



Secretome of human umbilical cord mesenchymal stem cell maintains skin homeostasis by regulating multiple skin physiological function

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Abstract

Skin is the largest organ in the body and the first defense to resist various diseases and external stimuli that easily cause infection and inflammation. Aseptic inflammation, barrier damage, and foreign aid pressure induce the destruction and damage to the skin microenvironment. Subsequently, it destroys the skin's physiological function, leading to the maintenance and circulation of steady-state imbalance and aggravating the process of skin disorders. Our study evaluated the therapeutic potential of the secretome of human umbilical cord mesenchymal stem cells (UC-CM) for dermatological diseases in adult human skin cells, ex vivo skin tissue, and a 3D skin model. Our data suggested several advantages of UC-CM due to (1) their low cytotoxicity and sensitization properties; (2) their anti-inflammatory capacity for treating inflammatory chronic cutaneous diseases; (3) their enhanced capacity of the skin barrier for treating abnormal barrier metabolism; and (4) their positive impact on restoring skin homeostasis due to effective regulation ability of skin physiological function including cell apoptosis, detoxification, and anti-aging. We thus envisage that the possibility of harnessing the therapeutic potential of UC-CM might benefit patients suffering from inflammatory skin disorders such as atopic dermatitis, acne, and psoriasis.

Keywords Human umbilical cord mesenchymal stem cell-derived conditioned medium (UC-CM) · Inflammatory skin disorders · Anti-inflammatory · Skin barrier · Skin homeostasis

Introduction

The human skin is the largest and most complex organ that serves as a physical barrier and protects the body from being invaded (Lephart 2016). Its dysfunction leads to the development of many skin diseases, especially inflammatory skin diseases (Spergel et al. 1998; Montero-Vilchez et al. 2021). Inflammatory skin diseases, such as atopic dermatitis (AD) (Shin et al. 2020; Leshem et al. 2020; Wallach and Taïeb 2014) and psoriasis (Ring et al. 2012; Eichenfield et al. 2014), reduce the quality of life, which is a major public concern (Spergel et al. 1998). A better clinical quality scheme is still challenging for the clinical management of some skin disorders, especially for patients

who have debilitating systemic autoimmune and inflammatory skin conditions and are non-responsive to regular therapies. Mesenchymal stem cell (MSC)-based therapies can be used for treating skin diseases, as they provide new cells for epidermal homeostasis and repairing injured tissue (Wong et al. 2015; Kim et al. 2017; Guo et al. 2020). However, some disadvantages limit their application, including low implantation efficiency, unexpected differentiation risk, risk of tumor formation, short half-life, difficulties in quality control, and high storage requirements (Shin et al. 2020; Park et al. 2018; Bhang et al. 2014). MSCs provide a tissue-specific microenvironment through paracrine signaling for regulating inflammation and regeneration (Kim et al. 2017). According to some studies, the secretome from MSC (MSC-Sec) exhibits the same characteristics as MSC and might replace live cell application (Caplan 2007) to provide an opportunity to treat skin disease.

MSC-Sec, also known as a conditioned medium, contains a large number of bioactive molecules secreted by MSCs, such as proteins, microRNA, growth factors, antioxidants, proteasomes, and exosomes (Damayanti et al. 2021). These factors facilitate communication between

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cells to help skin regeneration (Le Blanc et al. 2003; Tang et al. 2004; Damayanti et al. 2021). MSC-Sec therapy has key advantages over MSC-based applications of resolved safety profiles of tumorigenicity and emboli formation. It is also more practical for clinical applications since it is convenient and suitable for skin application scenarios. Based on their accessibility, umbilical cord-derived MSCs (UCMSCs), with higher proliferation capacity, immunomodulatory activity, plasticity, and self-renewal capacity, are the second leading source used in clinical studies (Xiang et al. 2020; Cruz-Barrera et al. 2020). Comprehensive preclinical research results have shown that the mechanism of UC-CM (secretome from UCMSCs) in skin application (Montero-Vilchez et al. 2021) includes rapid wound closure (Li et al. 2017), improved proliferation and migration of various dermal cells (Li et al. 2017), protection from UV radiation by a reduction in ROS formation (Liu et al. 2013), regulation of polarized macrophages (Zhang et al. 2020b), and inhibition of inflammation and fibrosis (Xiang et al. 2020; Zhang et al. 2020b). However, most of these studies focused on wound healing and rejuvenation, while lesser attention was paid to the progress in the treatment of inflammatory skin diseases. However, two clinical trials confirmed the potential of MSC-Sec in treating inflammatory skin diseases with positive results of improvements in skin barrier function in AD patients (Kim et al. 2020) and a decrease in psoriatic plaques in psoriasis scalp patients (Seetharaman et al. 2019).

The skin is critical for ensuring homeostasis, and the loss of the integrity of the skin occurs in several inflammatory skin diseases, particularly in AD (Alwan and Di Meglio 2021). Several preclinical and clinical studies have found that UC-CM helps to maintain tissue homeostasis by repairing the skin and providing immunity against inflammatory skin diseases (Robert et al. 2019; Fernández-Gallego et al. 2021; Mackenzie 1969; Liu et al. 2019). Despite the therapeutic success of UC-CM in clinical studies, the potential targets and mechanisms of action in maintaining skin homeostasis by regulating cell function and eliminating dysfunctional cells remain unknown. Thus, for inflammatory skin diseases, a better clinical quality scheme is required before more detailed scientific studies can validate its safety and effectiveness.

In this study, we determined the role of UC-CM in maintaining the integrity of the barrier, inhibiting inflammation, and regulating the skin microenvironment, which shapes cell functions, such as cell detoxification and ECM secretion. To support our hypothesis, we established UV-induced human skin cells and ex vivo skin tissue or sodium dodecyl sulfate (SLS)-induced 3D skin model and found that UC-CM is involved in many aspects of skin physiology, which has implications in inflammatory skin disease management.

Materials and methods

Isolation, culture, identification of human UCMSC, and UC-CM preparation

Human umbilical cords were harvested after obtaining written informed consent with ethic approval. UCMSC isolation was performed in the Current Good Manufacturing Practice (cGMP)-accredited laboratory and cultured in mesenchymal stem cells basic medium (MSC-BM, Beijing Yocon Biology Co., Ltd, China) supplemented with a serum-free replacement. In the fifth passage, UCMSCs (P5) were starved for 24 h and recultured for 48 h, then UC-CM were harvested for identity test. Subsequently, UC-CM was subjected to sequential centrifugation at 800 g for 5 min and 2000 g for 10 min to remove larger particles and cell debris and stored at -20°C for further analyses.

UCMSCs were analyzed using flow cytometry to examine the expression of cell surface markers for UCMSC characterization (Alhasan et al. 2016). The MSCs were labeled with mouse anti-human FITC-CD73 (Biolegend, USA), PE-CD90 (Biolegend, USA), PE-CD34 (Biolegend, USA), APC-HLA-DR (Biolegend, USA), PE-CD45 (Biolegend, USA), PE-CD105-PE (Biolegend, USA), and FITC-CD44 (Biolegend, USA). The flow cytometry data analysis software (Flowing Software, v2.5.1, www.flowingsoftware.com) was used to plot the results. In addition, Alizarin red staining was used for osteogenic identification, Oil red O staining for adipogenic differentiation, and Alcian blue and safranin O staining for chondrogenic differentiation.

Cell culture and UV irradiation

Unless indicated otherwise, cell lines were cultured in the following media: HaCaT and HUVEC were grown in DMEM (Invitrogen, USA) supplemented with 10% FBS (Invitrogen, USA), and HSF cells in DMEM supplemented with 15% FBS. All cell lines were purchased from ATCC and grown at 37°C , 5% CO_2 in a humidified atmosphere. For some experiments, cells were subjected to UVB irradiation at 70 mJ/cm^2 by a UV lighter (Waldmann, Schwenningen, Germany) when $> 90\%$ confluency was achieved (Kim et al. 2009).

Cytotoxicity assay

HaCaT, HUVEC, and HSF cells were seeded into 96-well microplates ($100\ \mu\text{L/well}$) at a density of 1×10^4 cells/well for 24 h. According to the manufacturer's instructions, UC-CM was treated for another 48 h and tested using cell counting Kit-8 (Beyotime, China). The absorbance of the

experimental group was recorded using molecular devices (Thermo, USA) at a 450-nm wavelength and reported as cell relative growth rate (RGR%) [(average optical density of test materials-Blank)/(average negative control-Blank) × 100%] to evaluate the cytotoxicity grade according to the 5-level (0–4) definition (see the notation under Table 1) of ISO-10993–5 (Cao et al. 2009).

Skin irritation test of multiple applications

The New Zealand white rabbits purchased from Shanghai Songjiang Chedun experimental animal breeding farm Co., Ltd (Shanghai, China) were used in this experiment, and the dorsal skin of each rabbit was divided into four regions for 0.5-mL applications of UC-CM and PBS, respectively, twice a day for 14 days for irritation test. The test regions' irritation scores were obtained by assessing the extent of erythema and edema based on the criteria reported previously (Paolino et al. 2002). Average irritation scores = (erythema reaction scores + dropsy reaction scores) / amount of animals.

Morphology and inflammatory factors testing of EpiKutis inflammatory model

EpiKutis[®] (3D epidermal skin model) was purchased from Guangdong Biocell Biotechnology Co., Ltd and treated as mentioned in Table 1. EpiKutis was transferred to a 6-well plate with 0.9-mL TA culture solution (Gibco, USA), and 25 μ L 0.2% SLS (Sigma, L6026, Germany) working solution was added to the surface of negative control (NC). In the sample group, 12.5 μ L 0.4% SLS and 12.5 μ L 100% UC-CM were added to the model surface. The model medium directly replaced the blank control (BC). After treatment, all the models were incubated at 37 °C, 5% CO₂ for 24 h, and the model culture solution was collected and stored in –80 °C refrigerator until ELISA analysis. IL-1 α (IL-1 α ELISA kit, Abcam, British), TNF- α (TNF- α ELISA kit, Abcam, USA), and PGE2 (PGE2 ELISA kit, Abcam, USA) analyses were carried out according to the manufacturer's instructions. Then, the remaining liquid inside and outside the models was cleaned, fixed by 4% paraformaldehyde, and sliced for H&E staining. The skin tissues were observed and documented using a microscope.

Table 1 Test program

Group	Stimulation	Model	Testing
Black control (BC)	–	EpiKutis [®]	Tissue activity
Negative control (NC)	SLS	EpiKutis [®]	Tissue morphology Inflammatory factors
UC-CM (CM)	SLS	EpiKutis [®]	Barrier-related protein

Immunofluorescence staining of EpiKutis

After SLS was treated, EpiKutis[®] (Guangdong Biocell Biotechnology Co., Ltd, China) was ring-cut and fixed with 4% paraformaldehyde for 24 h for further use. The sample was blocked for 1 h, stained with primary antibody (LOR antibody, Abcam, USA; FLG antibody, Abcam, USA), at 4 °C overnight, and followed with FITC-conjugated secondary antibody (Goat anti-rabbit IgG, Abcam, USA; Goat anti-mouse IgG, Abcam, USA) for 1 h at room temperature. Cell nuclei were stained with Hoechst 33342 (Invitrogen, Paisley, UK), observed, and documented under the fluorescence microscope (Leica; DM2500, Germany).

Reverse transcription-quantitative (real-time) polymerase chain reaction (RT-qPCR)

According to the manufacturer's instructions, the total RNA from the cells was isolated using an RNAPure Tissue kit (CW BIO, Beijing, China), and first-strand cDNA was synthesized using a ReverTra Ace- α [®] kit (Toyobo, Osaka, Japan). Primers were designed using DNAMAN software (Lynnon Biosoft, San Ramon, USA) as follows: human collagen I (Col I) sense, 5'-5'-GAG AGC ATG ACC GAT GGA TT and anti-sense, 5'-CCT TCT TGA GGT TGC CAG TC; human MMP-1 sense, 5'-TGG ACC TGG AGG AAA TCT TG and anti-sense, 5'-GGT ACA TCA AAG CCC CGA TA; human β -actin sense, 5'-CTTCC TGGGCATGGAGTC and anti-sense, 5'-GCCGATCCACACGGAGTA. RT-qPCR was performed using Ultra SYBR Mixture (CW BIO, Beijing, China) on a QuantStudio3 PCR detection system (Thermo Fisher Scientific, USA), and the transcriptional levels of target genes were quantified using the 2^{– $\Delta\Delta$ Ct} method (Livak and Schmittgen 2001) and expressed as the means \pm SD from triplicate experiments.

Enzyme-linked immunosorbent assay (ELISA)

HaCaT or HSF cells were exposed to UVB as described above, cleaned with PBS, and cultured with or without UC-CM for 48 h. The culture was collected and measured by human IL-1 alpha ELISA Kit (BOSTER, Wuhan, China), human TNF- α ELISA Kit (BOSTER, Wuhan, China), human Col I ELISA kit (Cusabio Biotech Co., Ltd, Wuhan, China), and human MMP-1 ELISA kit (Cusabio Biotech Co., Ltd, Wuhan, China) according to the manufacturer's instructions at a 450-nm wavelength.

Western blotting

Cells were exposed to UVB, re-cultured for 24 h, and all cells and lysed were collected using T-PER Tissue Protein Extraction Reagent (Thermo Fisher Scientific, Rockford,

IL). Proteins were quantified using a BCA Protein Assay Kit (Beyotime, China), and equal amounts of proteins were loaded onto 10% gradient SDS-PAGE. After membrane transfer, blots were blocked with 5% nonfat milk in PBS for 1 h at 37 °C, and incubated with mouse anti-human FLG or anti- β -tubulin I IgG1 (Sigma, USA) overnight at 4 °C, washed, and stained with respective HRP-conjugated secondary antibodies for 1 h at room temperature. Subsequently, signals were detected using a Chemi Doc XRS+ system (Bio-Rad, Hercules, CA).

Cellular senescence detection

Natural passage aging and UVB-treated HSFs were used to detect senescence. HSF cells were continuously cultured for 5, 15, and 30 generations. The P5 generation HSF cells were treated with UVB as described above. HSF cells cultured in 6-well plates were fixed and stained with senescence cells histochemical staining Kit (Sigma, USA) at 37 °C overnight and identified using light microscopy (OLYMPUS, Japan).

Intracellular ROS generation

ROS production was quantified using flow cytometry as previously described (Leoncini et al. 1991; Maresca et al. 2010). Briefly, cells were subjected to UVB as described previously and cultured for 12 h with or without UC-CM at different concentrations and detected using a DCFH-DA kit (Beyotime, China). Briefly, cells were incubated with DCFH-DA (1 mM) at 37 °C in the dark for 30 min, washed thrice immediately, and resuspended in PBS. Intracellular ROS production was measured using a flow cytometer (BD FACSCelesta, USA).

Ex vivo skin culture, UVA/UVB exposure, and Masson staining

Ex vivo skin was obtained from Guangdong Biocell Biotechnology Co., Ltd (Guangzhou, China). The freshly obtained skin tissue was immersed in 75% alcohol, washed for 30 s, and subsequently washed with PBS. The skin tissue was cut into small disks and placed into the culture mold, with the epidermis facing up and the dermis facing down. The culture mold was then transferred into a 6-well plate containing an isolated skin tissue culture medium (FSK4, Gibco, USA) and incubated at 37 °C with 5% CO₂. After 24-h incubation, isolated skin tissues were subjected to UVA 30 J/cm² and UVB 50 mJ/cm² for 4 days. After each irradiation, the culture medium of the negative control group was changed, and the sample group was replaced with the corresponding working fluid and added UC-CM (2 μ L) on the tissue surface. After 4 days of continuous irradiation, the skin tissue in vitro was cultured for 3 days, stained using Masson staining, and

observed under a microscope. Micrographs were processed using Image-Pro[®]Plus Image processing software to analyze the collagen fibers.

Statistical analysis

All experiments were performed in triplicates. The results were reported as the average value standard deviation unless otherwise stated. Data were statistically analyzed using the Prism 5 software program and compared using Student's *t*-test. Data with *p* < 0.05 was considered statistically significant.

Results

Characterization and identification of UCMSC

MSC activity affects the active secretome in a conditioned medium (UC-CM). Therefore, we first characterized the characteristic molecule markers of UCMSC and verified its differentiation ability in vitro by detecting adipogenic, osteogenic, and chondrogenic differentiation ability. Immunophenotypes of MSC showed a high expression of CD73 (99.8%), CD90 (100%), CD105 (98.4%), and CD44 (97.6%) and a low expression of CD45 (0.097%), CD34 (0.94%), and HLA-DR (0.015%) (Fig. 1a–b"). In addition, the positive results of Oil red O, Alizarin red S, Alcian blue, and safranin O staining proved the multi-lineage differentiation potential of UCMSC (Fig. 1c–f). Thus, this indicated that UCMSC used to produce UC-CM had good activity and characterization.

Detection of cytotoxicity and skin irritation of UC-CM

A cytotoxicity assay and multiple skin irritation tests at cell or animal levels were conducted to evaluate the safety of UC-CM in skin application. The cytotoxicity of UC-CM after 48-h cell treatment is presented in Table 2 and Fig. 1g. All experimental groups showed no (toxicity grade=0) or low cytotoxicity (toxicity grade = 1) for three cell types (HaCaT\HSF\HUVeC). The concentration of UC-CM under 50% could promote cell proliferation compared with negative control (NC), while at 70% or 100%, it could decrease cell proliferation in HaCaT and HSF cells, except HUVeC cells.

The results of the skin irritation test for multiple applications of UC-CM (14 days) in New Zealand rabbits are presented in Table 3 and Fig. 1h–h''', indicating no irritation to normal skins after continuous applications for 14 days compared with NC. In brief, the current experiment corroborated the low cytotoxicity and no-irritation of UC-CM to skin local applications, and thus the concentration of UC-CM at 50 or 100 percent was used in the subsequent experiments.

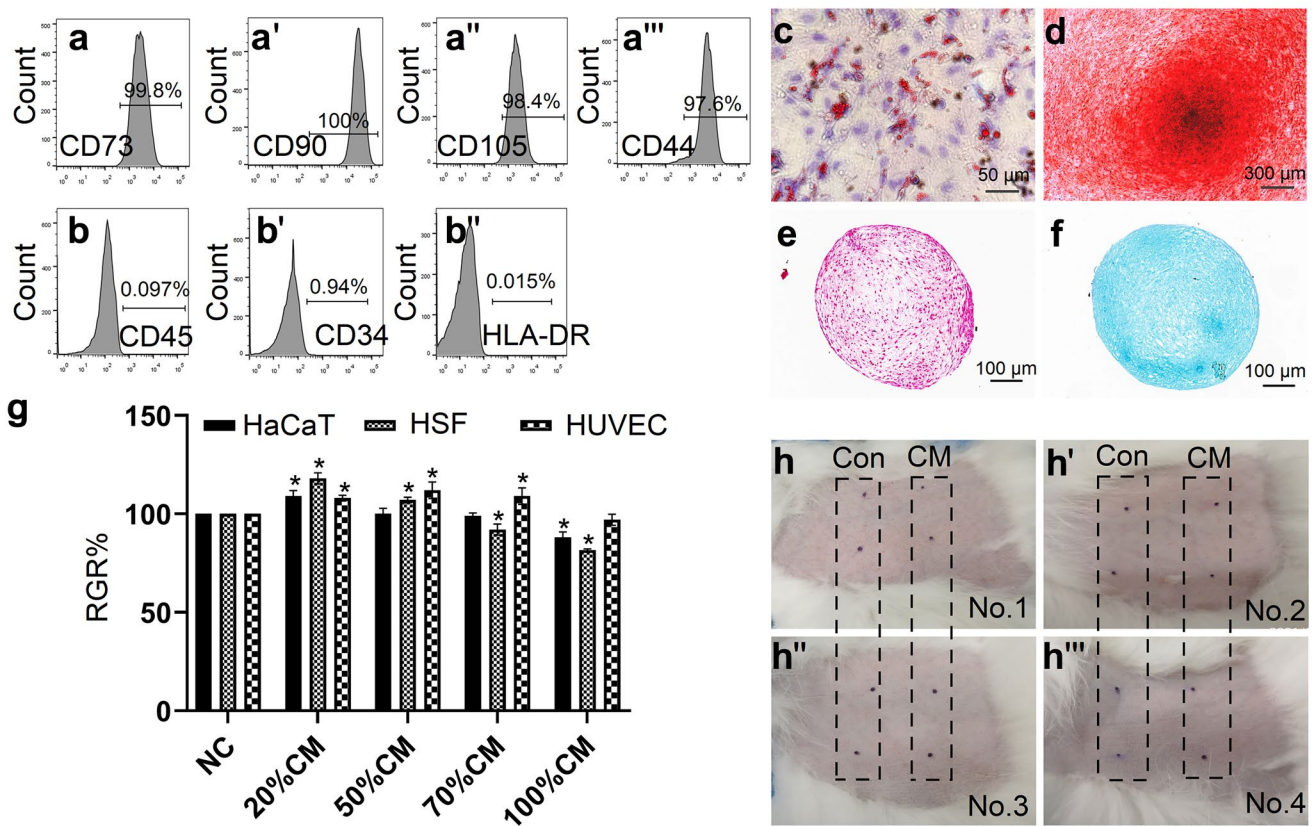


Fig. 1 Characterization of UCMSCs and safety evaluation of UC-CM. **a–b''** UCMSCs were positive for CD73, CD90, CD105, and CD44, and negative for CD45, CD34, and HLA-DR, as shown by flow cytometry analysis. The multi-lineage differentiation of UCMSCs into **c** adipocytes (Oil red O staining, upper left, scale bar: 50 μ m), **d** osteoblasts (Alizarin red S staining, upper right, scale bar: 300 μ m), **e** and **f** chondrocytes (safranin O staining, lower left and Alcian blue staining, lower right,

scale bar: 100 μ m). **g** Detection of cytotoxicity of UC-CM. HaCaT, HSF, and HUVEC cells were treated with or without UC-CM for 48 h. The proliferation of all cells was detected using the CCK8 test, and the relative growth rate (RGR%) was calculated. *vs. corresponding NC, $p < 0.05$. **h, h', h'', h'''** Skin irritation test of multiple applications. Photographs of multiple skin irritation tests in New Zealand rabbits at day 14. Control = PBS, CM = 100% UC-CM, $n = 4$

Therapeutic potential of UC-CM by anti-inflammatory effect and improving skin barrier function in the 3D dermal model

A 3D dermal model (EpiKutis®) exposed to 0.2% SLS treatment which can enhance oxidative stress and increase the

release of inflammatory factors, such as interleukin (IL)-1 α , prostaglandin E2 (PGE2), destroys the skin barrier and leads to the occurrence of skin inflammatory reaction was used to determine the therapeutic potential of UC-CM in the skin. Firstly, the effect of UC-CM on the tissue morphology and viability of the 3D skin model was examined. There was

Table 2 The relative growth rate of cells and cytotoxicity grade

Group	RGR (%)			Toxicity grade ^a		
	HaCaT	HSF	HUVEC	HaCaT	HSF	HUVEC
20% UC-CM	109*	118*	108*	0	0	0
50% UC-CM	100	107*	112*	0	0	0
70% UC-CM	99	92*	109*	1	1	0
100% UC-CM	88*	83*	97	1	1	1
NC	100	100	100	0	0	0

NC negative control

^aToxicity grade 0 for RGR% $\geq 100\%$, grade 1 for $80\% \leq RGR\% \leq 99\%$, grade 2 for $50\% \leq RGR\% \leq 49\%$, and grade 4 for $0\% \leq RGR\% \leq 29\%$

* vs NC, $p < 0.05$

Table 3 Average response scores of skin irritation for multiple applications ($n = 4$)

Number of animals	Skin irritation average score ^a of 14 days					
	Control			UC-CM		
	Erythema	Edema	Score	Erythema	Edema	Score
4	0	0	0	0	0	0

^a0 for no visible reaction, 1 for just present reaction, 2 for slight reaction, 3 for moderate reaction, and 4 for a severe reaction. Scores of < 0.5 could be considered as no irritation

no significant difference in the tissue viability of the 3D skin model when treated with UC-CM compared with the NC group (Fig. 2a). Simultaneously, H&E staining was performed on the model after culture to evaluate UC-CM's ability to repair barrier damage. As reported, 0.2% SLS resulted in visible tissue damage, with the stratum granulosum being no longer distinguishable from the stratum spinosum in EpiDerm cultures (Gibbs et al. 2002). In the 3D dermal model, compared with the BC group, the NC group had indistinguishable granular layer and spinous layer, loose and thickened cuticle (marked by yellow arrow), decreased number of living cell layers (marked by green arrow), and vacuoles appeared (black circle), indicating an effective SLS stimulation condition (Fig. 2b–b"). Compared with the NC group, even though the number of viable cell layers did not improve, the tissue morphology improved by fewer vacuoles and thinner cuticles after UC-CM administration.

Meanwhile, three important inflammatory factors were detected in the 3D skin model. The IL-1 and tumor necrosis factor (TNF) families of cytokines play important roles in immune regulation and inflammatory processes (Barksby et al. 2007; Hänel et al. 2013). PGE2 is a major inflammatory mediator involved in skin disorders (Heard 2020). As shown in Fig. 2c–e, the content of IL-1 α , TNF- α , and PGE2 in the NC group significantly increased ($p < 0.01$) compared with the BC group. Interestingly, the content of IL-1 α and TNF- α significantly decreased compared with the NC group, while the PGE2 content did not change significantly after UC-CM treatment in the 3D skin model. The content changes of two skin barrier proteins, filaggrin (FLG) and loricrin (LOR), in the 3D skin model by immunofluorescence staining were compared to directly evaluate the role of UC-CM in repairing damaged skin barrier function. Dysregulation of FLG and LOR leads to epidermal barrier

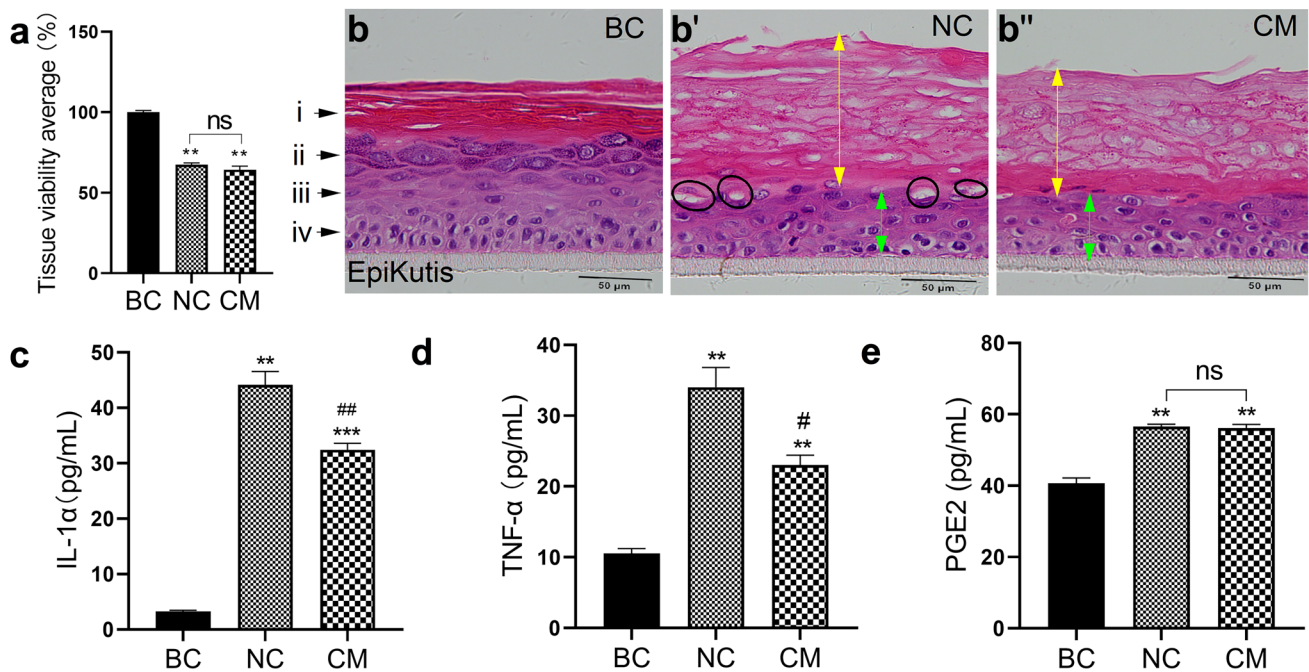


Fig. 2 Tissue viability, morphology, and inflammatory factor of 3D EpiKutis[®] model. Viability (a) and histological examination (b, b', b'') analysis of EpiKutis model. i: Stratum corneum; ii: granular layer; iii: spinous layer; iv: basal layer. Yellow arrow: cuticle; green arrow: living

cell layer; black circle: vacuoles. Scale bar: 50 μ m. BC, black control; NC, SLS treatment; CM, SLS + UC-CM treatment. ELISA for IL-1 α (c), TNF- α (d), and PGE2 (e). **vs BC, $p < 0.01$, *** $p < 0.001$; #vs NC, $p < 0.05$, ## $p < 0.01$; ns, not significant

defects, resulting in skin fragility and sensitivity (Bigliardi et al. 2016; Odou et al. 2001). LOR and FLG were detectable with a strong signal in the BC group (Fig. 3a and d) but undetectable in the NC group (Fig. 3b and e). It is worth noting that a positive signal of FLG (Fig. 3c and g, $p < 0.001$) and a negative result of LOR (Fig. 3f and h, ns) were found in the CM group compared with the NC group. The data suggested that IL-1 α , TNF- α , and FLG play an important role in anti-inflammatory and barrier repair by using UC-CM.

UC-CM abrogated the skin microenvironment disorder due to UV radiation by enhanced detoxification ability

Excessive exposure to UV can damage various biological molecules, induce inflammatory responses, and increase the oxidative stress level, contributing to the skin microenvironment imbalance. Firstly, UVB-irradiated HaCaT cells were used to confirm the results obtained in the 3D model. Cells showed a 147% ($p < 0.0001$) higher IL-1 and 165% ($p < 0.0001$) higher TNF- α than non-irradiated cells (Fig. 3i and j). Treatment with 50% or 100% UC-CM led to a 22% ($p < 0.05$) and 28% ($p < 0.01$) decrease in IL-1 level and a 29% ($p < 0.01$) and 43% ($p < 0.001$) decrease in TNF- α , respectively,

compared to the UV group. A decreased level of FLG was observed at protein level detecting by western blotting in the UV cell models associated with a slightly increased expression of FLG after UC-CM treatment (Fig. 3k).

The detoxification ability of the skin to deal with external pressure is reflected in its ability to eliminate stress responses, such as inhibition of cell apoptosis and ROS generation (Li et al. 2019). Thus, the detoxification ability of UC-CM at the cell level using UVB-irradiated HaCaT and HSF was discussed. Too little proliferation might yield thinner skin and loss of protection. As we found that UC-CM could improve cell proliferation (Fig. 1g), the study focused on its positive effects on decreasing cell apoptosis. The results showed that 50% and 100% UC-CM were able to protect the cells from UV-induced apoptosis (Fig. 4a–b''') with average percentage of apoptosis 2.59% ($p < 0.0001$) and 1.01% ($p < 0.0001$) in HaCaT cells (Fig. 4c) or 9.74% ($p < 0.0001$) and 2.92% ($p < 0.0001$) in HSF cells (Fig. 4d) compared with the sole UV treated group (7.48% in HaCaTs or 16.68% in HSFs). Furthermore, our data indicated that 50% or 100% UC-CM was able to inhibit ROS production with relative mean fluorescence intensity of 2.80 ($p < 0.01$) and 1.24 ($p < 0.001$) in HaCaTs (Fig. 5a and c) or 1.42 ($p < 0.01$) and 1.2 ($p < 0.01$) in HSFs (Fig. 5b and d) compared

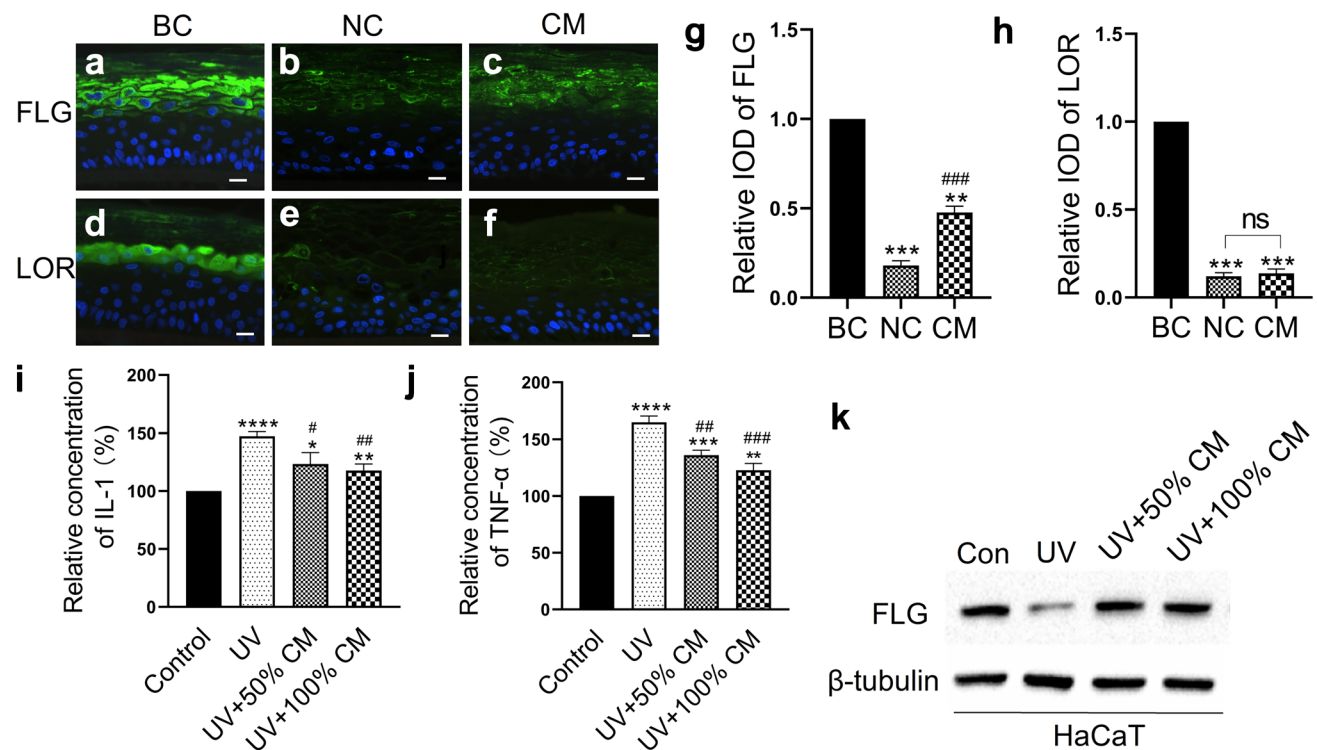
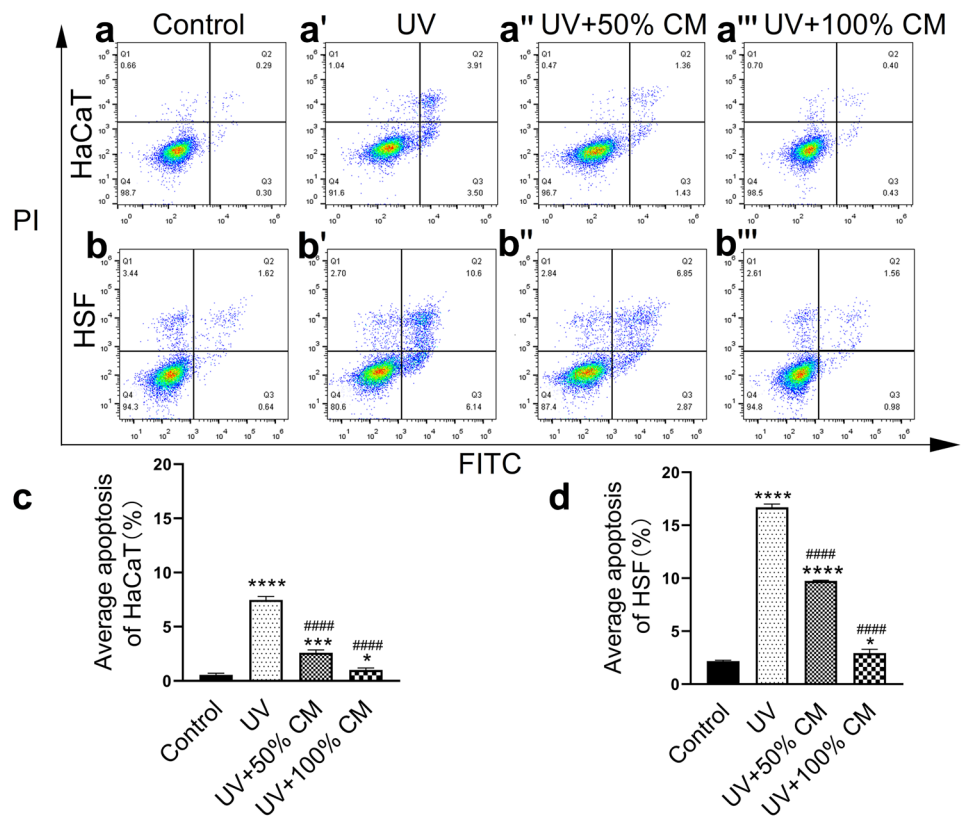


Fig. 3 Barrier protein expression of 3D EpiKutis[®] model and verification of UV-irradiated HaCaT cell. **a–f** Image of immunofluorescence staining of FLG and LOR and relative IOD analysis of FLG (g) and LOR (h). Scale bar: 20 μ m; BC, black control; NC, SLS treat-

ment; CM, SLS + UC-CM treatment. **i–j** ELISA for IL-1 and TNF- α in HaCaT cell. **k** Western blotting for HaCaT. Con, control. *vs BC, $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$; #vs NC, $p < 0.05$, ## $p < 0.01$, ### $p < 0.001$; ns, not significant

Fig. 4 Effects of UC-CM on cell function by reducing cell apoptosis. Analysis of cellular apoptosis by flow cytometry analysis **a–b'''** and apoptotic ratio of cells under different treatments (**c, d**). *vs control, $p < 0.05$, *** $p < 0.001$, **** $p < 0.0001$; ####vs UV, $p < 0.0001$



with the UV group, and reduce the oxidative toxicity caused by the increase of free radical production source.

UC-CM protect HSFs from oxidative stress and contribute to maintaining skin homeostasis

Oxidative stress produces inflammatory factors and senescent cells that destroy the microenvironment, finally accelerating extracellular matrix (ECM) disassembly, providing pressure for the healthy maintenance of the skin (Koentges et al. 2019; Xiong et al. 2016; Bao et al. 2020). ECM, mainly secreted by skin fibroblasts, plays a crucial role in maintaining skin elasticity, improving the skin microenvironment, and strengthening the epidermal barrier. ECM's ability for secretion and senescent cell deletion were evaluated in HSF cells. As one of the most important ECM components, collagen participates in cell physiological function. mRNA levels of *col-1* and *mmp-1*, two important contents of ECM responsible for collagen synthesis and decomposition, were evaluated using RT-qPCR. Our experiment indicated that UV irradiation significantly decreased the *col-1* expression and increased *mmp-1* expression (Fig. 6a and b). This phenomenon was reversed by UC-CM protection followed by upregulation of *col-1* and downregulation of *mmp-1* at the mRNA level, consistent at the protein level using ELISA assay (Fig. 6c and d). The above results were further confirmed by the Masson

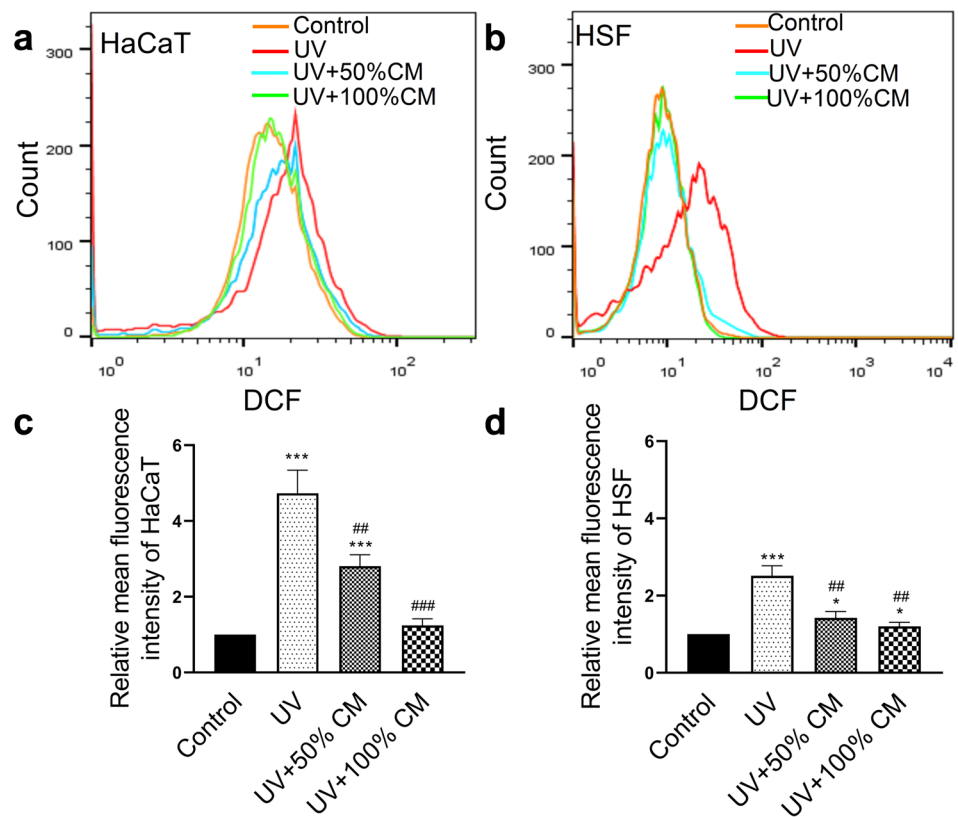
staining of collagen fibers in UV-irradiated ex vivo skin tissue. As shown in Fig. 6e–e' and f, the collagen fiber content in the NC group significantly decreased compared with the BC group, whereas it significantly increased in the CM group compared with the NC group. Thus, suggesting that UC-CM could protect skin from ultraviolet damage, adjust ECM composition, and strengthen skin defense.

The activity of SA- β -gal in natural passage HSF cells and UV-irradiated HSF cells with or without UC-CM treatment was evaluated. In young HSF culture, all groups' senescent HSF cell ratio was low. In the intermediate (P15) and senescent (P30) group, the senescent HSF cell ratio was lower than in the control group ($p < 0.05$ in P15; $p < 0.01$ in P30) after UC-CM treatment (Fig. 7a–c'' and d). Protection by 50% UC-CM decreased senescent ratio in UV-irradiated HSF cells ($p < 0.05$) than the UV group, while HSF cells exhibited similar trends to 100% UC-CM protection group ($p < 0.05$, Fig. 7e–e''' and f).

Discussion

MSC-Sec is considered to be a therapeutic agent because of its promising clinical features in skin disorders (Kim et al. 2020; Seetharaman et al. 2019). However, the mechanism by which they treat inflammatory skin disorders is still unclear.

Fig. 5 Effects of UC-CM on cell function by reducing ROS generation. Reactive oxygen species (ROS) intracellular level reduction and relative analysis in HaCaT cells (**a, c**) and HSF cells (**b, d**). Data are representative of three separate experiments. *vs control, $p < 0.05$, *** $p < 0.001$; ##vs UV, $p < 0.01$, ### $p < 0.001$



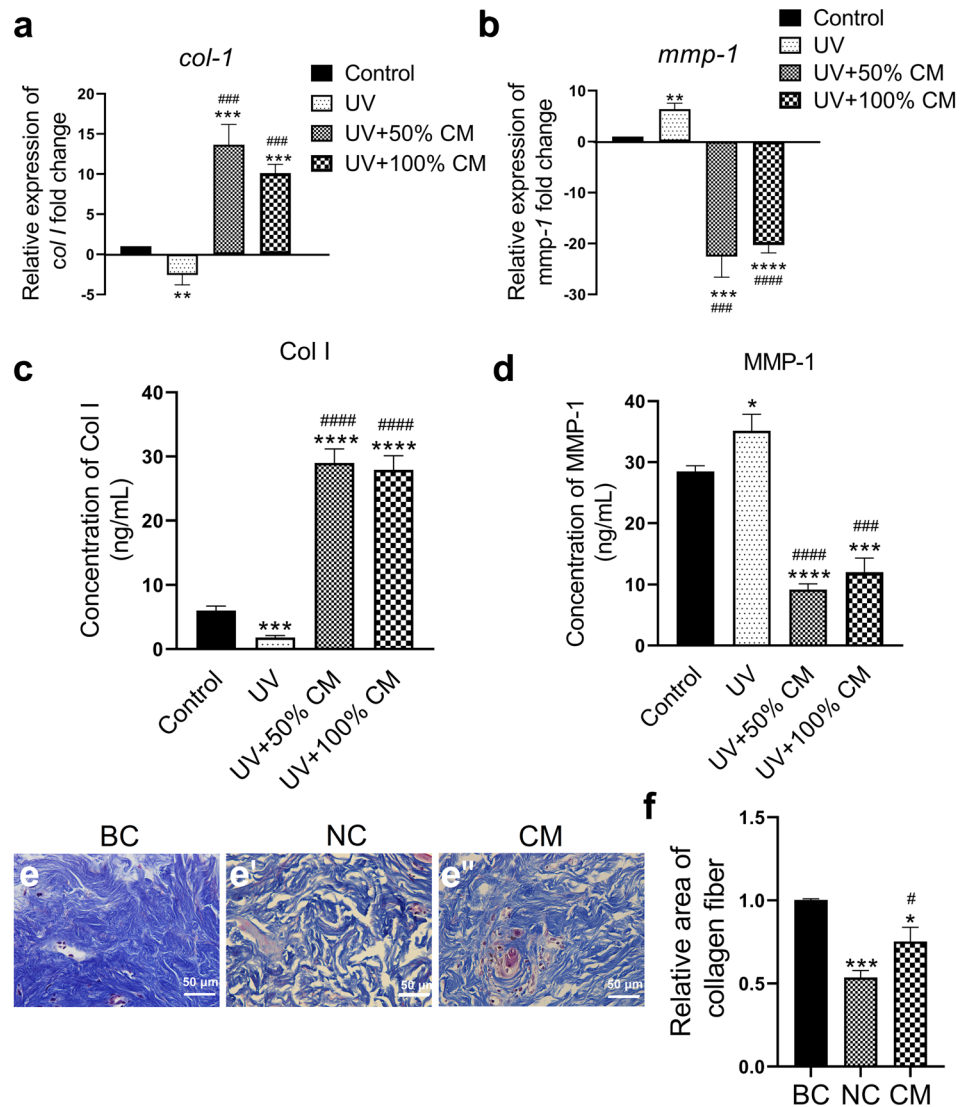
Defects in the development of the skin barrier occur in skin diseases, including AD and psoriasis (Elias et al. 2008; Hänel et al. 2013). As the main barrier in contact with the external environment, the intact skin barrier is crucial for the maintenance of tissue homeostasis (Akdis 2021). Inflammatory skin disease is associated with an imbalance in the internal environment of the skin, including biochemical defects, imbalance in immune responses, and abnormalities of the cytokine system (Farzanfar et al. 2018; Kim et al. 2020). Extrinsic damage, such as that caused by UVR, increases ROS formation; damages DNA, lipids, and proteins; destroys the extracellular matrix (ECM); and promotes cell apoptosis and cell aging, which accelerates inflammatory damage (Lephart 2016). Overall, skin homeostasis is a series of pathological events that occurs during crosstalk between cells and the microenvironment of the skin, and its imbalance initiates cell dysfunction, which leads to the occurrence and development of dermatological diseases. In this study, we addressed the therapeutic potential of the UC-CM for dermatological diseases using ex-vivo skin tissue and a 3D skin model to determine its ability to maintain skin homeostasis.

First, the secretome of UCMSC through the conditioned medium was obtained, and its safety was evaluated by performing cytotoxicity and skin irradiation tests with continuous applications for 14 days at the cell and animal

levels. Our results suggested that UC-CM could improve cell proliferation in HaCaT, HSF, and HUVEC cells under 50% concentration, as excessive concentration might change the composition of the basic culture medium when cultured. One hundred percent concentration of UC-CM application was found to be safe using New Zealand rabbits by conducting continuous multiple skin irritation experiments.

The SLS pre-treatment is a highly reproducible and effective procedure for barrier disruption (Döge et al. 2017). Additionally, SLS can cause the cells to release a large quantity of the inflammatory factor IL-1 α , a cytokine that is often used as an early marker for skin irritation (Gibbs et al. 2002; Müller-Decker et al. 1998; Welzel et al. 1996). We selected SLS irritation in EpiKutis[®] to mimic the skin environment with intensified inflammation and barrier disruption. The cultured tissues were evaluated by performing histomorphometry and determining tissue activity. We found that, although UC-CM did not improve tissue activity, the tissue morphology improved in the CM group. Basal cell vacuolization, which is the most prominent feature, was one of the primary morphological changes in interface dermatitis (Joshi 2013). After UC-CM treatment, a reducing vacuolar in living cells was found compared to the BC group. The stratum corneum (SC) in the epidermis acts in homeostatic control and blocks the formation of ROS (Rocha and Bagatin 2018). Exogenous

Fig. 6 Effects of UC-CM on ECM secretion. Quantitative RT-PCR for *col-1* (a) and *mmp-1* (b) and ELISA test for Col I (c) and MMP-1 (d). mRNA and protein levels of collagen I (Col I) and matrix metalloproteinase (MMP-1) with or without UC-CM treatment in UV-induced HSF cells were detected by RT-qPCR and ELISA. The relative transcriptional levels of target genes were quantified using the $2^{-\Delta\Delta C_t}$ method. The values shown are the means \pm SD from triplicate experiments. Image of Masson staining in skin tissue (e, e', e'') and relative statistical results (f). BC, black control; NC, SLS treatment; CM, SLS + UC-CM treatment; *vs corresponding control, $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$; #vs corresponding UV, $p < 0.05$, ### $p < 0.001$, #### $p < 0.0001$



stimulation promotes epidermal hyperplasia and mediates inflammation in human skin cells (Zhao et al. 2021). Compared to the SC of the BC group, the SC of the CM group was loose and thickened but was better than that of the NC group. To determine whether SC improvement occurred, we further examined the changes in two important proteins, including the filament-forming FLG and the structural protein LOR, which are responsible for the formation of a physically stronger barrier in SC.

Next, we examined the effect of UC-CM on the release of pro-inflammatory factors. Skin resident cells produce cytokines resulting in a complex network of signaling molecules that affect the quality of the skin barrier (Hänel et al. 2013). Based on the SLS stimulation of EpiKutis[®], UC-CM was found to mainly inhibit IL-1 α and TNF- α , but not PGE2 expression, in the SLS-induced 3D skin model in our study. The difference between the three cytokines is

that TNF- α and IL-1 α affect immune system-based inflammation processes, especially as an innate immune response, while PGE2 is a major factor in all types of skin inflammation (Ahuja et al. 2008; Gholijani et al. 2017; Agard et al. 2013). We speculated that UC-CM might promote skin immunity and affect the skin barrier through different regulatory pathways. This regulatory function makes IL-1 α and TNF- α have synergistic effects, which is different from the effects of PGE2. Previous studies have shown that IL-1 α and TNF- α have a strong synergistic effect on the expression of CXCL8 and b-defensin 2 (BD2) in psoriatic skin (Guilloteau et al. 2010). Some studies have also shown that PGE2 helps to mobilize skin dendritic cells (DCs) to drain lymph nodes (dLNs) in an IL-1-independent manner in skin infection (Krmeska et al. 2022). DC migration is necessary for initiating adaptive immunity, indicating that PGE2 also participates in adaptive immunity. Additionally, IL-1 α and TNF- α

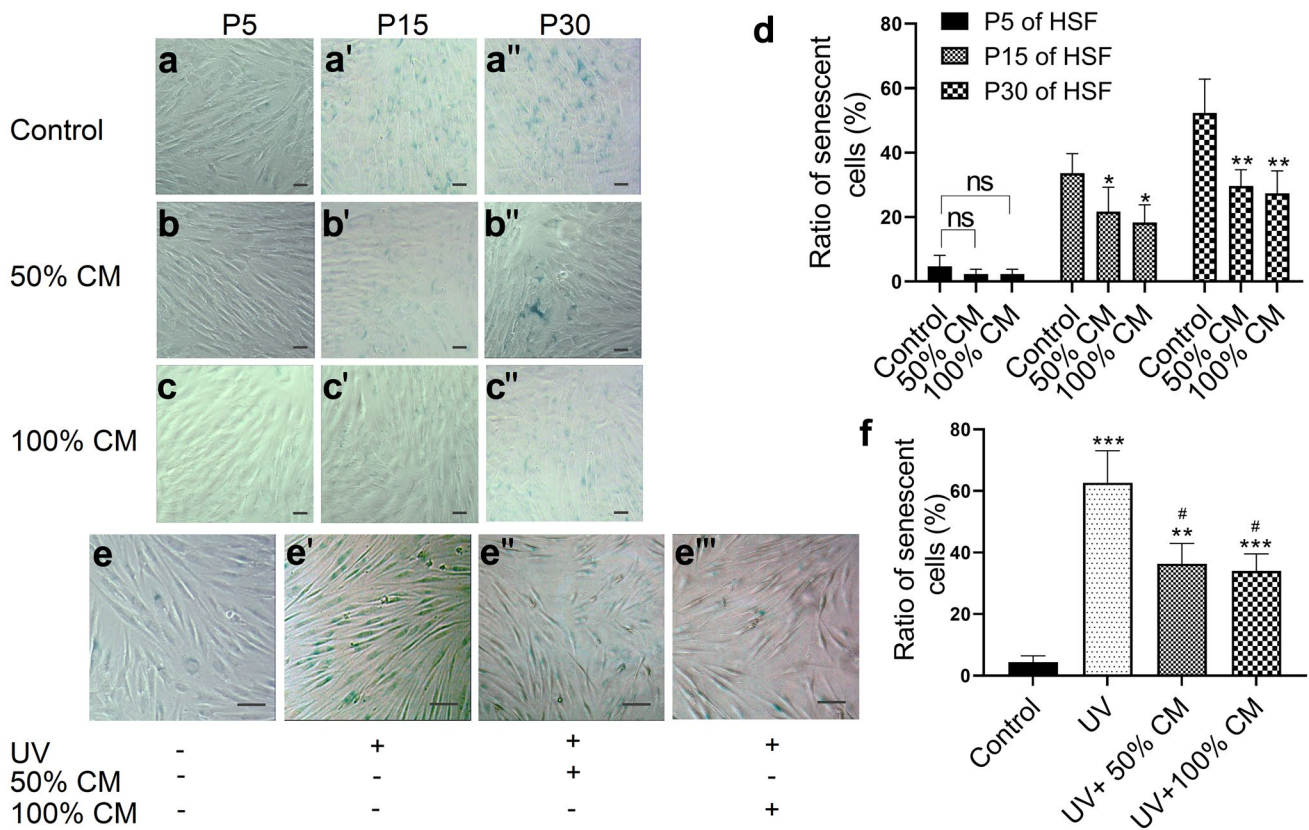


Fig. 7 Effects of UC-CM on cell function by reducing senescent cells. **a–c''** Morphology and SA-β-Gal expression of young (P5), intermediate (P15), and senescent (P30) HSF cells with or without UC-CM treatment, scale bar: 30 μm. **e, e', e'', e'''** Morphology and SA-β-Gal

are strong pro-inflammatory cytokines (Jin et al. 2021), and a decrease in their expression has positive therapeutic effects on recovering FLG involved in the formation of the skin barrier (Hänel et al. 2013). Specifically, TNF-α is expressed abnormally in psoriasis patients (Portugal-Cohen et al. 2012; Pasparakis 2012), and targeting it in psoriasis patients significantly reduces the disease symptoms (Pasparakis 2012). IL-1α leads to hyperkeratosis (O'Shaughnessy et al. 2010), and consistent with this concept, we found that an improvement in tissue morphology by thinner cuticles after UC-CM treatment might be due to a decrease in IL-1α levels (Fig. 2b''). The above-mentioned results support the therapeutic potential of UC-CM in the treatment of inflammatory skin diseases.

Disruption of the epidermal barrier is an important trigger in abnormal cutaneous inflammation (Zhang et al. 2020a). Pro-inflammatory cytokines can alter the expression of proteins involved in skin barrier function (Furue 2020); for example, TNF-α inhibits the expression of the FLG and LOR mRNAs in calcium differentiated keratinocytes (Kim et al. 2011). In many studies, FLG and LOR were found to be the main indicators of barrier function improvement

and had a synergistic effect. For example, *Elaeagnus* L gum polysaccharides (EAP) inhibited the production of several information mediators and increased FLG and LOR (Wang et al. 2021). The extracts of *Terminalia chebula* (TC), a deciduous tree, can increase the skin barrier establishment by upregulating the expression of the LOR and FLG mRNA and protein levels (Swindell et al. 2020). Our results suggested that UC-CM could increase the level of the barrier protein FLG without affecting the LOR expression, but both decreased in the NC group. LOR promotes the maturation of corneocytes and is an essential effector of cornification in the uppermost layer of stratum granulosum, (Ishitsuka and Roop 2021) while the primary function of FLG is to maintain corneocyte humidity. FLG participates in epidermal differentiation and contributes to the structural and functional integrity of the horny layer (Rocha and Bagatin 2018) and strongly influences the balance of the internal environment of the skin (Cabanillas and Novak 2016). During differentiation of the granular layer, FLG undergoes degradation that leads to the formation of the natural moisturizing factor (NMF), which is crucial for the maintenance of a healthy SC (Rocha and Bagatin 2018; Celleno 2018). On the other hand,

expression in HSF under UV irradiation after UC-CM treatment, scale bar: 50 μm. **d, f** Analysis of cellular senescence. *vs corresponding control, $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$; #vs UV, $p < 0.05$; ns, not significant

the degradation products of FLG facilitate skin hydration and an acidic environment, which are crucial for the optimal activity of enzymes involved in skin inflammation, lipid synthesis, and desquamation (Kim and Lim 2021). Abnormalities in the integrity of the skin barrier are mainly associated with reduced FLG production in AD (Scharschmidt et al. 2009; Kawasaki et al. 2012). We speculated that UC-CM might have medical value in the treatment of AD.

In the above-mentioned study, we found that UC-CM treatment can significantly reduce the expression of TNF- α in skin inflammation. While according to the report, TNF- α induces epidermal cell death and apoptosis (Zhao et al. 2021). Therefore, we hypothesized that UC-CM might enhance the defense function of the skin by improving the skin microenvironment. Subsequently, the therapeutic potential of UC-CM for dermatological diseases was evaluated based on cell functions, such as ECM secretion and cell detoxification. An increase in inflammatory factors was shown in UV-irradiated cell models, and FLG expression in HaCaT cells agreed with the 3D model. As reported, proteases, ROS, and pro-inflammatory cytokines induce tissue breakdown and impair physiological healing of the skin (Fisher 2005). New and old cells alternate to promote skin cell renewal via cell proliferation (Nguyen and Aragona 2021) and take rapid actions to recover cell functions, such as the reduction of reactive oxygen species (ROS) and removal of aging cells (Pignolo et al. 2020). In another study, we found that UC-CM can protect HaCaT and HSF cells from UV radiation by reducing ROS generation and cell apoptosis. On the other hand, we found activated SA- β -gal in both natural passage cultured old HSF cells and UVB-radiated young HSF cells, while UC-CM protection could significantly reduce the number of senescent cells. Our results suggested that UC-CM could effectively regulate and positively affect the restoration of skin homeostasis and enhance the anti-oxidation and anti-apoptosis ability of the skin to effectively resist endogenous damage.

UC-CM includes numerous cytokines (Ratajczak et al. 2012; Deng et al. 2018), which remodel the extracellular matrix in injured organs (Ratajczak et al. 2012). Collagen fiber is an important part of ECM. It is tough and plays an important role in maintaining skin tissue morphology and function (Shmulevich and Krizhanovsky 2021). The breakdown of collagen and the increase in MMP-1 are the key features of the pathology of various inflammatory skin diseases (Chung et al. 2001). According to our results, IL-1 α increased in the UV-induced cell models. IL-1 triggers the production of MMPs (Kalinski 2012), which can degrade collagen. Thus, we determined the expression of the Col-I and MMP-1 mRNA and protein levels. A positive regulation function of *col-1* and negative regulation of *mmp-1* was found

at the mRNA and protein level in our study, indicating the function of UC-CM in adjusting the components of ECM. Similar results were also obtained in the human adipose-derived mesenchymal stem cell-conditioned medium-treated UVB-induced human skin cells (Li et al. 2019).

A complex skin environment network affects the homeostasis, metabolism, and barrier function of the skin leading to the occurrence and development of dermatological diseases (Lephart 2016; Shmulevich and Krizhanovsky 2021; Ovadya et al. 2018). The most important aspect of our study was the examination of the role of UC-CM in regulating immune response, skin barrier, and cell function. Considering that this is the first study on the role of UC-CM in maintaining skin homeostasis, some doubts need to be addressed. For example, which aspect of UC-CM plays a key role in the above-mentioned functions. Cytokine balance is essential for efficient barrier formation, and the management of cytokine expression and activity is important for controlling and ameliorating the development of skin diseases (Vizoso et al. 2017). The balance between anti-inflammatory cytokines (such as tumor necrosis factor β 1 (Zagoura et al. 2012), IL-13, and neurotrophin-3 (Bermudez et al. 2016)) and pro-inflammatory cytokines (such as IL1 β (Zagoura et al. 2012), IL6 (Cantinieux et al. 2013; See et al. 2011), IL8 (Mirabella et al. 2011; Lee et al. 2011), and IL9 (Lee et al. 2011)) in UC-CM might determine the final effect. However, the UC-CM can provide exosomes (Exo). UC-Exo is produced and released by UCMSC and plays key immunomodulatory roles in the pathogenesis of various inflammatory skin diseases (Kim et al. 2019), such as psoriasis (Cheung et al. 2016; Jiang et al. 2019), AD (Kim et al. 2019), and systemic lupus erythematosus (SLE) (Kim et al. 2019). To summarize, the different regulatory abilities of UC-CM to perform immune regulation and mediate functional differences in barrier proteins need to be highlighted in greater detail.

In conclusion, our study highlighted the cosmetic or therapeutic applications of UC-CM on skin barrier dysfunction and abnormal skin homeostasis using ex vivo skin tissue, a 3D skin model, and human cells. We studied the biological effects of UC-CM on anti-inflammatory and skin barrier repair ability and also showed the importance of safety profiles. We found that cell function, which contributes to skin homeostasis, improved in UV-induced adult human skin cells after UC-CM treatment. We reported that the features of UC-CM in maintaining skin homeostasis might serve as an effective therapeutic strategy with a low risk of side effects. Thus, this therapeutic technique can overcome the limitations of inflammatory skin treatment in use.

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Declarations

Ethical approval and informed consent Human umbilical cords were harvested with written informed consent from the donor through the Yantai Yuhuangding Hospital with ethical approval ([2021]003). All procedures performed involving human participants in experiments were carried out per ethical standards of the institutional and/or national research committee and the 1964 Helsinki declaration and its later amendments or comparable ethical standards.

Conflict of interest The authors declare no competing interests.

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