A Novel Effect of Lipids Extracted from Vernix Caseosa on Regulation of Filaggrin Expression in Human Epidermal Keratinocytes

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Background: Vernix caseosa (VC), which is known as a unique human substance, is a biofilm that covers the skin of most human newborns. VC has many biological functions including anti-infective, skin cleansing and skin barrier repair. Objective: In the study, we purpose to investigate the novel effect of lipids extracted from VC on the regulation of filaggrin (FLG) expression and anti-inflammation in normal human epidermal keratinocyte (NHEK) cells. Methods: The lipids were extracted by chloroform/methanol (Folch method) and the major properties of fatty acid methyl esters were determined with gas chromatography-mass spectrometer. The relative viability of NHEK cells was evaluated by Cell Counting Kit 8 assay. The related expression of skin barrier protein was accessed with real-time quantitative polymerase chain reaction, Western blot and Immunofluorescence in NHEK cells with or without poly (I:C). Meanwhile, the changes of thymic stromal lymphopoietin (TSLP) and tumor necrosis factor alpha (TNF-α) are analyzed by enzyme-linked immunosorbent assay. Results: VC lipids mostly contained saturated and branched chains fatty acids. The expression of mRNA and protein of FLG were significantly increased after the supplement with lipid in NHEK cells. Meanwhile, lipids reversed the inhibition of poly (I:C) on FLG. Moreover, lipids suppressed the over secretion of TSLP and TNF-α induced by poly (I:C). Conclusion: These results indicate that lipids extracted from VC has positive effects on the expression of FLG and anti-inflammation, suggesting that lipids of VC may be used for a reference for novel therapeutic method in reducing and remedying skin disease like atopic disease. (Ann Dermatol 31(6) 611 ~ 620, 2019)

Keywords: Fatty acids, Inflammation, Keratinocytes, Vernix caseosa

INTRODUCTION

Atopic dermatitis (AD), which has approximately affected 25% of children and 3% of adults in the world, is a common, chronic and perennial skin inflammatory allergic disease. More seriously, the life quality of infant, children and teenagers with AD has been seriously destroyed. The main cause of AD contains genetic defects and exogenous factor stimulation. Genetic study has proved the significance of filament aggregation protein (filaggrin, FLG) gene in AD patient skin. The FLG, which assembles and aggregates keratin to form the keratin-matrix-complex that is an outer lipid envelope as an efficient barrier to prevention of water loss, mechanical damage, microorganism invasion, is a significant ingredient in stratum corneum (SC) which is important structure of the skin. Recent studies have showed that FLG null mutations play a key role in the pathogenesis of AD and FLG defect is aggravated by the imbalance of T helper 2 (Th2) cytokine in AD patients. Polyinosine-polyribocytidylid acid (poly I:C), a
common synthetic double-stranded RNA (dsRNA), can stimulate toll-like receptor-3 (TLR-3) to activate B cells and dendritic cells to secret pro-inflammatory cytokine that can bring out immune responses in AD patients6,7.

Vernix caseosa (VC), a white and cheesy biofilm, covers the SC of the skin of newborn when fetus go through the pregnancy stage of the last three months8. VC is a multi-functional skin cream, which participates in multiple physiological regulation. In utero, VC works as an effective waterproof film, protects the baby from harmful substance existed in amniotic fluid and takes part in the development of gastrointestinal and skin epidermis. After baby birth, VC can maintain low skin pH and significantly improves skin hydration. Bautista has found that adult forearm skin inside improved the capacity of moisture-binding after the treatment with VC10. Besides, VC-based creams are beneficial material to treat epidermal wounds, SC maturation and repair11.

The main ingredient of VC contains 80.5% water, 10.3% protein and 9.1% lipids in natural environment12. Lipids are critical matrix for the barrier of VC and nonpolar lipids are major component of the VC lipids. Sterol esters and triglycerides constitute main part of individual lipids, which are made up of straight long chain fatty acids and diverse methyl branched fatty acids. VC as a natural topical agent has similar structure with SC and the pioneer information of VC has been obtained. The lipids composition of SC mainly contain free fatty acids, which compared with VC is interesting. It has been found that a lot of fatty acids that were from phospholipids can regulates the levels of SC acidification and improve SC integrity14 and their 1-monoacylglycerols have antibacterial activity in vitro15. Boiten et al.16 and Tollin et al.17 applied a ceramide mixture based on the ingredient of the VC to facilitate SC barrier development. The anti-inflammatory effect of some fatty acids has been founded in many situations18,19. However, the effects of lipids extracted from VC on the normal human epidermal keratinocyte (NHEK) cells, especially on the structural protein FLG, had not been investigated. The objective of this study is to explore fatty acids composition of VC lipids and access its novel effects on the FLG and anti-inflammation in vitro.

MATERIALS AND METHODS

Sample collecting

VC was immediately gathered from six male and four female newborn subjects who we re provided at full term condition (weeks of gestation reached 40–42 weeks). VC samples were put in glass sample bottles and stored under −80°C condition. Blood-polluted samples were abandoned. The study was approved by the Ethics Committees of the Shanghai Clinical Research Center (IRB no. C181108) and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards. This article does not contain any studies with animals performed by any of the authors. We received the patient’s consent form about publishing all photographic materials.

Lipids extraction

Lipids extracted from vernix were obtained using the Folch method that is previously described20. Briefly, 1 g clean VC was added 20 ml of chloroform/methanol mixture (ratio of mixture: 2/1, V/V). After vortex for 15–20 minutes and centrifugation at 1,500 rpm for 1 min at room temperature, the liquids phase is collected. The solvent is washed with 0.2 volume of water. After centrifugation at 2,000 rpm for 10 min and discarding the supernatant, methanol and water were added in proportion of 1:1 volume ratio to the liquid phase. Finally, the lower level chloroform phase was dries with N₂ and samples were stored in −80°C condition until using.

Cell culture

NHEK cells were purchased from Guangdong Biocell Co., Ltd. (Guangdong, China) and cultured in Epilife medium (Gibco; Thermo Fisher Scientific, Waltham, MA, USA) with 60 μM Ca²⁺ and human keratinocyte growth supplement (HKGS; Gibco) in a humidified at condition of 95% air and 5% CO₂ at 37°C. The Epilife medium was changed every two days until the cells were used for the next experiment at 75% to 85% confluence. The cells were grown for no more than 8 cell generations.

Cell viability assay

The NHEK cells were seeded into 96-well plates (Nunc; Thermo Fisher Scientific) at a density 4×10³ cells/well. Then cells were incubated in serum-free culture medium for 12 hours. Cells were exposed to lipids and poly (I:C) (0–400 μg/ml) (Sigma-Aldrich, St. Louis, MO, USA) for 24 hours. After adding Cell Counting Kit 8 (CCK-8; Beyotime Biotechnology, Shanghai, China) 10 μl/100 μl, cell viability was accessed by reading absorbance at 450 nm (ID5 Plate Reader; Molecular Device, San Jose, CA, USA). According the results of the cells viability (%), suitable poly (I:C) and lipids concentrations were chosen. The calculation formula of viability is as follows:

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\text{Viability} (\%) = \frac{OD_{\text{treatment group}} - OD_{\text{blank background}}}{OD_{\text{control group}}} \times 100\%
\]

Lipids supplement and inflammatory stimulation

NHEK cells (2×10⁴ cells/ml) were seeded into 6-well
Lipids of Vernix Caseosa Regulate FLG Expression

Table 1. Primer designed for real-time quantitative PCR

| Gene name | Forward | Revise |
|-----------|---------|--------|
| FLG       | GCAAGGTCAAGTCCAGGAGA | CCCTCGGTTTCCACTGTCTC |
| TSLP      | ACTCTCGACTTGTGTCCTCCC | TGGCGAACATTTCTTTGGCG |
| TNF-α     | TATCCTGGGGGACCCCAATGT | AGCTTCTCCCACCGCAAAG |
| GAPDH     | CGGAGTCAAGCGATTTGGCTGTAT | AGCCCTTCCATGGTGTAAGAC |

PCR: polymerase chain reaction, FLG: filaggrin, TSLP: thymic stromal lymphopoietin, TNF-α: tumor necrosis factor alpha, GAPDH: glyceraldehyde 3-phosphate dehydrogenase.

Real-time quantitative polymerase chain reaction analysis

The treated cells were lysed with the TRIZOL reagent (Invitrogen Technologies, Carlsbad, CA, USA) following the manufacturer’s recommended protocol. The content and quality of total RNA were measured by Qubit 3.0 (Thermo Fisher Scientific). This target RNA fragment was chosen as a basic template for subsequent cDNA synthesis using the Prime script RT reagent Kit (Thermo Fisher Scientific). To measure related mRNA levels, SYBR Premix Ex Taq II (Takara Bio Inc., Dalian, China) were performed using the Light Cycler 96 system (Roche, Basel, Switzerland) following the manufacturer’s recommended protocol and primers were listed in Table 1. The data were analyzed using the delta cycle threshold method and $2^{-\Delta \Delta CT}$ method, and the expression levels of each gene were normalized to glyceraldehyde 3-phosphate dehydrogenase (GAPDH). All values are reported as means±standard deviation (SD).

Western blotting

The NHEK cells were risen with pre-cold phosphate-buffered saline (pH 7.4) to inhibit protease activity. Total soluble protein extracted using M-PER Mammalian Protein Extraction Reagent (Thermo Fisher Scientific) with a protease inhibitor mixture (1:100; Roche Applied Science, Basel, Switzerland). The protein concentration was measured by a bicinchoninic acid (Pierce; Thermo Fisher Scientific) kit according to the instructions. The 12% sodium dodecyl sulphate-polyacrylamide gel electrophoresis was used to separated total proteins (20 μg), then gel was transferred onto a polyvinylidene fluoride (PVDF) membrane using iBlot 2 system (Thermo Fisher Scientific). The PVDF membrane was blocked with 5% bovine serum albumin (BSA) for 1 hour. After blocking, the PVDF membrane was incubated in 0.1% Tris buffered saline with Tween 20 including primary antibody anti-FLG (1:2,000; Thermo Fisher Scientific), anti-GAPDH (1:3000; Invitrogen Technologies) overnight at 4°C. Next, the membrane was washed and incubated with horseradish peroxidase-conjugated secondary antibody (1:15,000; Abcam, Cambridge, UK) for 1 hour. Protein expression was detected using iBright L1000 image system (Thermo Fisher Scientific). The results of protein expression were quantified with densitometry normalized to GAPDH by iBright analysis software (Thermo Fisher Scientific).

Immunofluorescence assay

After treatment, the treated cells were added 4% paraformaldehyde to fix cells for 30 minutes at room temperature. After fixation, cells were incubated in 0.05% Triton X-100 for 30 minutes to permeate cell. Then, the cells were blocked with 5% BSA buffer and incubated with primary FLG antibody (1:500; Abcam) overnight at 4°C. The cells were treated with an AlexaFlour 488-conjugated secondary Goat anti-rabbit IgG (1:100; Invitrogen Technologies) for 2 hours in the dark conditions. The nuclei were dyed by incubation with 4',6-diamidino-2-phenylindole (1:1,000 dilution; Thermo Fisher Scientific) for 20 minutes at room temperature. The results of cells were observed with EVOS FL autoII fluorescence microscope (Life Technology, Boston, MA, USA) and analyzed using the EVOS browser imaging software.

Measurement of secreted thymic stromal lymphopoietin and interleukin-1 α by enzyme-linked immunosorbent assay

After poly (I:C)-stimulated NHEK cells were treated with different concentrations of lipid for 24 hours, the samples were collected from the cell supernatants. Tumor necrosis factor alpha (TNF-α), thymic stromal lymphopoietin (TSLP) were determined by enzyme-linked immunosorbent assay
Gas chromatography-mass spectrometer of fatty acid methyl esters

The lipids extracted from VC were converted into fatty acid methyl esters by incubation in 10% vitriol (H₂SO₄)/methanol for 3 hours at 70°C. After adding 2 ml n-hexane (Sigma-Aldrich), 1 ml of ultrapure water was added to aid stratification, and the supernatant was aspirated for detection. The methyl esters were analyzed by gas chromatography-mass spectrometer (GC-MS; Agilent 7890B; Agilent Technologies, CA, USA) containing a 5977 MSD connected with a HP-5MS (5% phenyl, polymethyl siloxane column) (30 m×250 mm×0.25 μm). A 1 μl specimen was injected into the GC inlet with a syringe. The split ratio was set as 1 : 20 and the temperature of inlet was kept at 250°C. Initial column temperature was 100°C, rising at a rate of 2°C/min, holding for 2 minutes, then rising at a rate of 2°C/min, 220°C holding 3 minutes, at last rising to 290°C at a rate of 5°C/min, maintaining for 2 minutes. The carrier gas, helium, utilized at a flow rate of 1.0 ml/min. Mass charge ratio scanning was performed from 50 to 500 m/z with electron ionization (70 eV). The area of each peak under the curve was automatic integration and the relative percentage was obtained compared with integrated using ChemStation software. Database similarities (%) higher than 85 was considered acceptably for correct identification.

Statistical analyses

All the values have been reported in terms of means±SD. The mean values were calculated based on data from at least three independent replicate experiments. The data were analyzed using the Student’s t-test. p-value less than 0.05 was considered statistically significant. All statistical analyses were performed using SPSS ver. 22.0 (IBM Corp., Armonk, NY, USA).

RESULTS

Cytotoxicity of lipids and poly (I:C) in NHEK cells

The CCK-8 assay was used to evaluate the safety of lipids and poly (I:C). As is shown in Fig. 1, NHEK cells were exposed to different concentration of lipids extracted from VC and poly (I:C; 400, 200, 100, 50, 25, 12.5, 6.25, 0 μg/ml) and cell vitality was decreased in a dose-dependent manner. Cells cytotoxicity induced by lipids was significantly increased at concentrations greater than 100 μg/ml. So, we used 100 μg/ml, 50 μg/ml, 25 μg/ml as the safe concentration of lipids. Poly (I:C), as an inflammatory stimulator, a dose-dependent decreased cell vitality also has been found (Fig. 1B). When NHEK cells were exposed to 25 μg/ml poly (I:C), the cell vitality was significantly decreased. When the concentration of poly (I:C) was greater than 50 μg/ml, the cell vitality kept a low level. Thus, we choose 10 μg/ml poly (I:C) as a suitable concentration for further experiments.

Fig. 1. The effect of lipids extracted from vernix caseosa (VC) and Poly (I:C) on normal human epidermal keratinocyte (NHEK) cells viability. (A) NHEK cells were exposed on lipids at the different concentrations (400, 200, 100, 50, 25, 12.5, 6.25, 0 μg/ml) for 24 hours. (B) NHEK cells were exposed on poly (I:C) at the different concentrations (400, 200, 100, 50, 25, 12.5, 6.25, 0 μg/ml) for 24 hours. Cell viability was assessed by Cell Counting Kit 8 assay. The results are representative of five independent experiments. Bars means standard deviations. *p<0.05.
The effect of lipids on the expression of FLG gene and protein expression

Real-time quantitative polymerase chain reaction (RT-qPCR) and western blot were used to measure the changes of FLG protein expression induced by lipids with or without poly (I:C) in vitro. In the NHEK cells, stimulation with lipids at 25, 50, 100 μg/ml for 24 hours increased the expression of FLG, which were less than positive control (15 μM WY14643). Fig. 2A presented that the 50 μg/ml lipids group significantly (p < 0.01) increased the FLG mRNA level compared with poly (I:C)-treated control group. Meanwhile, the protein expression of FLG was significantly increased than normal control group (Fig. 2B). In Fig. 2A, there is a clear decreasing trend of gene expression of FLG in poly (I:C)-treated group compared with poly (I:C)-untreated. The addition of lipids with 25, 50, 100 μg/ml effectively reversed inhibition of poly (I:C) on FLG gene expression (Fig. 2A). Fig. 2C showed that 50 μg/ml lipids significantly improved FLG expression compared with normal control and poly (I:C)-treated group. The result of immunofluorescence 50 μg/ml lipids significantly (p < 0.01) increased the FLG protein expression under normal condition in vivo. Meanwhile, the addition of poly (I:C) significantly decreased FLG expression (p < 0.05). Further, the FLG protein expression significantly improved after the treatment of 50 μg/ml lipids for 24 hours (Fig. 3).

Lipid suppress the secretion of TNF-α and TSLP in NHEK

NHEK cells were treated with 10 μg/ml poly (I:C) to induce the inflammatory response and simultaneously exposed to lipids at various concentrations. TNF-α and TSLP are important in the development of inflammatory response, which were measured by RT-qPCR (Fig. 4A) and ELISA (Fig. 4B) in NHEK cells. The levels of TNF-α and TSLP was significantly increased when NHEK cells were exposed to 10 μg/ml poly (I:C) for 24 hours (Fig. 4). But treatment with 25 μg/ml, 50 μg/ml, and 100 μg/ml lipids significantly decreased the mRNA level of TNF-α and TSLP (Fig. 4A). Further, the results of ELISA illustrated that the protein level of TNF-α and TSLP significantly reduced after the treatment of 50 μg/ml, 100 μg/ml lipids com-

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**Fig. 2.** The positive effect of lipids on filaggrin (FLG) mRNA and protein in normal human epidermal keratinocyte cells. (A) The levels of FLG mRNA increased after treatment with positive control (WY14643) and different lipids under poly (I:C) untreated and poly (I:C) treated condition; real-time quantitative polymerase chain reaction. (B) The FLG protein expression increased after treatment with positive control (WY14643) and lipids; Western blot. (C) Quantitative processing of FLG protein expression. Data are presented in graphical form that shows the fold change compared with control group and relative density. The values are the mean ± standard deviation of at least three independent experiments in duplicate. GAPDH: glyceraldehyde 3-phosphate dehydrogenase, NC: normal control. *p < 0.05 compared with normal control, **p < 0.01, #p < 0.05 compared with poly (I:C)-treated group.
Fig. 3. Lipids improved the expression of filaggrin (FLG) with or without poly (I:C) in normal human epidermal keratinocytes; immunofluorescence ×200. The data are expressed as relative density in graphical form (FLG/glyceraldehyde 3-phosphate dehydrogenase [GADPH]). Bars mean standard deviations. DAPI: 4',6-diamidino-2-phenylindole. *p<0.05 compared with normal control, **p<0.05 compared with poly (I:C)-treated group, ***p<0.01.

pared with poly (I:C)-treated group (p<0.01) (Fig. 4B). These results suggest that lipids extracted from VC could negatively control the inflammatory responses by suppressing TNF-α and TSLP.

The composition of VC fatty acids
As shown in Table 2, the composition of fatty acid methyl esters (FAME) was analysis by GC-MS. The high contribution originated from 14:0, 16:0, 16:1, 18:0, and 18:1. These contributed more than 10% to the total fatty acids in our samples. The major methylated fatty acids constituents of lipid were methyl tetradecanoate (49.83%), Tetradecanoic acid (8.2%), hexadecanoic acid, methyl ester (7.29%), Methyl stearate (2.27%), etc. Other minor methylated fatty acid compounds were detected in the vernix extract including tridecanoic acid, methyl ester (0.11%), pentadecanoic acid, methyl ester (0.23%), 9-hexadecenoic acid, methyl ester (0.21%), 9,12-octadecadienoic acid methyl ester (0.26%), methyl 9-(Z)-octadecenoate (0.95%). Some other fat-soluble alkanes compounds were also determined. However, some branched methylated fatty acids were detected included tridecanoic acid, 12-methyl-, methyl ester (0.92%), tetradecanoic acid,12-methyl-, methyl ester (1.31%), methyl(Z)-hexadec-9-enoate (0.28%), methyl -hexadecanoic acid-methyl ester (0.24%).

DISCUSSION
The proFLG presented in stratum granulosum is degraded to generate FLG that is a critical role for the integrity of the SC and calcium is involved in the procession of proFLG degradation during terminal epidermal differentiation. Finally, FLG is degraded to produce its partial component amino acids, a mixture of hygroscopic amino acids, called natural moisturizing factor that is contributed to SC acidification and hydration. It has been founded that the skin presents the degressive expression of FLG in patients with AD. So, FLG deficiency has a key role in pathogenesis of AD, which improves the risk of microbial infection and has an impact on the integrity of the epidermis. There are many novel therapeutic approaches to treat AD or other atopic diseases by adjusting FLG expression.

VC is a noteworthy biofilm that covers the skin of most human fetus and newborns. Because the increasing impacts in water-holding capacity, skin inflammation and immunity, VC that contains abundant lipids have been studied in many fields. High internal phase water-in-oil emulsions designed as VC analogues have better semi-permeability and water-handling properties than conventional lipids, which is closed to native vernix. Vernix caseosa mimetic, another vernix analogue, can improve the skin
barrier repair and increase the capacity of SC moisturization\textsuperscript{25}. Further, some studies indicated that there are some differences of lipid compositions in lesion and AD skin compared with health skin\textsuperscript{26}, which is related to levels and chain length of fatty acids. Besides, the changes of lipid composition and organization had been investigated in human skin SC after topical treatment with free acids\textsuperscript{27}. Previous studies have showed that palmitoleic acid (C16:1) and linoleic acid (C18:2) have powerful antimicrobial and antiviral activity and unsaturated fatty acids can upgrade the expression of FLG\textsuperscript{28}. In our study, FAME species mainly contains saturated, monounsaturated (C14:0, C16:0 and C16:1, C18:1) and branched fatty acids (Table 2), which is well line with recent report\textsuperscript{29}. But, there is no study to investigate whether lipids extracted from VC have an impact on FLG in vitro. We first quantified effects in mRNA and protein expression of FLG in NHEK cells treated with different concentrations of lipids (Fig. 2). Our study showed that the lipids extracted from VC can increase the expression of gene and protein expression of FLG in NHEK cells.

Epidermal keratinocytes play a central role in immune reaction and inflammatory\textsuperscript{30}, in which dsRNA brings out a series of responses. TSLP, an interleukin (IL)-17-like cytokine, is overexpressed by keratinocytes in lesion skin with AD\textsuperscript{31}, which has a downregulatory effect on FLG by signal transducer and activator of transcription 3 and extracellular signal-regulated kinase-depend pathways\textsuperscript{32}. Howell et al.\textsuperscript{33} has reported that Th2 cytokines can improve viral replication by downregulation of the secretion of LL-3 in the lesion skin with AD, which indicated that Th2 and poly I:C synergistically stimulate the release of TSLP. Rouaud-Tinguely et al.\textsuperscript{34} had demonstrated that IL-4 and IL-13 can down-regulate the expression of FLG and
Table 2. Fatty acid methyl esters

| No. | Time | Molecular Name | Area (%) |
|-----|------|----------------|----------|
| 1   | 2.76 | C_{10}H_{22} Decane | 0.50     |
| 2   | 3.04 | C_{12}H_{26} Dodecane | 2.44    |
| 3   | 6.05 | C_{20}H_{42} Eicosane | 1.96    |
| 4   | 7.00 | C_{20}H_{44} Heneicosane | 2.37   |
| 5   | 7.46 | C_{28}H_{58} Octacosane | 0.43   |
| 6   | 9.11 | C_{14}H_{28}O_{2} Tridecanoic acid, methyl ester | 0.11   |
| 7   | 9.42 | C_{13}H_{30}O Benzophenone | 0.15   |
| 8   | 10.61 | C_{16}H_{34}O 2-hexyl-1-Decanol | 0.12 |
| 9   | 10.99 | C_{15}H_{30}O Tridecanoic acid, 12-methyl-, methyl ester | 0.92 |
| 10  | 12.27 | C_{15}H_{30}O Methyl tetradecanoate | 49.83   |
| 11  | 13.53 | C_{15}H_{30}O Tetradecanoic acid | 8.20 |
| 12  | 14.39 | C_{16}H_{32}O Tetradecanoic acid,12-methyl-methyl ester | 1.31 |
| 13  | 15.76 | C_{17}H_{32}O Pentadecanoic acid, methyl ester | 0.23 |
| 14  | 17.22 | C_{17}H_{32}O 9-Hexadecenoic acid, methyl ester | 0.21 |
| 15  | 18.20 | C_{17}H_{32}O Hexadecanoic acid, methyl ester | 0.91 |
| 16  | 18.79 | C_{17}H_{32}O Methyl(Z)-hexadec-9-enoate | 0.28 |
| 17  | 19.68 | C_{17}H_{32}O Hexadecanoic acid, methyl ester | 7.29 |
| 18  | 21.07 | C_{19}H_{32}O n-Hexadecanoic acid | 0.38 |
| 19  | 22.81 | C_{19}H_{36}O methyl -Hexadecanoic acid-methyl ester | 0.24 |
| 20  | 27.67 | C_{19}H_{36}O 9,12-Octadecadienoic acid methyl ester | 0.26 |
| 21  | 28.05 | C_{19}H_{36}O Methyl 9-(Z)-octadecenoate | 0.95 |
| 22  | 29.64 | C_{19}H_{36}O Methyl stearate | 2.27 |
| 23  | 37.46 | C_{27}H_{46}O Cholesterol | 0.26 |

the over-expression of the Th2 was showed up in acute AD skin. The activation of TSLP will have an negative impact on innate immunity and trigger a nuclear transcription factor (κ B)35. In this study, poly I:C mimicked viral dsRNA to stimulate the massive expression of TSLP (Fig. 4). Meanwhile, our study demonstrated that lipids significantly inhibits the production of TSLP induced by poly (I:C) in NHEK cells. Interestingly, previous studies observed that self-dsRNA from damaged skin can trigger the production of TSLP by TLR-3 pathway, which has been founded as a potential mechanism for inhibition of inflammation36. We speculate that lipids extracted from VC can interfere the TLR-3 pathway to decrease the production of TSLP responses induced by dsRNA. We also demonstrated that lipid significantly decrease the impact of poly (I:C) on FLG genes in vivo (Fig. 3). Additionally, poly (I:C) also stimulates normal NHEKs to release pro-inflammatory cytokines that will in turn stimulate other related cells to produce various adhesion molecules and cytokines such as IL-1 and IL-8, which can accrue inflammatory cells to lead inflammatory reaction. Our study showed that TNF-α significantly increased induced by 10 μg/ml poly (I:C). In recent study, TNF-α alone or combined with Th2 cytokines can affect barrier lipid properties and cause the reduction in the distribution of free fatty acids37. In our study, the mRNA and protein levels of TNF-α significantly decreased after the treatment of different concentration of lipids.

There are some limitations in this study. First, which components of lipids increase the mRNA and protein expression of the FLG has not been investigated. Furthermore, the mechanism for inducing the inhibition of poly (I:C) on FLG is not known. So, additional experiments are going on to handle these limitations.

In conclusion, our findings indicate that VC lipids have some beneficial effects on improving skin barrier function through increasing expression of FLG and regulating the NHEK cells inflammation by inhibiting the secretion of TSLP and TNF-α. Therefore, lipids extracted from VC may be used for a reference for novel therapeutic method in reducing and remedying skin disease like atopic disease.

CONFLICTS OF INTEREST

The authors have nothing to disclose.

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