

Skin Microbiota and Cosmetics Influence

Subjects: Microbiology

Contributor: Thomas Latire, Mathilde FOURNIERE, Gilles Bedoux

Dermatological and cosmetics fields have recently started to focus on the human skin microbiome and microbiota, since the skin microbiota is involved in the health and dysbiosis of the skin ecosystem. Amongst the skin microorganisms, *Staphylococcus epidermidis* and *Cutibacterium acnes*, both commensal bacteria, appear as skin microbiota sentinels. These sentinels have a key role in the skin ecosystem since they protect and prevent microbiota disequilibrium by fighting pathogens and participate in skin homeostasis through the production of beneficial bacterial metabolites. These bacteria adapt to changing skin microenvironments and can shift to being opportunistic pathogens, forming biofilms, and thus are involved in common skin dysbiosis, such as acne or atopic dermatitis.

Keywords: skin microbiote ; cosmetics ; active compounds

1. Introduction

Research on human microbiota in dermatology started in the 1950's with Kligman, with the improvement in cell culture ^[1]. In 2008, the National Institutes of Health launched the five-year Human Microbiome Project with the objective to sequence human microbiome components. Emergence of new technologies, such as next-generation sequencing (NGS), enable the comprehensive study of the human microbiome ^{[2][3]}. The microbiome corresponds to the set of genomes of microorganisms in symbiosis with the human host, including microorganisms, alive or dead, and free DNA, whereas microbiota refers to microorganisms living in or on a defined ecosystem ^[4]. Human microbiota, mainly established on the skin, oral and vaginal mucosa, as well as in the respiratory, urinary and gastrointestinal tracts, have fundamental roles in health and diseases ^{[5][6]}. The human adult skin microbiota is comprised of diverse microorganisms, including bacteria, fungi ^{[7][8]}, yeasts, viruses ^[9], archaea ^[10] and mites, principally *Demodex* ^[11].

The skin, the largest organ of the human body, is a complex and dynamic ecosystem. Its outer epidermal layer, the stratum corneum, is the first physical barrier that prevents chemical substances or pathogenic microorganisms' entrance, fluid evaporation and body heat loss ^[12]. The stratum corneum is composed of 75–80% proteins, mainly keratins and membrane proteins; 5–15% lipids; and 5–10% unidentified compounds ^{[13][14]}. For the establishment of this microbiota, the skin provides essential nutrients, such as amino acids from the hydrolysis of proteins, fatty acids from the stratum corneum, sweat, lipid hydrolysis or sebum and lactic acids from sweat ^[15]. Microorganisms are based on this stratum corneum and on the deeper cutaneous appendages ^{[16][17]}. The skin microbiota, made up of millions of commensal microorganisms, e.g., 1 million/cm², is the second largest microbiota of the human body in mass ^[18]. Cutaneous bacteria belong to four main phyla among the thirty-six known ^[17]. The average skin body distribution of these main bacteria phyla, detected on 20 diverse skin sites of 10 healthy individuals, were found to be Actinobacteria at 51.8%, Firmicutes at 24.4%, Proteobacteria at 16.5% and Bacteroidetes at 6.3% ^{[19][20]}. Procurement of the skin microbiota occurs in the early stage of birth. In utero, the skin is sterile, devoid of any microorganism, and is colonized a few minutes after birth by commensal microorganisms of the mother, depending on the childbirth method ^{[21][22][23]}. This colonization process in the neonatal stage is essential for the establishment of immune tolerance towards commensal microorganisms ^{[1][24]}. Microbiota colonization continues during growth until reaching an equilibrium state in adulthood ^[1].

An increase in scientific investigations into skin microbiota has inevitably led to the emergence of related studies in the cosmetic industry ^[25], which has now become unavoidable in the cosmetic market. Indeed, skin microbiota is involved in the maintenance of a healthy cutaneous barrier. Skin and microorganisms live in symbiosis and microorganisms help to maintain the skin barrier, the immune system and limit pathogenic microorganism growth ^[26]. However, an imbalance in skin microbiota, called dysbiosis, is correlated with skin pathological diseases, such as acne with the loss of phylotype diversity of *Cutibacterium acnes* and atopic dermatitis with the increase in pathogenic *Staphylococcus aureus* and commensal *Staphylococcus epidermidis*, as well as with non-pathological diseases, such as sensitive and dry skins ^[4].

The principal challenges of such research are (1) the comprehension of each microorganism's role in healthy and beautiful skin; (2) the description of the microbiota of weakened skin with the precise determined cause or consequence of the correlation between dysbiosis and skin pathologies; and (3) evaluation studies of the active ingredients' effect on skin microbiota. The maintenance, protection and restoration of microbiota diversity and equilibrium, as well as prevention of skin dysbiosis, are the emerging claims of cosmetic products. These products may include prebiotics, probiotics, post-biotics or active ingredients with demonstrated effects.

2. Influence of Cosmetics on Skin Microbiota

Intra- and inter-individual multiplicities in microbiota and chemical composition highlighted in the study of Bouslimani et al. (2015) are a huge challenge for the cosmetics industry and skincare development. Molecules associated with skincare or hygiene products last on the skin after their use despite several washings and these products might alter molecular and bacterial diversity [20].

Cosmetic ingredients used that are either functional ingredients, such as preservatives, oils and emulsifiers, or active ingredients, impact the skin microbiota and require attention. Indeed, conventional skincare or hygiene products such as soap, gel and cream contain preservatives and natural and synthetic chemicals that impact microbiota even if these effects are not fully investigated in detail for now [26][27]. Preservatives, such as phenoxyethanol, parabens, and methylisothiazolinone, are known to inhibit the survival of skin commensal bacteria such as *S. epidermidis* [28]. This alteration phenomenon depends on the residual activity of the preservative in the cutaneous environment [27]. Functional ingredients like oils, emulsifiers, fatty acids, gelling agents, thickeners and basic cosmetics for skincare, which both improve skin hydration, modulate microbiota diversity [29][30][31][32][33].

Dermo-cosmetic companies must undergo studies on skin microbiota while developing new ingredients or products in order to ensure consumers that their products maintain, improve a healthy microbiome, or restore a healthy skin-microbiome balance in case of a disturbed microbiome [34][35][36]. In 2019, the first certification of being "Microbiome-friendly", set up by "MyMicrobiome", appeared for a final cosmetic product. This certification is to validate that the product is contamination-free, that specific bacteria of the targeted area will be unharmed, that the microbiome diversity is preserved and that the skin balance is not disturbed (not by the suppression of commensals nor by the stimulation of pathogenic bacteria). For instance, a study in 2018 evaluated three different face washes, two "everyday" products and one 100% natural product, applied on 32 women's upper volar forearm, for their effects on skin microbiome diversity, along with skin pH, moisture and trans-epidermal water loss (TEWL), washing twice a day for 4 weeks. Volunteers were divided into three groups according to skin characteristics: skin pH acid/normal/alkaline, very dry/dry/moist skin and very healthy/healthy/normal/stressed/critical skins, and each was assigned one product. All groups exhibited an increase in alpha diversity (species richness via operational taxonomic unit count and species diversity via the Shannon index) over time and their skin moved to a "healthier" state. The study suggested that synthetic ingredients modified the microbiota diversity, especially within the first two weeks [33]. A more recent study (2019) tested the ability of *S. epidermidis* to metabolize the functional ingredients commonly found in dermo-cosmetics formulae with native oils and waxes, fatty acid esters, fatty acid alcohols, fatty alcohols ethers, fatty acids and other [31]. The tested substances were formulated in water-in-oil and oil-in-water emulsions at concentrations commonly used in cosmetics—10% for oils, 5% for emulsifiers and 2% for fatty acids; the gelling agents and thickeners did not affect the in vitro bacterial growth of commensal *S. epidermidis* [31], and thus are microbiota friendly for this commensal. Future investigations should be performed to determine the effect of functional ingredients on skin microbiota diversity.

The cosmetic industry targets the potential of skin microbiota with more and more studies conducted on the research of active ingredients targeting skin microbiota and the assessment of their action mode: skin benefits, microbiota balance and bacteria physiology, such as induction or suppression of metabolic pathways, adhesion, biofilm formation, growth kinetics, virulence factors, quorum sensing, etc. [1][37][38]. Active cosmetic ingredients that target skin microbiota can be classified into the following categories:

- active ingredients, algal- or plant-based, and thermal water-based, which are not a nutrient source for microorganism;
- prebiotics: nutrients that confer a health benefit with modulation of structure and functionality of the host microbiota in topical application for the cosmetic sector [26]. Cosmetic prebiotic approaches are to maintain healthy skin microbiota, or improve the skin microbiota composition by limiting or reducing pathogen growth and in the same time preserve or stimulate commensal bacteria growth [39][40][41][42];
- probiotics: fragmented bacteria that confer health benefits to the host. Cosmetic products with "probiotics" or "probiotic ingredients" often contain non-viable bacteria, products of bacterial fermentation or cell lysates, which do not require changes in the preservative ingredient system [26]. Nevertheless, cosmetic products containing fragments of microorganisms as probiotics require care regarding safe production. For now, a strict definition of a probiotic in

cosmetic products has not been established and these products should only follow European Cosmetic Regulation 1223/2009 [43];

- post-biotics: bacterial metabolites and or cell wall components released by probiotic microorganisms [27].

The major identified applications of active cosmetic ingredients targeting particularly *S. epidermidis* and *C. acnes* are (1) promotion of commensal metabolism and/or bacterial diversity with the ratio *S. epidermidis*/*C. acnes* for limitation of pathogen invasion; (2) reduction of pathogen growth, virulence and biofilms; and (3) modulation of the skin microenvironment and modulation of the immune responses since the cause–consequence link between the skin pathologies and microbiota dysbiosis is not yet established. These claims in dermo-cosmetics are based on the evaluation methods: “Evaluation Methods of Skin Microbiota Targeting *Staphylococcus epidermidis* and *Cutibacterium acnes* from a Cosmetics Perspective”.

References

1. Dréno, B.; Araviiskaia, E.; Berardesca, E.; Gontijo, G.; Sanchez Viera, M.; Xiang, L.F.; Martin, R.; Bieber, T. Microbiome in healthy skin, update for dermatologists. *J. Eur. Acad. Dermatol. Venereol.* 2016, 30, 2038–2047.
2. Oh, J.; Byrd, A.L.; Deming, C.; Conlan, S.; NISC Comparative Sequencing Program; Kong, H.H.; Segre, J.A. Biogeography and individuality shape function in the human skin metagenome. *Nature* 2014, 514, 59–64.
3. Huttenhower, C.; Gevers, D.; Knight, R.; Abubucker, S.; Badger, J.H.; Chinwalla, A.T.; Creasy, H.H.; Earl, A.M.; Fitzgerald, M.G.; Fulton, R.S.; et al. Structure, function and diversity of the healthy human microbiome. *Nature* 2012, 486, 207–214.
4. Grice, E.A.; Segre, J.A. The skin microbiome. *Nat. Rev. Microbiol.* 2011, 9, 244–253.
5. Dethlefsen, L.; McFall-Ngai, M.; Relman, D.A. An ecological and evolutionary perspective on humang-microbe mutualism and disease. *Nature* 2007, 449, 811–818.
6. Wilson, M.; Houpt, E.R. An introduction to the human–microbe symbiosis. In *Microbial Inhabitants of Humans: Their Ecology and Role in Health and Disease*; Cambridge University Press: Cambridge, UK, 2005; pp. 1–50. ISBN 9780511735080.
7. Findley, K.; Oh, J.; Yang, J.; Conlan, S.; Deming, C.; Meyer, J.A.; Schoenfeld, D.; Nomicos, E.; Park, M.; Kong, H.H.; et al. Topographic diversity of fungal and bacterial communities in human skin. *Nature* 2013, 498, 367–370.
8. Gao, Z.; Perez-Perez, G.I.; Chen, Y.; Blaser, M.J. Quantitation of major human cutaneous bacterial and fungal populations. *J. Clin. Microbiol.* 2010, 48, 3575–3581.
9. Foulongne, V.; Sauvage, V.; Hebert, C.; Dereure, O.; Cheval, J.; Gouilh, M.A.; Pariente, K.; Segondy, M.; Burguière, A.; Manuguerra, J.-C.; et al. Human Skin Microbiota: High Diversity of DNA Viruses Identified on the Human Skin by High Throughput Sequencing. *PLoS ONE* 2012, 7, e38499.
10. Probst, A.J.; Auerbach, A.K.; Moissl-Eichinger, C. Archaea on Human Skin. *PLoS ONE* 2013, 8, e65388.
11. Lacey, N.; Kavanagh, K.; Tseng, S.C.G. Under the lash: Demodex mites in human diseases. *Biochemist* 2009, 31, 20–24.
12. Robinson, P.J. Skin. In *Encyclopedia of Toxicology*, 3rd ed.; Elsevier: Amsterdam, The Netherlands, 2014; pp. 283–309. ISBN 9780123864543.
13. Wilkes, G.L.; Brown, I.A.; Wildnauer, R.H. The biomechanical properties of skin. *CRC Crit. Rev. Bioeng.* 1973, 1, 453–495.
14. Proksch, E.; Brandner, J.M.; Jensen, J.-M. The skin: An indispensable barrier. *Exp. Dermatol.* 2008, 17, 1063–1072.
15. Wilson, M. The Indigenous Microbiota of the Skin. In *The Human Microbiota in Health and Disease: An Ecological and Community-Based Approach*; Garland Science, Ed.; Taylor & Francis: Cambridge, UK, 2018; pp. 86–95. ISBN 978-0815345855.
16. Karkman, A.; Lehtimäki, J.; Ruokolainen, L. The ecology of human microbiota: Dynamics and diversity in health and disease. *Ann. N. Y. Acad. Sci.* 2017, 1399, 78–92.
17. Kong, H.H.; Segre, J.A. Skin Microbiome: Looking Back to Move Forward. *J. Investig. Dermatol.* 2012, 132, 933–939.
18. Byrd, A.L.; Belkaid, Y.; Segre, J.A. The human skin microbiome. *Nat. Rev. Microbiol.* 2018, 16, 143–155.
19. Grice, E.A.; Kong, H.H.; Conlan, S.; Deming, C.B.; Davis, J.; Young, A.C.; Bouffard, G.G.; Blakesley, R.W.; Murray, P.R.; Green, E.D.; et al. Topographical and Temporal Diversity of the Human Skin Microbiome. *Science (80-)* 2009, 324, 1190–1192.

20. Bouslimani, A.; Porto, C.; Rath, C.M.; Wang, M.; Guo, Y.; Gonzalez, A.; Berg-Lyon, D.; Ackermann, G.; Moeller Christensen, G.J.; Nakatsuji, T.; et al. Molecular cartography of the human skin surface in 3D. *Proc. Natl. Acad. Sci. USA* 2015, 112, E2120–E2129.
 21. Dominguez-Bello, M.G.; Costello, E.K.; Contreras, M.; Magris, M.; Hidalgo, G.; Fierer, N.; Knight, R. Delivery mode shapes the acquisition and structure of the initial microbiota across multiple body habitats in newborns. *Proc. Natl. Acad. Sci. USA* 2010, 107, 11971–11975.
 22. Capone, K.A.; Dowd, S.E.; Stamatas, G.N.; Nikolovski, J. Diversity of the human skin microbiome early in life. *J. Investig. Dermatol.* 2011, 131, 2026–2032.
 23. Ladizinski, B.; McLean, R.; Lee, K.C.; Elpern, D.J.; Eron, L. The human skin microbiome. *Int. J. Dermatol.* 2014, 53, 1177–1179.
 24. Mueller, N.T.; Bakacs, E.; Combellick, J.; Grigoryan, Z.; Dominguez-Bello, M.G. The infant microbiome development: Mom matters. *Trends Mol. Med.* 2015, 21, 109–117.
 25. ALCIMED. Skin Microbiome—The Development of a Science that will Transform the Cosmetic Industry; ALCIMED: Paris, France, 2016.
 26. Sfriso, R.; Egert, M.; Gempeler, M.; Voegeli, R.; Campiche, R. Revealing the secret life of skin-with the microbiome you never walk alone. *Int. J. Cosmet. Sci.* 2019, 42, 116–126.
 27. Holland, K.T.; Bojar, R.A. Cosmetics: What is their influence on the skin microflora? *Am. J. Clin. Dermatol.* 2002, 3, 445–449.
 28. Wang, Q.; Cui, S.; Zhou, L.; He, K.; Song, L.; Liang, H.; He, C. Effect of cosmetic chemical preservatives on resident flora isolated from healthy facial skin. *J. Cosmet. Dermatol.* 2019, 18, 652–658.
 29. Staudinger, T.; Pipal, A.; Redl, B. Molecular analysis of the prevalent microbiota of human male and female forehead skin compared to forearm skin and the influence of make-up. *J. Appl. Microbiol.* 2011, 110, 1381–1389.
 30. Lee, H.J.; Jeong, S.E.; Lee, S.; Kim, S.; Han, H.; Jeon, C.O. Effects of cosmetics on the skin microbiome of facial cheeks with different hydration levels. *Microbiologyopen* 2017, 7, e00557.
 31. Dobler, D.; Schmidts, T.; Wildenhain, S.; Seewald, I.; Merzhäuser, M.; Runkel, F. Impact of Selected Cosmetic Ingredients on Common Microorganisms of Healthy Human Skin. *Cosmetics* 2019, 6, 45.
 32. Callewaert, C.; Hutapea, P.; Van de Wiele, T.; Boon, N. Deodorants and antiperspirants affect the axillary bacterial community. *Arch. Dermatol. Res.* 2014, 306, 701–710.
 33. Wallen-Russell, C. The Role of Every-Day Cosmetics in Altering the Skin Microbiome: A Study Using Biodiversity. *Cosmetics* 2018, 6, 2.
 34. Lopes, E.G.; Moreira, D.A.; Gullón, P.; Gullón, B.; Cardelle-Cobas, A.; Tavaría, F.K. Topical application of probiotics in skin: Adhesion, antimicrobial and antibiofilm in vitro assays. *J. Appl. Microbiol.* 2017, 122, 450–461.
 35. Shokryazdan, P.; Siew, C.C.; Kalavathy, R.; Liang, J.B.; Alitheen, N.B.; Faseleh Jahromi, M.; Ho, Y.W. Probiotic potential of *Lactobacillus* strains with antimicrobial activity against some human pathogenic strains. *BioMed Res. Int.* 2014, 2014, 927268.
 36. Khmaladze, I.; Butler, É.; Fabre, S.; Gillbro, J.M. *Lactobacillus reuteri* DSM 17938—A comparative study on the effect of probiotics and lysates on human skin. *Exp. Dermatol.* 2019, 28, 822–828.
 37. Percoco, G. Methods and results in 3D skin biopsies. In Proceedings of the The Microbiome of the Skin-New Avenues of Research, in cosmetics, Amsterdam, The Netherlands, 17 April 2018.
 38. Hillion, M. Interactions peau/microbiote cutané: Étude du microbiote cutané cultivable et influence de produits cosmétiques sur la virulence bactérienne. Apports de la technique de spectrométrie de masse MALDI-TOF. Ph.D. Thesis, Thèse de l'Université de Rouen, Évreux, France, 2013.
 39. Krutmann, J. Pre- and probiotics for human skin. *J. Dermatol. Sci.* 2009, 54, 1–5.
 40. Bojar, R.A. Studying the human skin microbiome using 3D in vitro skin models. *Appl. In Vitro Toxicol.* 2015, 1, 165–171.
 41. Baldwin, H.E.; Bhatia, N.D.; Friedman, A.; Eng, R.M.; Seite, S. The role of cutaneous microbiota harmony in maintaining a functional skin barrier. *J. Drugs Dermatol.* 2017, 16, 12–18.
 42. Seite, S.; Misery, L. Skin sensitivity and skin microbiota: Is there a link? *Exp. Dermatol.* 2018, 27, 1061–1064.
 43. Parlement Européen et du Conseil. Règlement (CE) No 1223/2009 du PARLEMENT Européen et du Conseil du 30 Novembre 2009 Relatif Aux Produits Cosmétiques; Office des publications de l'Union européenne: Luxembourg, 2009.
-

