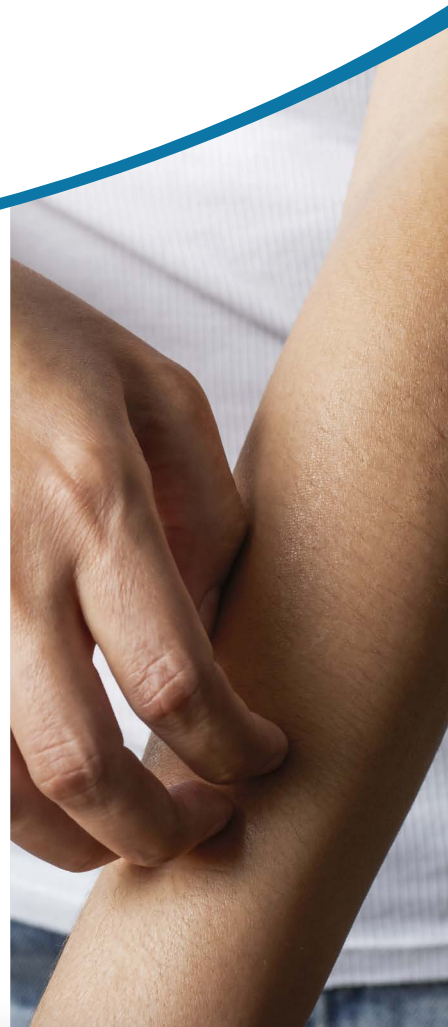


UPDATES ON DERMATOLOGY

SKIN BARRIER AND ITS ECOSYSTEM





Stéphane FAUVERGHE
NAOS International Medical Director

Dear All,

I am very pleased to present you the sixth edition of BIODERMA Updates Series dedicated to updates in Dermatology.

For 3 years now, BIODERMA has been regularly organizing international events dedicated to Dermatology, for dermatologists and all healthcare professionals interested in Dermatology, always presented by renowned experts in their field.

*In our approach to promote the development of knowledge in Dermatology, we have the pleasure to propose you this new publication, that is the summary of the BIODERMA Symposium held during the World Congress of Dermatology in Singapore in July 2023: **Ecobiology: Dialogue between skin barrier and its ecosystem** with Enzo Berardesca from the USA, Kenji Kabashima from Japan, Brigitte Dréno from France and myself as speakers.*

During this symposium, Enzo Berardesca presented: Skin barrier, an organ still in adaptation. Kenji Kabashima delivered a lecture about the environmental triggers and the skin barrier. The lecture of Brigitte Dréno, our chair, was about the skin microbiome, a new actor in cutaneous neurogenic inflammation. And, finally, I presented: Atopy, an ecobiological approach for a targeted skin care formula ecobiological approach of sun protection: to reinforce the natural mechanisms of the skin.

I wish you all an enjoyable, enriching and interesting reading.

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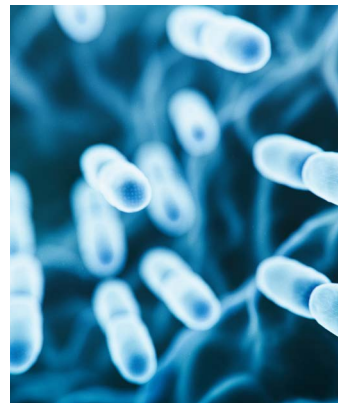


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SPEAKERS'S SHORT BIOGRAPHIES



Enzo BERARDESCA
USA

Enzo BERARDESCA is director of the dermatology department of inflammation and immuno-infectiousness at the Institut de dermatologie de San Gallicano, IRCCS, Rome. He has published more than 400 scientific books and 8 books. From a scientific point of view, he has been involved for many years on the integument, using non-invasive methods to conduct efficacy and safety studies on topical products. He is scientific director of Dermakos magazine.

Enzo Berardesca, obtained his training at the University of Pavia and received the M.D. degree in 1979. He served as resident and dermatologist in the Dept of Dermatology, IRCCS Policlinico S. Matteo, Pavia from 1982 to 1987, as research assistant in the Dept. of Dermatology, University of California School of Medicine in San Francisco, USA in 1987.

From 1988 to 2001 he has been at the Dept. of Dermatology of the University of Pavia,

head of the Dermatoallergology Unit and of the Skin Bioengineering Lab.

He is member of the editorial board of Skin Pharmacology, Skin Research and Technology, The American Journal of Clinical Dermatology and the Journal of Cutaneous and Ocular Toxicology. He is member of the Society for Investigative Dermatology, the European Society for Dermatological Research, the Italian Group for Research on Contact Dermatitis (GIRDCA), and vice-chairman of the European Group For standardization of Efficacy Measurements of Cosmetics (EEMCO group).

His current major research interests are irritant dermatitis, barrier function and noninvasive techniques to investigate skin physiology with particular regard to skin color and racial differences in skin function, sensitive skin and efficacy evaluation of topical products.



Kenji Kabashima
JAPAN

Kenji Kabashima, MD, PhD, graduated from Kyoto University in 1996. He further honed his medical and dermatology skills through training at the US Naval Hospital in Yokosuka, Kyoto University Hospital, both in Japan, and at the University of Washington Medical Center in the US.

He initiated his research on bioactive lipid mediators at Kyoto University and received his PhD there. Subsequently, he pursued his studies at University of California San Francisco (UCSF) and the University of Occupational and Environmental Health in Japan.

Currently, Dr. Kabashima holds several prominent positions, including Professor and Chair of the Department of Dermatology at Kyoto University Graduate School of Medicine, Senior Principal Investigator at A*STAR (Singapore), and Visiting Consultant at the National Skin Centre (Singapore). He also serves as President of the Japanese Society for Investigative Dermatology.

His research focuses on understanding the underlying mechanisms of inflammatory skin diseases such as atopic dermatitis, contact dermatitis, and psoriasis, as well as exploring 3D visualization of the skin using two-photon microscopy and drug development.



Brigitte DRÉNO

France

Prof. Brigitte DRÉNO is a Dermato-Oncologist. She is also Vice President of the Scientific and technical Culture of Nantes University France. She leads a research project included in the investments for the future hospital-university research in health (RHU) focusing on burn and regenerative dressing. She works also the field of melanoma, drug-induced cutaneous adverse events and microbiome and acne. She is the director of an INSERM INCITE research team.

She is Founding member of the European Association of Dermato Oncology (EADO), past president of the French Society of Dermatology and of the French college of dermatology teachers, Treasurer of the International League of Dermatological Societies (ILDS). In addition, member of AAD, ADA, International Society for Cutaneous Lymphomas, Skin Cancer Foundation.

She has published over 900 articles referenced in PubMed (H Index 83), participated to the redaction of chapters of several books. She built the first GMP cell and gene therapy hospital unit in France in Nantes.

She is an Editorial Board Member of JEADV, Acta Dermatology, European Journal of Cancer Prevention, International Journal of women's Dermatology and Editor of the quarterly medical review (La Presse Médicale). She has obtained several awards such as the Award of ILDS, the Prize of the International Society for Cutaneous Lymphomas, the Prize of the French victories of medicine. She has been granted to the grade of Knight of the Legion of Honor by the President of the French Republic, as well as Chevalier in the Order of Academic Palms of Nantes' s University and has been elected recently member of the French Academy of Medicine.

THE SKIN BARRIER, AN ORGAN STILL UNDERGOING ADAPTATION

ENZO BERARDESCA,

Philip Frost Dept. of Dermatology University of Miami, Miami, USA

The human skin forms a protective barrier against the external environment and is our first line of defense against physical assaults (mechanical injury, UV-irradiation), microbial assaults (bacteria, fungus, virus), and chemical assaults (irritants, allergens). It also defines our outward appearance, protects our internal tissues and organs, and acts as a sensory interface. It has important homeostatic functions such as reducing water loss and contributing to thermoregulation of the body. Skin barrier disruption can lead to increased transepidermal water loss (TEWL) and dryness.



The skin barrier is a living system adapting and evolving itself throughout our life. At birth, the skin barrier is not fully complete, and the beginning of postnatal life is a period for active adaptation and maturation of the cutaneous structure and functions⁽¹⁾. In infancy (first year after birth), skin structures and functions are similar to those of adults (excluding the inactive apocrine sweat glands).

Androgens may influence the barrier function, as testosterone perturbs epidermal permeability barrier homeostasis⁽²⁾. Development of the skin barrier before birth is increased by estrogens and decreased by testosterone. In experimental models, the inhibition of androgen production stimulated barrier recovery⁽²⁾. While lower TEWL and increased skin reactivity were observed in some women as a function of the menstrual phase⁽³⁻⁵⁾, no significant gender-differences in TEWL, hydration, pH or sebum were detected in other studies^(6, 7). Skin repair is improved by estrogens; but repairing capabilities decrease after menopause⁽⁸⁾. Increased pain sensitivity to noxious stimuli, increased chronic pain, and lower analgesic effects of drugs have been observed in women, as a function of the menstrual cycle⁽⁹⁾.



With aging, and particularly in women after menopause, the skin becomes dryer and less flexible, with increased TEWL, fragmentation of collagen and elastin, and with fewer but larger corneocytes⁽¹⁰⁻¹²⁾. As a result, skin barrier structure, permeability barrier function, epidermal calcium gradient, epidermal lipid synthesis and stratum corneum (SC) lipid processing, cytokine production and response after aggression, SC acidity, SC hydration, and antimicrobial barrier are changed or disturbed⁽¹³⁾. Hydration becomes uneven in aged skin, and ethnic differences in skin dryness were found between Caucasian, African, American, Chinese, and Mexican women, with a higher percentage increase in Caucasian women^(14, 15).

Epidermal dysfunction, compromised permeability homeostasis, reduced stratum corneum hydration and elevated skin surface pH predispose to the development of aging-associated cutaneous and extracutaneous disorders, including eczematous dermatitis, pruritus, and xerosis⁽¹⁶⁾. Alterations in epidermal function can lead to the development of a chronic, low-grade systemic inflammation termed “inflammaging”,

which is linked to the development of aging-associated systemic disorders⁽¹⁶⁾.

As the epidermis forms a microbial, physical, chemical, immunological, and neuro-sensory barrier between the internal and external environment, it is important to consider different barriers rather than just a single physical barrier⁽¹⁷⁾.

The microbial barrier, or microbiome, is made of skin microorganisms which form the first barrier against the environment through various mechanisms of colonization resistance, including resource exclusion, direct inhibition, and/or interference⁽¹⁸⁾. The skin microbiota also contributes to the differentiation and epithelialization of the physical skin barrier (*Figure 1*). Microbes boost the chemical barrier of the skin by producing lipases that digest sebum triglycerides to free fatty acids, which amplifies the acidity of skin and restricts colonization by transient and pathogenic species. Finally, microbes stimulate innate and adaptive immune defenses, such as the release of antimicrobial peptides, induction of neonatal tolerance, and development of protective immunity⁽¹⁹⁾.

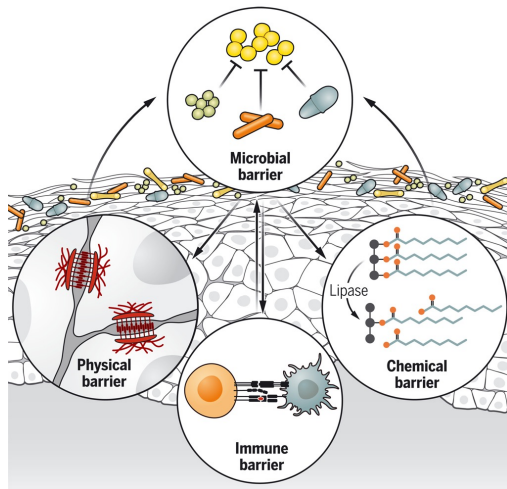


Figure 1. The microbial barrier influences the different skin barrier functions⁽¹⁹⁾

Structural cells such as keratinocytes, fibroblasts and adipocytes contribute to barrier immunity. Specialized immune cells present in the skin include mononuclear phagocytes, such as Langerhans cells, dermal macrophages and dermal dendritic cells, in addition to the resident memory T cells⁽²⁰⁾. Skin barrier immunity changes with age,

and the skin immune composition becomes altered, with reduced Langerhans cells, decreased antigen-specific immunity and increased regulatory populations such as Foxp3+ regulatory T cells. Taken together, these alterations result in decreased skin barrier immunity in the elderly⁽²⁰⁾.

The neurosensory barrier is modulated by unmyelinated C-fibers that express endothelin receptors A and B (ETA and ETB), the capsaicin and heat receptor (TRPV1), and the cold receptors (TRPM8 and TRPA1) (Figure 2). Ligand stimulation induces a burning or itching sensation. Upon sensitization of these sensory neuroreceptors by inflammatory mediators, such as bradykinin, prostaglandins or nerve growth factor, receptor activation is enabled, which possibly underlies sensitive skin. Stress may induce mast cell degranulation, perpetuating the activation of ETA and ETB. Depending on the temperature, TRPV1 expressed in keratinocytes may increase or delay skin barrier recovery, possibly contributing to sensitive skin.

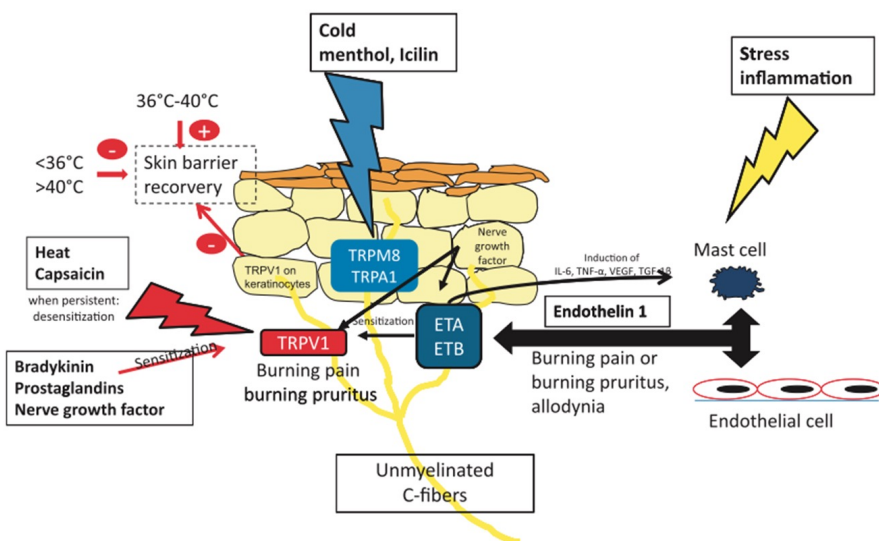


Figure 2. The neurosensory barrier⁽²⁰⁾

IN CONCLUSION

The skin barrier is a living system adapting and evolving throughout our lifetime. There are several barriers, i.e., microbial, physical, chemical, immunological, and neuro-sensory, which act on different pathways of skin physiology and can modulate/interfere with local and systemic reactions. Finally, a healthy barrier is a healthy skin and a healthy body.

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ENVIRONMENTAL TRIGGERS AND THE SKIN BARRIER

KENJI KABASHIMA, M.D.

Kyoto University, Japan

The skin is exposed to various pathogens including bacteria, viruses, proteins, and chemicals (Figure 1). The role of the physical barrier is to block pathogens and thus to prevent infection. Conversely, the role of the immunological barrier (Figure 2) is to eliminate proteins and chemicals and to prevent allergies.

Allergens include proteins of biological origin and chemicals of non-biological origin. In healthy skin, the epidermis has two sets of physical barriers, the stratum corneum (SC) and the tight junctions (TJs)⁽¹⁾. These two

barriers prevent the outside-in penetration of external antigens and the inside-out leakage of internal constituents. For skin appendages, such as hair follicles and sweat glands, TJs are the main barrier to block pathogens⁽²⁾. The size threshold molecular weight, for the TJ barrier, is < 1.000 Da, thus, tacrolimus, gentamicin, steroids, fluorescein isothiocyanate (FITC), 2,4-dinitrofluorobenzene (DNFB), and water are able to penetrate this barrier, but not big pathogens and pollens. Protein antigens are very large and have difficulty penetrating the normal skin. However, some small haptens FITC can easily penetrate the barrier⁽²⁾.

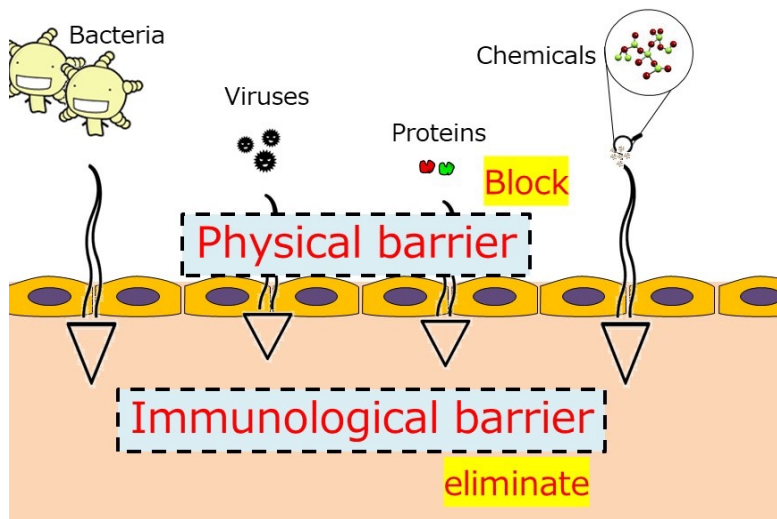


Figure 1 External exposome factors⁽¹⁾

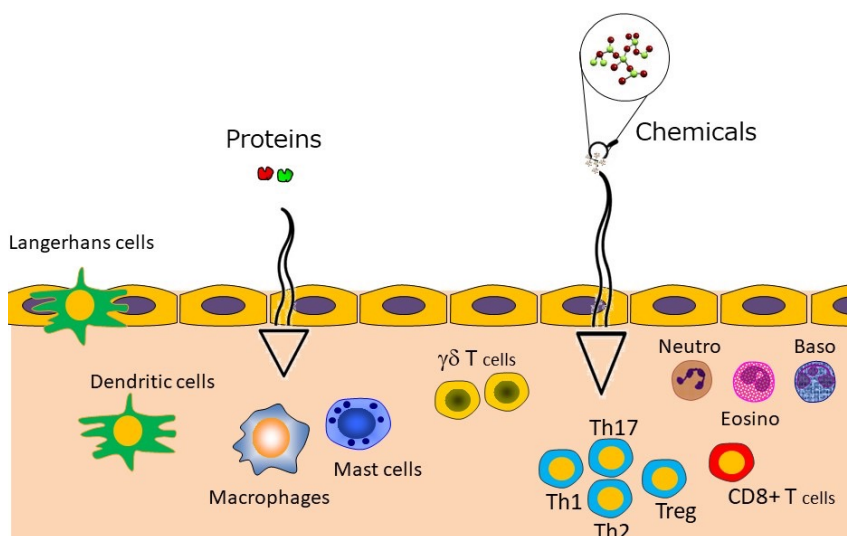


Figure 2 Immunological barrier of the skin⁽¹⁾

Disruption of the skin barrier triggers inflammatory and immunological processes. Protein antigens induce delayed-type hypersensitivity (DTH), such as edematous erythema following insect bites. With chemicals, papules and vesicles observed in eczema are examples of contact hypersensitivity (CHS).

Once a pathogen has entered the skin, the immune response is induced, and dermal dendritic cells migrate more quickly than Langerhans cells to capture antigens. Moreover, neutrophils start to migrate rapidly inside the dermis, faster than T cells. After hapten application, dendritic cells (DCs) exhibit cluster formation around post-capillary venules. Effector T cells are activated within these DC clusters, called inducible skin-associated lymphoid tissue (iSALT), and produce cytokines⁽³⁾.

Allergen exposure through the epidermis can initiate systemic allergy and predispose individuals to the development of one or more atopic diseases, including atopic

dermatitis (AD or eczema), food allergy, asthma, and allergic rhinitis, via the so-called atopic march (Figure 3)⁽⁴⁾. Two independent mutations in the gene encoding filaggrin (FLG) are associated with AD development⁽⁵⁾. FLG is first produced as a polymer (profilaggrin), then becomes a monomer which aggregates to keratin fibers in epithelial cells. Upon degradation, FLG products account in part for the water-holding capacity and maintenance of acidic pH of the SC, both crucial for the epidermal barrier homeostasis by regulating activity of multiple enzymes that control desquamation, lipid synthesis and inflammation⁽⁶⁾. Thus, FLG is an essential protein for SC integrity. Skin barrier impairment in AD patients leads to an increase in transepidermal passage of chemical antigens, and then to chronic inflammation. Repeated elicitation of contact hypersensitivity induces a shift in cutaneous cytokine milieu, from a T helper cell type 1 to a T helper cell 2 profile⁽⁷⁾.

Protein antigens can penetrate the skin barrier in AD patients through small wounds, the protease activity of antigens, injections,

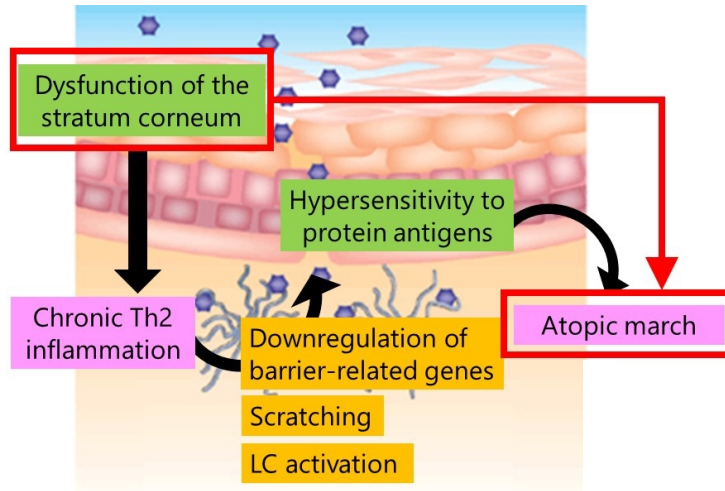


Figure 3 Allergens trigger atopic dermatitis⁽⁴⁾

and/or chronic inflammation. Allergy to latex, a protein sensitization occurring through small skin wounds, can sometimes lead to allergies to banana, avocado, kiwi, chestnuts, etc., called "latex-fruits syndrome"⁽⁸⁾. Pollens have high protease activity, and pollen sensitization can result in pollen-food syndrome⁽⁹⁾. A relationship between tick bites and sensitization to galactose- α -1,3-galactose (α -Gal), with the development of a red meat allergy as a secondary phenomenon has been shown⁽¹⁰⁾. In Japan, jellyfish stings have been linked to sensitization to poly- γ -glutamic acid and, thus, allergy to natto, made with fermented soybeans, which can cause late-onset anaphylaxis⁽¹¹⁾.

Chronic inflammation is the most important mechanism of protein antigen sensitization in AD. Th2 cytokines have the potency to disrupt both the SC and the tight junction barriers^(12, 13). Th2 cytokines can also cause itch by directly stimulating itch sensory neurons⁽¹⁴⁾. Thymic stromal lymphopoietin (TSLP) is an itch mediator, and epithelial cells directly communicate to cutaneous sensory neurons via TSLP to promote itching^(15, 16).

Once type 2 inflammation occurs, Langerhans cells start to migrate in the epidermis quite aggressively. Activated Langerhans cells extend their dendrites upward beyond the tight junction and can directly capture the external protein antigens⁽¹⁷⁾. Thus, dysfunction of the stratum corneum can lead to chronic Th2 inflammation, which induces antigen-specific IgEs, leading to hypersensitivity to protein antigens in other organs. Downregulation of barrier-related genes directly induces scratching and Langerhans cell activation to further induce Type 2 inflammation. As dysfunction of the stratum corneum and hypersensitivity to protein antigens result in the atopic march, it is very important to maintain SC integrity. Gene mutations are responsible for the Netherton syndrome (SPINK5), SAM syndrome (desmoglein 1), and peeling skin syndrome (corneodesmosin), all involving disruption of the keratinocyte barrier function leading to AD-like skin inflammation, high serum IgE level, and high incidence of food allergy. In Japan, the daily application of moisturizer during the first 32 weeks of life has been shown to reduce the risk of AD/eczema in infants⁽¹⁸⁾.

IN CONCLUSION

The skin barrier is composed of the physical (SC and TJs) and immunological barriers to stop environmental aggressors. In terms of atopic dermatitis, the filaggrin-related physical barrier (plus type 2 inflammation) is known to be important. Protein antigens have difficulties penetrating the intact skin barrier. Type 2 skin inflammation causes hypersensitivity to protein antigens. Finally, controlling skin inflammation is important to prevent the development of atopic march.

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NEUROGENIC INFLAMMATION AND MICROBIOME

BRIGITTE DRÉNO, M.D., PH.D.

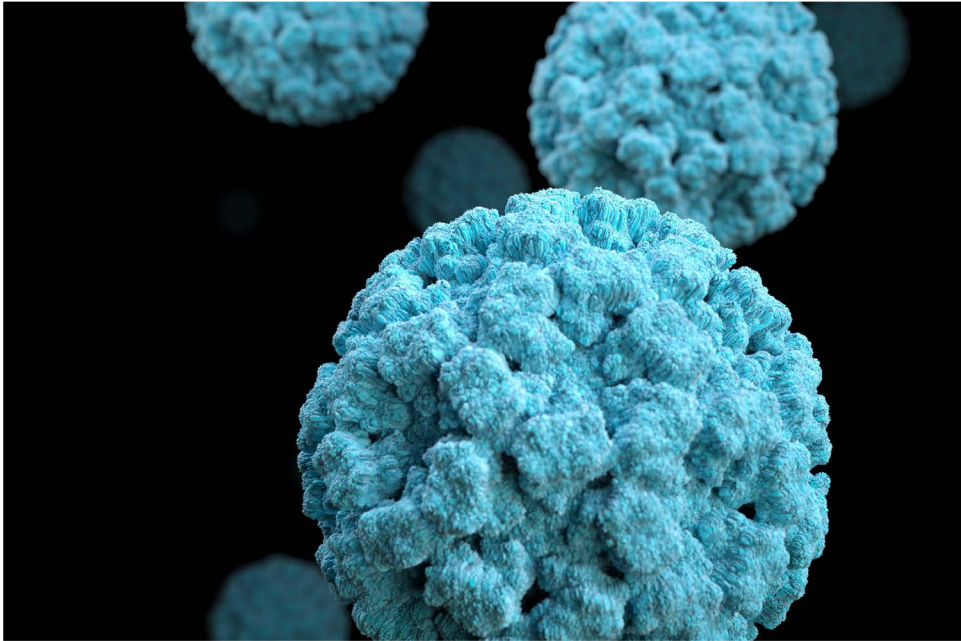
Nantes University, France

Various interactions exist between the skin microbiome and cytokine-triggered and neurogenic inflammations.

Of these 3 actors, the first one is the skin microbiome. As a collection of genomes of microorganisms (bacteria, fungi, viruses) residing on the skin surface, the microbiome plays an important role in human health⁽¹⁾.

The skin microbiome changes over the life course⁽²⁾. *In utero*, the skin is sterile; the initial colonization of the skin occurring during delivery disappears by 6 weeks of age. Then, the skin-like profile is enriched with *Staphylococcus* and *Corynebacterium* species. At puberty, there is a major shift as sex hormones drive maturation of sebaceous glands, causing a proliferation of lipophilic *Cutibacterium Acnes* (*C. acnes*) and *Malassezia* species. In adults, the skin microbiome stability is remarkable, given the continuous disturbances induced by lifestyles, environment, etc. *C. acnes* and the pilosebaceous follicle are likely to be major stabilizers of this effect.

The microbiome is the guardian of skin barrier functions and the first barrier against the environment. It maintains skin homeostasis and controls the skin barrier epithelialization, with the aryl hydrocarbon receptor, a transcriptional factor, playing a major role. The microbiome boosts the skin chemical barrier, produces lipases, digests sebum triglycerides to free fatty acids, and amplifies skin acidity, thus restricting colonization by pathogenic bacteria. It modulates skin immunity and protects against pathogens, and releases extra cellular vesicles and antimicrobial peptides directly or by activating skin cells. For example, *C. acnes*, which represents >50% of bacterial species, produces propionic acid involved in skin odor, stimulates expression of β -defensin 2, and synthesizes vitamin B12 of which a deficiency is associated with hyperpigmentation. *Staphylococcus hominis* produces an antimicrobial peptide (AMP) with a unique inhibitory activity against *S aureus*. *Staphylococcus lugdunensis* produces lugdunin, an AMP-inducing production of the antimicrobial peptide LL-37. *Micrococcus luteus* degrades pollutants and isomerizes urocanic acid, which plays a role in UV protection. *Staphylococcus Epidermis* metabolizes sphingomyelin to produce ceramides^(3, 4).



Bacteria interact among themselves by secreting extracellular vesicles (EVs). *C. acnes* constitutively releases extracellular EVs, increases proliferation of keratinocytes, modulates differentiation of keratinocytes with a decrease in keratin 10 and desmocollin 1 and an increase in filaggrin, and controls some commensal bacteria⁽⁵⁾.

The second actor is cytokine-triggered inflammation. During activation of the innate immunity, the following pattern recognition receptors (PRRs) are expressed either at the surface of skin cells (keratinocytes) or intra-cellularly [toll-like receptors (TLRs), peroxisome-activated receptors (PARs), nod-like intracellular receptors (NLRs 1-3), and rig-like intracellular receptors (RLRs virus)]. AMPs, the natural “antibiotic-like” peptides, are secreted by skin cells⁽¹⁾.

Loss of diversity of the microbiome leads to dysbiosis. In acne, dysbiosis can be linked to the loss of diversity of *C. acnes* phylotypes, with a predominant phylotype: IA1⁽⁶⁾. Loss of diversity of *C. acnes* activates innate immunity⁽⁷⁾. In an *in vitro* skin explant study assessing the impact on innate immune system (IIS) activation after *C. acnes* phylotype diversity loss, innate immune markers (IIMs) were significantly upregulated when incubated with phylotype IA1 alone, compared with the combination IA1 + II + III⁽⁷⁾.

The restoration of microbiome diversity helps to suppress inflammation via the down-regulation of innate immunity. The activation of innate immunity is associated with the secretion of cytokines. Anakinra (Kineret®), a recombinant and slightly modified version of the human interleukin 1 receptor antagonist protein, appears to be a promising option in the treatment of severe acne.

The 3rd actor, neurogenic inflammation, is a process involving the release of neuropeptides from sensory nerves, leading to inflammation⁽⁸⁾. The activation of sensory nerve fibers triggers the release of neuropeptides, such as substance P (SP) and calcitonin gene-related peptide (CGRP). Neuropeptides promote vasodilation, edema, and recruitment of immune cells, resulting in inflammation. Signals from the brain influence skin function, and vice versa, and the skin microbiome plays a crucial role in the skin-brain axis. Afferent fibers, unmyelinated C-fibers, myelin-type A fibers and autonomic nerve fibers are present in the skin with a dense distribution throughout all of its layers. The main neuropeptides released from these fibers include: substance P (SP), calcitonin gene-related peptide (CGRP), neuropeptide Y, natriuretic peptides (NP), and catecholamines (CT).

Interactions occur between nerve fibers and bacteria. *Bacillus cereus* was the first bacteria to show the effect of SP on a cutaneous bacteria⁽⁹⁾. *B. cereus*, a low-level skin bacterium, is responsible for acute cutaneous infections. *In vitro*, SP stimulates the cytotoxicity of *B. cereus* on HaCaT keratinocytes by acting on neurokinin 1 receptors (TACR1). Keratinocytes overproduce collagenase, and *B. cereus* overproduces superoxide dismutase.

Interactions also occur between neuromediators and bacteria. In a healthy microbiome, bacteria interact strongly with each other via production of AMPs modulated by the degradation of enzymes such as peptidases. Neuromodulators modify the interactions between pathogen and commensal bacteria. SP and CGRP stimulate virulence

in *Staphylococcus aureus*, increase adhesion to epidermal cells and production of virulence factors by *Pseudomonas aeruginosa*, and increase biofilm formation in *S. epidermidis* and *P. fluorescens*. Interactions also occur between SP, CGRP and *S. epidermidis*. The thermo-unstable ribosomal elongation factor (EfTu) is translocated to the *S. epidermidis* surface through the mechanosensitive channels MscL. SP diffuses through the bacterial wall in *S. epidermidis*, links to EfTu, inducing the production of a biofilm. CGRP links to DnaK, leading to an increase in *S. epidermidis* virulence. Finally SP and CGRP increase the virulence of *S. epidermidis*. *S. aureus* is not sensitive to CGRP. Exposure of *S. aureus* and *S. epidermidis* to SP leads to increasing their cytotoxicity to keratinocytes through the overexpression of chemokines mRNAs, CCL5, CXCL1, and IL8. These results suggest that SP and CGRP modulate the activity of many diverse cutaneous Gram-positive bacteria on the skin.

Natriuretic peptides (NPs) and catecholamines (CT) are released by blood, capillary endothelial cells, C-tactile fibres and sympathetic C-fibres. NPs and CT mainly modulate *S. aureus*, *S. epidermidis* and *C. acnes* biofilm formation. For *C. acnes*, NP activity is different according to the phenotype. NP and CNP provide an important competitive advantage to *C. acnes* against *S. aureus* for biofilm formation. The activity of natriuretic peptides is dependant on temperature: at 37°C, the development of *S. epidermidis* biofilms is stimulated, whereas that of *S. aureus* is inhibited; at 33°C (skin temperature), the opposite effect can be observed. In the skin, NPs serve as thermostats to regulate biofilm formation activity by the different bacteria.

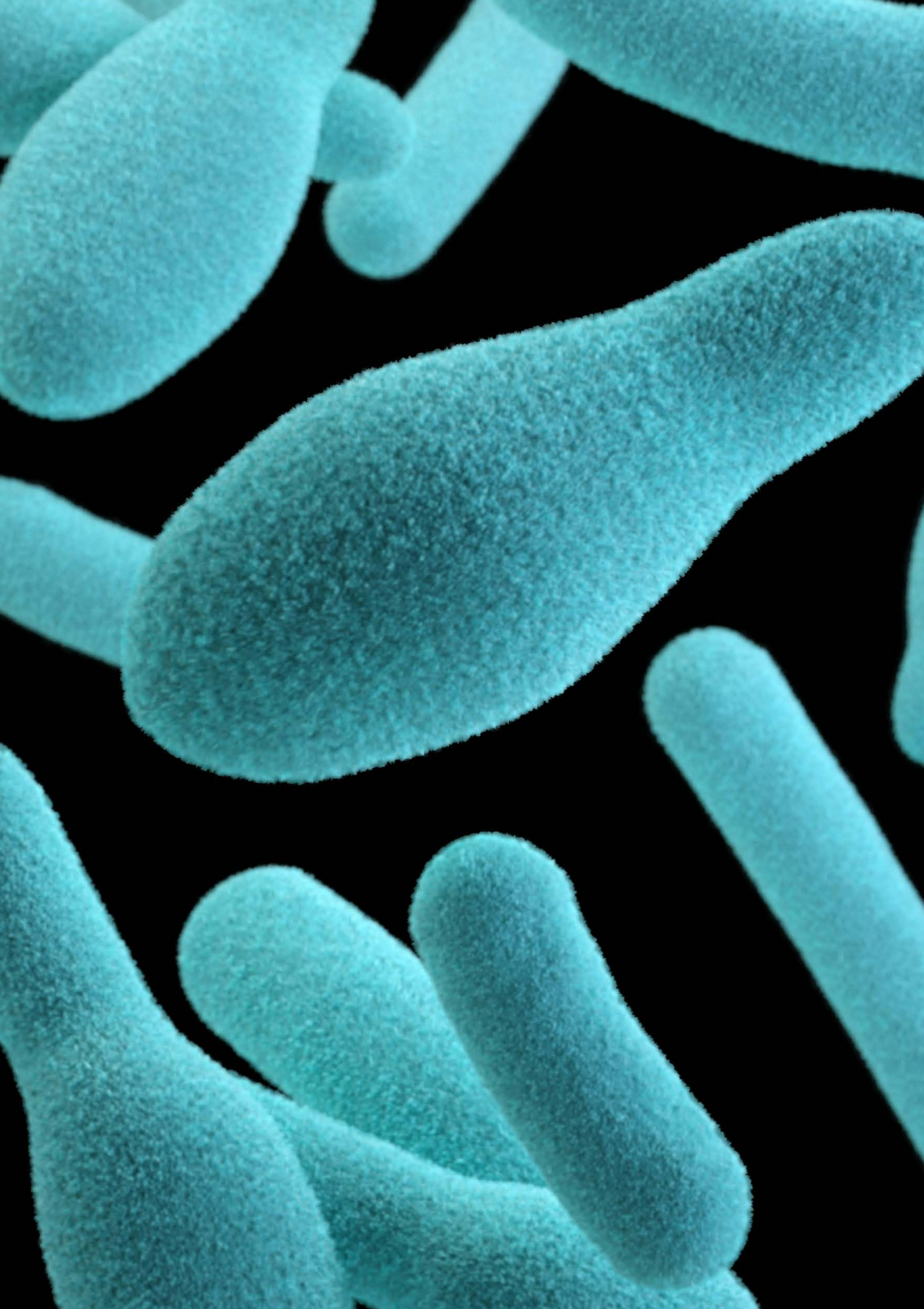
Bacteria are able to interfere with cutaneous physiology by producing molecules showing total or partial homology with skin. Neurotransmitters secrete molecules and are capable of acting as neurotransmitters. *S. epidermidis* and *C. acnes* are able to biosynthesize histamines causing pruritis. Corynebacteria synthesize glutamate, a neurotransmitter released in skin by primary sensory neurons.

Psychological stress induces alteration of the skin barrier function, increasing the severity of group A *Streptococcus pyogenes* infections. This alteration is due to an increase in endogenous glucocorticoids, which inhibits epidermal lipid synthesis by decreasing lamellar body secretion and decreases expression of defensins (cathelin-related AMP and β -defensin 3)⁽¹⁰⁾. SP and its degrading enzymes are involved in the pathogenesis of acne, which in turn might partially explain the pathologic significance of neurogenic and psychogenic aspects in the disease process^(11, 12). SP increases adhesion of both bacteria to the keratinocytes, secretion of enterotoxin C2 by *S. aureus*, and formation of biofilm by *S. epidermidis*⁽¹³⁾.

Strains of the *C. acnes* type III lineage are associated with a skin condition called progressive macular hypomelanosis (PMH)⁽¹⁴⁾. Skin commensal bacteria *S. epidermidis* and its by-product LTA promote melanocyte survival by inducing upregulation of TRAF1, CASP14, CASP5, and TP73. On the other hand, *C. acnes* can inhibit UVB-irradiated melanocyte survival by increasing apoptosis⁽¹⁵⁾. In rosacea, cathelicidin processing is disturbed, resulting in peptide fragments causing inflammation, erythema and telangiectasias⁽¹⁶⁾. Atopic dermatitis (AD) is a good example of combined cytokinetic and neurogenic inflammation.

Antimicrobial peptides LL-37, β -defensins, and dermicidin are decreased in AD skin, favoring *S. aureus* colonization (role of IL-13, IL-31, IL-4). *S. aureus* grows poorly in acidic conditions (normal skin pH=5.3) but grows much better in AD, where pH is higher. *C. acnes*, which maintains the acidic pH of the skin by secreting propionic acid, is decreased in AD. Neurogenic inflammation via SP and natriuretic peptides increase the adherence of *S. aureus* to the stratum corneum. In addition, filaggrin deficiency (genetic or acquired from TH2 skewing), leads to irregular corneocytes facilitating *S. aureus* adherence. *S. epidermidis* and *C. acnes* produce histamine whose role in itching and pruritis is well known. The skin microbiome can control inflammation in AD. Proteases and phenol-soluble modulins- α (PSM α) secreted by *S. aureus* cause epidermal proteolysis and skin barrier damage in mice, which promotes inflammation and itching. *S. hominis* secretes peptides that inhibit PSM α produced by *S. aureus*⁽¹⁷⁾. Inhibition of *S. aureus* activity by clinical *S. hominis* isolate correlates with the prevention of skin barrier damage and inflammation.

The skin microbiome opens the door to bacteriotherapy. Based on *ex vivo* findings of the modulation of inflammation with monoclonal antibodies to the cAMP factor 2, injection of the cAMP factor-targeted acne vaccine directly into acne lesions was proposed for potential use in the future⁽¹⁸⁾. Acnes bacteriophages exist in the pilosebaceous unit, and a potential phage therapy targeting only *C. acnes* phylotype implicated in acne (1A1 mainly) has also been proposed for future research^(19, 20). The rationale for a potential role of probiotics (live microorganisms) or prebiotics is based on their potential to correct dysbiosis. Antimicrobial peptides are produced by commensal bacteria of the microbiome.



IN CONCLUSION

The microbiome opens the door to ecobiological health science. By studying the relationship between living organisms and their environment (exposome and other organs), this interdisciplinary field combines the principles and methodologies from biology and environmental science. Through a comprehensive understanding of the permanently moving network that drives our body health and an ecobiological approach particularly adapted to the skin, the final goal of global health and harmony of the body could be reached.

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ATOPY: AN ECOBIOLOGICAL APPROACH FOR A TARGETED SKIN CARE FORMULA

STÉPHANE FAUVERGHE, M.D.

NAOS Medical Director, Lyon, France

Since its beginning, BIODERMA has chosen an ecobiological approach to develop products that favor biomimetic ingredients and act on skin-own mechanisms and causes rather than on clinical signs alone. One of the objectives of BIODERMA has been to develop a skin care formula that is specifically adapted to atopic dermatitis (AD).

In AD, an impaired skin barrier leads to allergen penetration, a decrease in natural moisturizing factor linked to filaggrin deficiency, an increase in TEWL linked to deficiency in the hydrolipidic film, and in an inadequate ceramide/cholesterol ratio. This generates skin inflammation and results in the clinical signs and symptoms of AD. Cutaneous bacterial dysbiosis is linked to the loss of diversity and an increase in the *S. aureus* biofilm, particularly during flare-ups^(1, 2).

Both EU and US guidelines underline the importance of emollients in AD management^(1, 3, 4). Current national and international recommendations suggest at least a once-daily application of moisturizers⁽⁵⁻⁷⁾, and in appropriate quantity depending on the formulation^(8, 9).

Lack of compliance is an issue as only one-third of atopic patients comply with their topical treatment, and half of the patients use less emollient than the recommended quantity^(10, 11). Thus, it is necessary to propose emollients combining clinical efficacy, inflammation management, itching relief for quality of life (QOL), dermatological safety, optimal tolerance for all skin types, and adapted texture and sensoriality to encourage compliance.

Atoderm® Intensive baume was developed by BIODERMA as an ecobiological approach to manage atopy. This balm contains Lipigenium™, composed of biomimetic lipids and phytosphingosine, activates ceramide neosynthesis, restores filaggrin neosynthesis, and thus helps rebuild the skin barrier⁽¹²⁻¹⁴⁾. Palmitoylethanolamide (PEA), Atoderm® Intensive baume's second ingredient, is a biomimetic fatty acid known to regulate pruritus and improve skin comfort and quality of life of AD patients^(15, 16). PEA provides relief from itching by acting on TSLP. In one study, after 3 weeks of use, there was a 70% reduction in itching score ($p < 0.0001$), 94% of patients reported a decrease in their urge to scratch, and 88% of patients said that itching had stopped durably⁽¹⁷⁾.



AD is characterized by a dysbiosis, with a sharp decline in microbial diversity. During AD flares, biofilm-growing *Staphylococcus aureus* emerges, in strict association with disease severity, as the major colonizer in AD skin lesions⁽¹⁸⁾.

Atoderm® Intensive baume enhances skin barrier therapy, by limiting *S. aureus* adhesion on human corneocytes and inhibiting the formation of the *S. aureus* biofilm, thus preserving the skin microbiotic balance by acting on causes. Atoderm® Intensive baume decreases the amount of bacteria on the skin surface by 99.48%⁽¹⁹⁾. Moreover, Atoderm® Intensive baume prevents flares⁽²⁰⁾. In that monocentric, double-blind, randomized, placebo-controlled study, 130 subjects (aged 6 months to 15 years) with moderate AD (SCORAD 15-40) were treated for 6 months with topical corticosteroids or tacro/picrolimus in combination with Atoderm® Intensive baume or a basic emollient (placebo). After 6 months, there were significant improvements in quality of life (QOL), with

an 89% increase in QOL score, in SCORAD (51% improvement), and in PO-SCORAD (55% improvement)⁽²⁰⁾. In addition, more than ¾ of the patients had no relapse during the 6 months, and for the other 25% the time between flare-ups was 20 days or more. Finally, flare up severity was cut by half. Another study was conducted to assess the efficacy of Atoderm® Intensive baume on skin dryness during and after an outbreak of AD in subjects aged over 3 months and treated with topical corticosteroids⁽²⁰⁾. In this multicentric, prospective observational study, 125 subjects (>3 months old) with light to severe AD received 1 or 2 applications /day of Atoderm® Intensive baume on the face and body for 2 months⁽²⁰⁾. There were significant improvements in QOL, with an 85% decrease in pruritus and an 89% decrease in insomnia after 2 months. Dryness was reduced by 81% and scales by 92%⁽²⁰⁾. At the end of the study, 85% of the parents stated that they were no longer affected by their child's skin problem, and 86% felt that it did not impact their own sleep any longer⁽²⁰⁾.

IN CONCLUSION

Atoderm® Intensive baume supports AD treatment during flare-ups. As a concomitant skin care, it helps the skin to rebuild its barrier and is extremely well tolerated without burning/tingling sensations. Between flare-ups, Atoderm® Intensive baume regulates itching and the inflammatory response, helps the skin strengthen its barrier, and improves the quality of life of patients and families. In both situations, it is very important to use the appropriate quantity of emollient. Atoderm® Intensive baume is an ecobiological approach that combines high efficacy, high tolerance and high compliance, for the well-being of the skin of both patients and their families.

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