California, USA) was cloned into a lentiviral vector with puromycin selection (System Biosciences). Missense mutations were created by PCR mutagenesis. Plasmid constructions were verified by DNA sequencing and restriction enzyme analysis.

**Histology, immunofluorescence, and immunohistochemistry.** Standard histological PFA fixation, paraffin embedding, and immunostaining protocols were performed. Briefly, immunohistochemical analysis was performed on 5-μm-thick sections of mouse skin after antigen retrieval (Vector Laboratories) using the following primary antibodies: mouse monoclonal anti-Ki67 (1:100; Novocastra); rabbit polyclonal anti–keratin 14 (1:100; Covance); rabbit polyclonal anti–keratin 10 (1:100; Covance); rabbit polyclonal anti–loricrin (1:100; Covance); mouse monoclonal anti-p16 (1:100; Santa Cruz Biotechnology Inc.); and rabbit monoclonal keratin 17/19 (1:100; Cell Signaling Technology). Immune complexes were detected with secondary antibodies conjugated with either Cy3, fluorescein isothiocyanate (Jackson ImmunoResearch), or HRP (Vector Laboratories). After staining, the images were directly analyzed on an AxioM1 microscope equipped with a CCD digital camera (Carl Zeiss).

Relative protein expression was quantified on a 0 to 3 scale: 0 indicates no staining; 1 signifies weak staining; 2 indicates intermediate staining; and 3 signifies strong staining. A score was obtained every 250 μm (p16INK4A) of epidermis (minimum of 5 mm) for each sample and then averaged. The number of Ki-67+ cells was counted every 250 μm of epidermis (minimum of 5 mm) for each sample and then averaged.

**NF-κB cellular localization.** p65-deficient cells harboring a p65 promoter driving p65-dsRed fusion protein were provided by Markus Covert (Stanford University). An assay was performed as previously described (47).

**Radiation dermatitis.** Radiation dermatitis was induced as previously described (32). Briefly, a 2-cm² area of skin on the back was irradiated locally at 6 Gy daily for 10 consecutive days. Ionizing irradiation was administered using a Philips RT-250 200kVp x-ray machine (12.5mA, half value layer, 1.0 mm Cu). To induce dermatitis, unanesthetized mice were placed in individual lead boxes with skin protruding through a cut-out window at the rear of each box. Skin reaction to radiation was assessed according to the RTOG score system (48). Grade 0: no change over baseline. Grade 1: follicular, faint, or dull erythema/epilation/dry; desquamation/decreased sweating. Grade 2: tender or bright erythema, patchy moist desquamation/moderate edema. Grade 3: confluent, moist desquamation other than skin folds, pitting edema. Grade 4: ulceration, hemorrhage, necrosis.

**Hydrogen peroxide experiments.** Cells at 100% confluence were treated for 48 hours with medium containing 0.5% serum. On the day of the experiment, DMEM (Hyclone) containing no serum was used to treat the cells with 90 μM hydrogen peroxide (CVS) and then stimulated with the appropriate concentration of TNF-α (10 ng/ml; Roche).

**Statistics.** A 2-tailed Student’s t test was used to determine significance, with P values of less than 0.05 considered significant. Higher levels of significance (P < 0.01; P < 0.001) are indicated in the text. For Figure 4C, we followed the nonparametric analysis of ordinal data method (49, 50). The method used was a repeated-measures ANOVA for ordinal data. We considered two factors: treatment (control vs. HOCI) and time (14, 20, and 30 days) and an interaction term (treatment × time). The treatment factor allowed us to study differences between the control and HOCI groups, the time factor allowed us to study differences among the three time points, and the interaction term allowed us to determine whether differences between the control and HOCI groups were consistent over time. The analysis was implemented using SAS Macro (SAS Institute Inc.) (50).

**Study approval.** Experiments involving mice were reviewed and approved by the IACUC of Stanford University. Mice were treated in accordance with NIH guidelines for the humane care of animals.

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