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THE ROLE OF EPIDERMAL BARRIER DYSFUNCTION  
AND CUTANEOUS MICROBIOME DYSBIOSIS  
IN THE PATHOGENESIS AND MANAGEMENT OF  
ACNE VULGARIS AND ROSACEA

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# Dysbiosis, (Barrier) Dysfunction, and Dermatoses: A Chicken-and-Egg Dilemma

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**A**cne vulgaris (AV) and rosacea are two of the most common dermatoses diagnosed and managed by dermatologists.<sup>1,2</sup> Despite this and our improved understanding of the unique pathogenesis of each, there has been little focus on general skin care and how it may affect physiologic functioning of the epidermis until recently. Under optimal homeostatic conditions, the skin generally provides a physical, chemical, and immune layer that deftly balances moisture, temperature, and biochemical substrates.<sup>3</sup> These factors create biomes specific to different areas of the body including the skin, teeth, and internal/external mucosa, where various microorganisms (including bacteria, viruses, and fungi) can thrive.<sup>3</sup> Given that bacteria alone outnumber human cells more than 10 to 1 and have over 100 times more genes than humans, the microbiome, and the factors that support it, cannot be ignored.<sup>4</sup> While alterations in epidermal barrier function and the disruption of the cutaneous microbiome (CM) have been studied in a few dermatoses, most notably atopic dermatitis, these may not be the first to come to mind when treating AV and/or rosacea.

One of the most prominent pillars of AV pathogenesis is the follicular colonization with *Cutibacterium acnes*. Although typically a commensal organism, virulent strains of *C. acnes*—phylogroup IA1 in particular—have been associated with AV and worsening acne severity.<sup>5</sup> With the onset of puberty, there is also a notable change in epidermal barrier function as measured by increased transepidermal water loss (TEWL).<sup>6</sup> TEWL is further compounded in patients with AV, which may be up to 50% greater depending on acne severity.<sup>7</sup>

Rosacea pathogenesis is thought to be a complex, interconnected web of dysregulated innate and adaptive immunity combined with hyperreactive neurovasculature. In practice, many of the secondary criteria for rosacea (eg, burning, stinging, edema, dryness)<sup>8</sup> are also symptoms of epidermal barrier deficiency. Multiple studies have demonstrated that the addition of physiologic moisturizers and/or cleansers to standard-of-care regimens may augment treatment outcomes for mild-moderate disease.<sup>9-13</sup> Unlike AV in which one organism has been implicated, rosacea has been associated with multiple organisms including bacteria (eg, *Streptococcus epidermidis*, *Bacillus oleronius*, and *Helicobacter pylori*), *Demodex* species, a decreased abundance of *C. acnes*, and even the general overgrowth of bacteria in the small intestine (SIBO).<sup>14</sup> Although a singular perpetrator has not been conclusively identified, multiple studies have found dysbiosis within the rosacea cutaneous (and enteric) microbiome.

Both AV and rosacea have varying degrees of epidermal barrier dysfunction, which may be further altered by topical and even systemic therapies. Similarly, AV and rosacea have unique cutaneous microbiomes and microbiome derangement that can be both directly and indirectly affected by common treatment options, including the indiscriminate use of broad-spectrum antibiotics. With derangements at baseline that may be further exacerbated by current therapeutic paradigms, discussion of proper skin care is close to becoming a cornerstone of care. Every acne and rosacea encounter should endeavor to discuss key factors in moisturizers and cleansers: physiologic to slightly acidic pH; scent-/fragrance-free; emollients to soften the skin; humectants, such as glycerin and hyaluronic acid, to attract water; occlusives like petrolatum to seal them in, and proper lipid composition to potentially supplement those deficient in disease states.

A final interesting concept to consider: the chicken and the egg. A healthy microbiome is dependent on a functioning epidermis with adequate moisture and nutrients for survival and balance. In turn, a healthy microbiome can quell virulent organisms and provide a suitable pH to optimize biochemical and immunological epidermal processes. In AV and rosacea, there are data to suggest that both dysbiosis and epidermal barrier dysfunction have roles to play. But does one precede (or even supersede) the other? And, more importantly, if dermatologists have the potential to repair and even restore both, how much might it matter in clinical practice? While research continues to elucidate the role of both dysbiosis and barrier dysfunction in acne and rosacea, it behooves us to employ quality skin care to restore and repair so the chicken and egg concept is irrelevant. No visit for either disease is complete until quality skin care has been discussed.

## REFERENCES

1. Landis ET, Davis SA, Taheri A, Feldman SR. Top dermatologic diagnoses by age. *Dermatol Online J*. 2014;20(4). doi:10.5070/d3204022368.
2. Peck GM, Roberson FA, Feldman SR. Why do patients in the united states seek care from dermatologists? *Dermatol Ther*. 2022;12(4):1065-1072. doi:10.1007/s13555-022-00706-0.
3. Baldwin HE, Bhatia ND, Friedman A, Eng RM, Seite S. The role of cutaneous microbiota harmony in maintaining a functional skin barrier. *J Drugs Dermatol*. 2017;16(1):12-18.
4. NIH HMP Working Group, Peterson J, Garges S, et al. The NIH Human Microbiome Project. *Genome Res*. 2009;19(12):2317-2323. doi:10.1101/gr.096651.109.
5. Cunliffe WJ, Gould DJ. Prevalence of facial AV in late adolescence and in adults. *Br Med J*. 1979;1(6171):1109-1110. doi:10.1136/bmj.1.6171.1109
6. Pappas, A., Dunn, K., Cula, G.O., et al. Barrier and microbiome changes in the facial skin of children as they approach puberty. Presented at: 26th European Academy of Dermatology and Venerology (EADV) Congress: P006513; September 17, 2017; Geneva Switzerland.
7. Yamamoto A, Takenouchi K, Ito M. Impaired water barrier function in AV. *Arch Dermatol Res*. 1995;287(2):214-218. doi:10.1007/BF01262335
8. Gallo RL, Granstein RD, Kang S, et al. Standard classification and pathophysiology of rosacea: The 2017 update by the National Rosacea Society Expert Committee. *J Am Acad Dermatol*. 2018;78(1):148-155. doi:10.1016/j.jaad.2017.08.037.
9. Pelle MT, Crawford GH, James WD. Rosacea: II. Therapy. *J Am Acad Dermatol*. 2004;51(4):499-514. doi:10.1016/j.jaad.2004.03.033
10. Draelos ZD. Cosmetics in acne and rosacea. *Semin Cutan Med Surg*. 2001;20(3):209-214. doi:10.1053/sder.2001.27556
11. Del Rosso JQ. The use of moisturizers as an integral component of topical therapy for rosacea: clinical results based on the assessment of skin characteristics study. *Cutis*. 2009;84(2):72-76.
12. Subramanyan K. Role of mild cleansing in the management of patient skin. *Dermatol Ther*. 2004;17 (Suppl 1):26-34. doi:10.1111/j.1396-0296.2004.04s1003.x
13. Draelos ZD, Green BA, Edison BL. An evaluation of a polyhydroxy acid skin care regimen in combination with azelaic acid 15% gel in rosacea patients. *J Cosmet Dermatol*. 2006;5(1):23-29. doi:10.1111/j.1473-2165.2006.00219.x
14. Holmes AD. Potential role of microorganisms in the pathogenesis of rosacea. *J Am Acad Dermatol*. 2013;69(6):1025-1032. doi:10.1016/j.jaad.2013.08.006

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# The Role of Epidermal Barrier Dysfunction and Cutaneous Microbiome Dysbiosis in the Pathogenesis and Management of Acne Vulgaris and Rosacea

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## ABSTRACT

**Background:** Dysregulation of either the cutaneous microbiome (CM) or epidermal barrier function (EBF) is thought to play an increasingly important role in acne vulgaris (AV) and rosacea pathogenesis.

**Objective:** To review the literature regarding epidermal barrier dysfunction (EBD) and cutaneous dysbiosis in AV and rosacea and provide clinical pearls for dermatologists.

**Methods:** A Medline literature search was performed for relevant literature regarding EBD and dysbiosis and either AV or rosacea. An expert consensus panel was then convened to discuss article merits and distill findings into clinical pearls.

**Results:** Final review included 138 articles. Puberty may alter natural stratum corneum lipid ratios, instigating and/or exacerbating EBD in AV. Patients with severe AV have an abundance of virulent *Cutibacterium acnes* phylotype IA1. EBD may manifest as classic signs of rosacea and improve with treatment. While several microbial populations are dysregulated in rosacea, the effect from any singular species is unclear. Current AV and rosacea treatment regimens may mitigate inflammation but may also indiscriminately damage CM and EBF. Physiologic moisturizers and cleansers that harness pre-/pro-/postbiotics may have a role in restoring CM, EBF, and potentially improving dermatosis severity.

**Limitations:** Limited prospective clinical trial data especially regarding over-the-counter (OTC)/non-prescription skincare products.

**Conclusion:** Appropriately developed prescription and OTC preparations may selectively influence the microbiome and potentially maintain/restore EBF. By understanding this relationship, dermatologists will be better able to educate patients on the importance of appropriate skin care.

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## INTRODUCTION

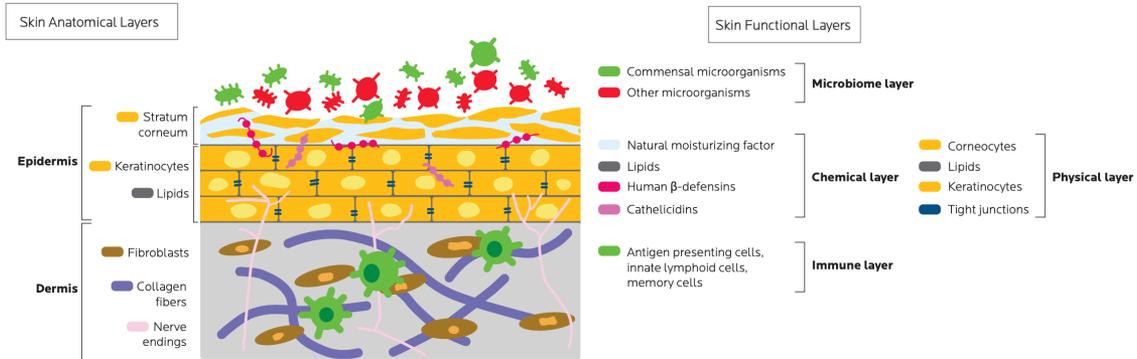
The cutaneous microbiome (CM) is a sprawling ecosystem comprising various microorganisms and their metabolites that dwell on every surface of the human body.<sup>1-4</sup> Together, the human body and microbiome create a functional synergy that is increasingly thought to modulate local and systemic inflammation.

While CM composition varies with moisture, sebaceous gland activity, and anatomical region, these microbes exist symbiotically with other commensal organisms and the surfaces on which they dwell, receiving necessary nutrients (eg, amino acids, fatty acids, moisture) while providing

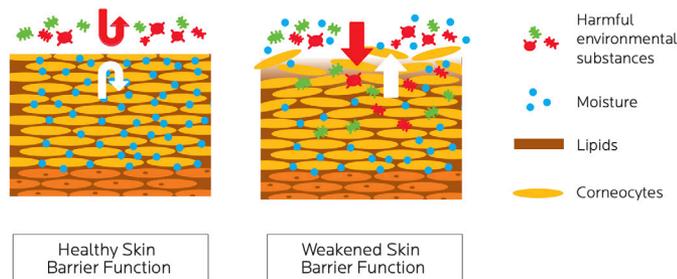
a slightly acidic pH, priming host immune response (eg, induced antimicrobial peptides [AMP] and bacteriocins) and outcompeting classic pathogenic organisms (eg, pathogenic biofilm-producing, strains of *Streptococcus epidermidis* and *Staphylococcus aureus*).<sup>4-10</sup> In this regard, the CM functions as a distinct, organized, and unique layer atop the stratum corneum (Figure 1).<sup>11</sup>

When appropriately functioning, the CM and the epidermal barrier exist in homeostasis.<sup>12-15</sup> The loss of epidermal barrier function (EBF), dysbiosis, or both may result in a dysregulated inflammatory response, as is classically seen in atopic

**FIGURE 1.** Skin barrier anatomy and function. In addition to division by anatomic layers, the skin can also be divided into functional layers (including a physical, chemical, immunological, and microbiome layer) that work in conjunction to promote epidermal (barrier) function.



**FIGURE 2.** Epidermal barrier dysfunction. A functioning epidermal barrier is capable of keeping harmful substances out, retaining moisture, minimizing inflammation, and promoting effective innate immune function. Insults may disrupt the anatomic and/or functional layers and propagate inflammation through the skin.



dermatitis (AD) (Figure 2).<sup>13</sup> Despite increasing understanding in how EBD and dysbiosis may affect inflammation, therapeutic options are limited and still poorly understood.

The purpose of this paper is to review the literature regarding (1) the role that dysbiosis and EBD play in the pathogenesis of acne vulgaris (AV) and rosacea and (2) provide pearls and potential therapeutic interventions to restore CM homeostasis and EBF.

**TABLE 1.**

Post-Meeting Summative Survey Results		
Statements		Mean Score
There were sufficient data to conclude that:	ACNE is a barrier deficiency disorder	4.0
	ROSACEA is a barrier deficiency disorder	4.7
	ACNE is associated with an alteration of the healthy microbiome	5.0
	ROSACEA is associated with an alteration of the healthy microbiome	4.5
	Water and therefore moisturizers are prebiotics	4.2
	Quality skin care can help to restore/normalize the healthy microbiome	4.5
	Normalization of a healthy microbiome can improve ACNE	4.7
	Normalization of a healthy microbiome can improve ROSACEA	4.3

Mean score was calculated from a 5-point Likert scale with 1 being "Strongly Disagree" and 5 being "Strongly Agree"

## MATERIALS AND METHODS

A panel of United States (US) dermatologists recognized for their contributions and prior expertise in the diagnosis and management of AV/rosacea, and/or history of academic achievement were convened on August 4<sup>th</sup>, 2021.

Prior to meeting, a Medline literature search was performed for original studies, meta-analyses, clinical guidelines, and reviews regarding "epidermal barrier dysfunction"; "epidermal barrier function"; "transepidermal water loss"; "cutaneous microbiome"; "dysbiosis"; "probiotics"; "prebiotics"; and "postbiotics". The Boolean term "AND" was used to find intersections between these phrases and either "acne vulgaris" or "rosacea".

Authors discussed topics including: the role of EBD and dysbiosis in AV and rosacea; the role of skin care in AV and rosacea; and agent/ingredients which may complement current standard-of-care prescription regimens.

Results of the discussion and a follow-up survey containing 8 summative statements voted on by panelists (using a 5-point Likert scale with 1 being "Strongly Disagree" and 5 being "Strongly Agree") that was performed on October 14<sup>th</sup>, 2021 are presented below.

## RESULTS

All 8 post-meeting statements received a mean score of  $\geq 4$  (Table 1). The authors emphasized that, although potentially limited by temporal or logistic constraints, dermatologists should endeavor to discuss skin care during every AV/rosacea visit. Optimal skin care may improve both diseases, potentially with minimal pharmacotherapy. Furthermore, poor choices made by an uninformed patient may derail an otherwise ideal therapeutic approach.

### Acne Vulgaris

#### Barrier Dysfunction

Epidermal barrier dysfunction (EBD) is well-known in AV as adverse effects of over-the-counter (OTC) products, and prescription and adjunctive procedural therapies.<sup>16</sup> These effects, clinically noted as xerosis, abnormal sensations (eg, stinging, burning, tingling, pruritus), pain, tightness, and/or an irritant dermatitis, are all signs of increased transepidermal water loss (TEWL) and may be partially mitigated with concurrent use of appropriate moisturizers.<sup>16-21</sup>

Data suggest AV patients may have pre-existing EBD.<sup>22-25</sup> Untreated Japanese male AV patients ages 14-26 (moderate, n=11; mild, n=25) had significantly greater TEWL that correlated with AV severity than age/gender-matched controls (n=29) (TEWL g/m<sup>2</sup>/h $\pm$ SD: Control 10.3 $\pm$ 2.4; Mild, 14.4 $\pm$ 2.5; Moderate, 16.8 $\pm$ 3.8,  $P<.01$ ).<sup>22</sup>

TEWL may worsen as patients approach puberty, regardless of AV, with one study finding among 132 healthy children (boys, n=67; girls, n=65) ages 6-13 that TEWL was significantly greater by age 9 for girls and 11 for boys vs their 6-year-old counterparts ( $P<.05$ ).<sup>25</sup>

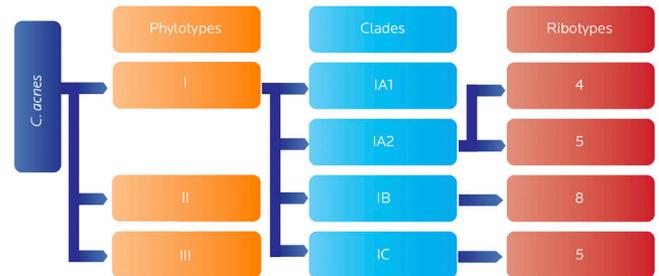
Compared against an internal ceramide standard, a decrease in total stratum corneum (SC) ceramides in AV ( $\mu\text{g}\pm\text{SD}$ : moderate 3.4 $\pm$ 0.45; mild 4.07 $\pm$ .87) vs control (6.49 $\pm$ 0.98) ( $P<.05$ ) had a significant negative correlation with increased TEWL.<sup>23</sup> After isolating total ceramide and sphingosine from other SC lipids, there was also a significant negative correlation with decreasing total ceramide ( $P<0.1$ ) and free sphingosine ( $P<0.1$ ) with worsening AV severity.<sup>23</sup> Of note, similar ceramide derangements have been found in and are thought to pathologically contribute to EBD in psoriasis and AD.<sup>26</sup>

Together these data suggest increases in TEWL may be a result of ceramide imbalances due to adrenarche-induced ceramide synthetic dysfunction. Although these changes may not be specific to AV, EBD may be associated with, and potentially precede and/or exacerbate, clinically-evident AV.

#### Microbiome Dysbiosis

In AV, there are two broadly distinct CMs: superficial and follicular.<sup>27</sup> Superficial facial and truncal CM are dominated

**FIGURE 3.** *Cutibacterium acnes* phylotypes. Single and multi-locus sequencing have found phylotypes IA1 to be predominantly associated with additional virulence factors that instigate/propagate acne vulgaris while phylotypes II (and III) are thought to be predominantly commensal species.



by *Staphylococci spp.* (eg, *S. epidermidis*) which account for >27% of bacteria, while anaerobic *Propionibacteria* (including *Cutibacterium acnes*) account for <2%.<sup>28</sup> Conversely, the physiologically-healthy follicular CM is more homogenous, with *C. acnes* comprising 89% to 94% of the bacterial population, functioning as a commensal organism in healthy skin.<sup>14,29</sup> This disparity may suggest that, unlike in the superficial CM, the sebum-abundant follicular milieu may self-select for lipophilic organisms and that the absence of microbiota species diversity could be the norm.<sup>14</sup>

Counterintuitively, *C. acnes* follicular colonization is one of the 4 dogmatic factors of AV pathogenesis and may correlate with severity.<sup>25,30-32</sup> During puberty there is a significant increase in absolute abundances of organisms<sup>25</sup> and relative abundance of *C. acnes* that parallels increase in sebum production.<sup>33-35</sup> Conversely, the decreased incidence and prevalence of AV (~3-5% of individuals) by the fifth decade of life coincides with up to a 49.9% of the reduction in *Propionibacterium spp.* and may be due to age/age-related changes, a significant decrease in sebum production, and increase in overall CM diversity.<sup>36-38</sup>

*C. acnes* phylotypes are distinct between individuals with and without AV (Figure 3).<sup>27,29,39-42</sup> *C. acnes* phylotype IA1 (and to a lesser degree IA2) represents 55-67% of all *C. acnes* in mild and severe AV (respectively), whereas, in healthy skin, IA1 exists alongside other phylotypes including IA2, II, and III.<sup>42</sup> This homogenization of *C. acnes* phylotypes may correlate with increasing AV severity, as one study found phylotype IA1 comprised 84.4% and 95.6% of all *C. acnes* on the face and back (respectively) of individuals with severe AV compared with healthy individuals who had ~39% phylotype IA1 and ~43% phylotype II.<sup>43</sup>

Compared with *C. acnes* phylotype II, a commensal strain, phylotype IA1 (Figure 3) has pro-inflammatory capabilities via increased activation of toll-like receptor 2 (TLR2) Th1/Th17 axis, decreased interleukin-10 (IL-10) response,<sup>29,44</sup> and a combination of virulence factors including: increased triacylglyceride lipase activity generating short-chain fatty acids,<sup>45,46</sup>  $\beta$ -hemolytic,

membrane pore-forming toxins termed Christie-Atkins-Munch-Petersen (CAMP) factors that may stimulate TLR-2;<sup>47,52</sup> plasmids that promote bacterial adhesion and biofilm formation;<sup>29</sup> lyases that degrade hyaluronic acid and the extracellular matrix, propagating inflammation;<sup>46</sup> porphyrins that induce oxidative stress/damage;<sup>53,54</sup> and reduced loss of clustered regularly interspaced short palindromic repeats (CRISPR) loci allowing increased horizontal acquisition of virulent traits.<sup>55</sup> Furthermore, virulent *C. acnes* may also have an increased resistance to erythromycin, clindamycin, and tetracycline classes of antibiotics.<sup>56,57</sup> This presents potential concern for indiscriminate, inadvertent eradication of commensal organisms (including *C. acnes* phylotype II and III) within already pathologic follicular and superficial CM, allowing any remaining phylotype IA1 to flourish.

While *C. acnes* typically dominates the follicular microbiome, even in AV-prone skin, it does coexist with other microbes (eg, *Malassezia spp.* fungi, *S. epidermidis*, and bacteriophages).<sup>58</sup> Of particular interest is *S. epidermidis*, which has shown potential to limit *C. acnes* growth in vitro,<sup>59</sup> and (*C. acnes* specific) bacteriophages.<sup>60-63</sup> These bacteriophages, which are significantly more abundant in healthy skin vs. acne-prone skin ( $P<.05$ ) may modulate the microbiome in vitro by facilitating the exchange of virulence factors and preferentially killing specific bacteria.<sup>14,60,61</sup>

Harnessing the natural “predator-prey” relationship of these organisms may have potential as a targeted, non-antibiotic AV therapy.<sup>59</sup> Studies also suggest that isotretinoin may achieve its durable therapeutic response by inducing a “sebaceous drought,” resetting the follicular microbiome, and allowing a more diverse CM to repopulate.<sup>64-66</sup>

## Rosacea

### Barrier Dysfunction

Many of the clinical signs of EBD (hyperirritability, burning, stinging, and sensitivity to common OTC skincare products) are hallmarks of rosacea.<sup>17,67,68</sup> EBD may also explain rosacea triggers, including extreme climates and UV exposure and increased likelihood of positive patch test (and risk of allergic/irritant contact dermatitis).<sup>67,69</sup> This may be due in part to a significantly more alkaline central facial skin (pH 5.7 vs 5.2,  $P=.026$ ) and materially different skin surface lipid composition.<sup>70</sup>

EBD in rosacea appears localized to areas of (and those adjacent to) inflammation.<sup>71-73</sup> In a study of unaffected nasolabial-fold and volar-arm skin in rosacea patients ( $n=28$ ) and age/gender-matched controls ( $n=32$ ), nasolabial-fold skin in rosacea patients had materially higher TEWL than healthy controls (21.1 vs 16.8,  $P=0.127$ ) and significantly lower SC hydration (SCH) (16.5 vs 24.4,  $P<.001$ ); these trends were not observed in the volar forearm.<sup>71</sup> Evaluations using topically-applied lactic acid to elicit irritation (ie, lactic acid stinger test [LAST]) have found individuals with erythematotelangiectatic-predominant (ETR) and papulopustular-predominant (PPR) rosacea to be significantly more likely to have a positive (75% vs 18.8%,  $P<.001$ ) and greater LAST response ( $P<.001$ ).<sup>72,74</sup> Worsening TEWL and SCH also correlated with LAST scores and patients’ perception of sensitive skin.<sup>72</sup>

EBD may also correlate with rosacea severity. Not only does concomitant use of more physiologic pH cleansers and moisturizers minimize adverse effects of prescription therapy, but they may also have additive effects in reducing rosacea severity (Table 2).<sup>75-79</sup> A 6-week investigative trial with oral minocycline 100 mg daily, in the absence of topical intervention, has also

TABLE 2.

Epidermal Barrier Outcomes Following Topical and Systemic Management of Rosacea

Study	Study Type & Size (n)	Rosacea Phenotype & Severity	Participant Age	Sex (%F)	Study Length	Treatment Regimen	Results
Del Rosso <sup>77</sup>	Split Face (102)	PPR, mild-moderate	20-39: n=29 40-59: n=52 >60: n=21	85	7 days	Twice daily: Azelaic acid 15% gel and cleanser to full face. Moisturizer to only right half of face.	Significant reduction in CSS with moisturizer use vs baseline ( $P=.008$ ) and vs side without moisturizer ( $P=.015$ )
Subramanyan <sup>78</sup>	RCT (70)	--, moderate	--	--	4 weeks	Metronidazole. Synthetic detergent (syndet) vs soap bar	Significantly less patient self-reported itching, irritation, tingling with syndet vs soap bar ( $P=--$ )
Draelos et al <sup>79</sup>	RCT (67)	PPR and ETR, mild-severe	Range 19-66	100	12 weeks	Twice daily: Azelaic acid 15% gel Gluconolactone cleanser (4%) and moisturizer (10%) vs nonstandardized patient self-selected products	improved facial erythema (week 8, $P=.012$ ; week 12, $P=.001$ ), itching (week 8, $P=.045$ ), stinging (week 12, $P=.029$ ). self-reported significant improvement ( $P<.05$ ) in skin sensitivity, dryness, texture, and smoothness compared with baseline vs controls
Ní Raghallaigh et Powell <sup>80</sup>	Open-label, case-control (29)	PPR, --	< 40: n=6 40-55: n=12 >55: n=11	48.2	6 weeks	Once daily: Minocycline 100 mg	Significant improvement in erythema ( $P=.003$ ) and hydration vs baseline (Central face: 41.4 vs 38.5, $P=.041$ ; Cheeks: 42 vs 37.4, $P=.012$ )

-- Data not reported. CSS, cumulative symptom score (comprising itching, burning, stinging, tingling); ETR, erythematotelangiectatic-predominant rosacea; PPR, papulopustular-predominant rosacea; RCT, randomized controlled trial.

been shown to significantly improve erythema ( $P=.003$ ) and SCH (Central face,  $P=.041$ ; Cheeks,  $P=.012$ ).<sup>80</sup> This may suggest that underlying (systemic) inflammation may exacerbate rosacea severity and EBD in a positive-feedback loop.

#### Microbiome Dysbiosis

Rosacea has several potential key microbes: *Demodex spp.* (*Demodex folliculorum*, *Demodex brevis*), *Bacillus oleronius*, *Staphylococcus epidermidis*, and *Cutibacterium acnes*.<sup>81</sup>

*Demodex spp.*, specifically *D. folliculorum* and *D. brevis*, are commensal mites that dwell within the pilosebaceous follicle and consume sebum.<sup>82</sup> The highest concentrations of *Demodex spp.* have been found in areas typically affected by rosacea (ie, cheeks, forehead).<sup>83</sup> A 5-year epidemiological study of 3213 patients found PPR was associated with a mean *Demodex* density (Dd) of 36 D/cm<sup>2</sup> (range: 8-112 D/cm<sup>2</sup>) compared with  $\leq 5$  D/cm<sup>2</sup> in normal skin.<sup>84</sup> Non-invasive skin samples have found *Demodex spp.* more frequently in individuals with rosacea vs healthy age-matched controls (96% vs 74%,  $P<.01$ ); in one study the Dd was  $\geq 5.7$ x in rosacea patients.<sup>82</sup> This appears to be unique to rosacea as other inflammatory conditions of the face failed to show excess *Demodex* presence.<sup>83</sup> Clinicohistological studies have found mixed results but suggest that heavy *Demodex spp.* burdens may cause follicular rupture, dermal invasion, and cell-mediated reaction with granuloma formation along with activation of a plurality of inflammatory signals (including matrix metalloproteinase-9 (MMP-9), interleukin-8 (IL-8), tumor necrosis factor alpha (TNF-alpha), and LL-37).<sup>82,85</sup> However, it is unclear if *Demodex spp.* instigates inflammation in rosacea or that underlying inflammatory changes enable proliferation of *Demodex spp.*<sup>86</sup>

*Bacillus oleronius* is a non-commensal gram-negative bacteria that has been cultivated from *Demodex* mite in PPR patients and is capable of stimulating a mononuclear cell response more readily in rosacea patients (n=16/22, 73%) vs healthy controls (n=5/17, 29%,  $P=.01$ ).<sup>87</sup> 62 and 83 kilodalton (kDa) antigens isolated from *B. oleronius* have been found to preferentially stimulate serum reactivity in ETR individuals (n=21/26, 80%) over controls (n=9/22, 40%;  $P=.004$ ) as well as in individuals with ocular involvement.<sup>88,89</sup> *Demodex spp.* count was noted to be significantly greater in rosacea patients (6.6 $\pm$ 9.0 vs 1.9 $\pm$ 2.2,  $P=.014$ ), suggesting *Demodex* may be a vector for *B. oleronius*-induced inflammation.<sup>89</sup> Inflammatory cascades downstream of *B. oleronius* include many factors associated with *Demodex*-associated inflammation, including MMP-9, IL-8, TNF-alpha.<sup>90</sup>

*S. epidermidis* is a gram-positive coccus that, although capable of mitigating pathogenic *S. aureus* strain activity,<sup>91,92</sup> has capacity for virulence.<sup>93,94</sup> Cultures taken from PPR patients have found *S. epidermidis* within pustules but not from unaffected ipsilateral cheek skin.<sup>95</sup> Furthermore, studies comparing *S. epidermidis* strains isolated from untreated PPR patients with controls found

*S. epidermidis* from PPR pustules were significantly more likely to have  $\beta$ -hemolytic activity than from control isolates and also to secrete different (potentially virulent) proteins at higher temperatures (ie, 37°C vs 30°C).<sup>96</sup> Such temperatures are not uncommon in the vasodilated cheeks of a rosacea patients.

The lack of *C. acnes* may also play a role in rosacea.<sup>97,98</sup> A small study (n=58) comparing superficial cutaneous swabs from individuals with PPR (n=15), ETR (n=21) with controls (n=22) found significantly less *C. acnes* isolated from the bilateral malar cheeks in rosacea patients (ETR: 27.3%; PPR: 23.3%) compared with controls (62.6%,  $P<.01$ ).<sup>98</sup> Another study found that healthy individuals age  $\geq 60$  have significantly less ( $P=.018$ ) *C. acnes* than those  $<60$ .<sup>97</sup> These observations may explain why AV is more prevalent among teens while another unrelated facial dermatosis, rosacea, is more common in those in the fourth/fifth decade of life.

The authors also note that there is growing evidence of “cross-talk” between the enteric microbiome and CM in rosacea patients.<sup>99-105</sup> Population studies have found individuals with rosacea were more likely to have celiac disease (Hazard Ratio 1.46, 95%CI 1.11-1.93,  $P<.001$ ), Crohn’s disease (HR 1.45, 95%CI 1.19-1.77,  $P<.001$ ), ulcerative colitis (HR 1.19, 95%CI 1.02-1.39,  $P<.001$ ), and irritable bowel syndrome (IBS) (HR 1.34, 95%CI 1.19-1.50,  $P<.001$ ).<sup>100</sup> A randomized-control trial using rifaximin (a non-systemic, intestinal-limited antibiotic) for the management of small intestine bacterial overgrowth (SIBO) found that rosacea lesions completely cleared (n=20/28) or greatly improved (n=6/28) after treatment with rifaximin compared with those who received placebo (no change, n=18/20; worsened, n=2/20;  $P<.001$ ).<sup>102</sup> Other studies have also implicated alterations in the relative abundance of different species in rosacea including those of the genera *Gordonia* and *Geobacillus*, *Corynebacterium*, *Actinomyces*, *Vellonella*, and *Chloroplast*.<sup>103-105</sup>

Although these findings may imply a correlation between disease and dysbiosis, they cannot yet establish causality as it is unclear if dysbiosis is a primer or symptom of inflammation.<sup>104</sup> Furthermore, studies have not consistently identified a single organism tying dysbiosis to rosacea, suggesting that dysbiosis itself may be sufficient to stimulate inflammation.<sup>103,104</sup>

#### Current Therapeutic Regimens and the Microbiome *Acne Vulgaris*

Topical retinoids and benzoyl peroxide (BPO) are mainstays of mild-moderate AV management.<sup>106</sup> BPO has been found to increase TEWL and deplete SC levels of  $\alpha$ -tocopherol.<sup>107</sup> Topical retinoids have been found to transiently thin the SC, increase cell turnover, and increase TEWL.<sup>18</sup> Clinically, patients perceive these effects as irritation, inflammation, and xerosis, which may be partially alleviated by concurrent application of a gentle, non-comedogenic moisturizer.<sup>19-21</sup> Topical treatment may also influence the microbiome (at least in the acute [post-]

treatment period) by creating an inhospitable environment for some microbes and a boon for others, with yet unclear long-term post-treatment implications.<sup>108,109</sup> Despite this, it should be noted these medications are all capable of (at least transiently) improving acne severity.

Systemic antibiotic therapy is the cornerstone for moderate-severe AV therapy.<sup>106</sup> While broad-spectrum antibiotics may achieve transient improvements, they may also induce long-term CM alterations. In one longitudinal prospective study, 4 women ages 25 to 32 with recently diagnosed AV were given oral minocycline 100 mg twice daily for 4 weeks. Superficial cutaneous swabs found a 1.4-fold reduction in *C. acnes* counts with a trend towards *C. acnes* abundance recovery 8 weeks after minocycline discontinuation, but a sustained reduction in *Lactobacillus spp.* and *Corynebacterium spp.* over the same time interval.<sup>110</sup>

Isotretinoin is one of the few prescription AV therapies that consistently achieves durable response.<sup>106,111</sup> Several studies suggest this may be due in part to modulations of the microbiome.<sup>64-66,109</sup> CM samples post-isotretinoin therapy have found up to a 100-fold decrease in *C. acnes* colonies (including strains resistant to erythromycin, clindamycin, and tetracycline) 1 month after completion of 18-week isotretinoin course<sup>65</sup> and increased diversity of other taxa.<sup>66</sup> This CM recalibration may be a result of an isotretinoin-induced sebaceous drought that induces a microbial survival “bottleneck,” which allows non-virulent *C. acnes* strains to repopulate the follicles.<sup>64</sup>

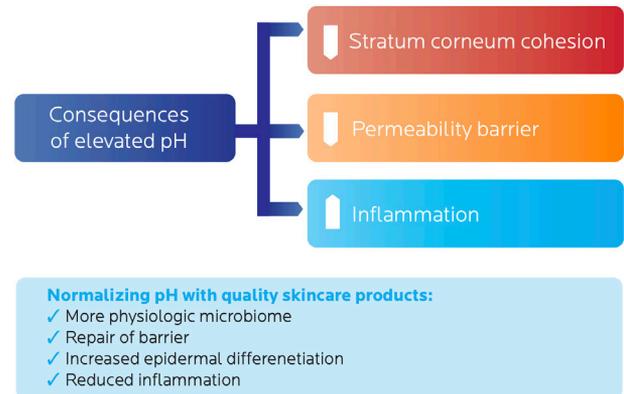
#### Rosacea

Systemic antimicrobials have demonstrated efficacy in management of rosacea, primarily PPR.<sup>112,113</sup> In a longitudinal cohort study of 12 PPR patients treated with doxycycline, superficial skin swabs found a significant 3.43-fold increase in *Weissella confusa* relative abundance ( $P=0.008$ )<sup>97</sup> and also a material change in the predominant genera with a baseline composition of *Staphylococcus* (28%), *Cutibacterium* (13%), *Pseudomonas* (9%), *Corynebacterium* (8%), *Acinetobacter* (7%), and *Snodgrassella* (6%) being replaced by *Staphylococcus* (22%), *Stenotrophomonas* (33%), *Corynebacterium* (8%) and *Cutibacterium* (7%) after 6 weeks of doxycycline 100 mg twice-daily.<sup>97</sup> The significance, if any, of this microbiome shift is unknown.

#### Present Practices and Potential Paradigms

Dermatologists may consider the nature between EBD and dysbiosis akin to an interdependent positive feedback loop. However, clinically, the directional relationship between dysbiosis and EBD may be irrelevant if patients are counseled on therapeutic options with the potential to improve both anomalies. The authors note that skincare and OTC products, including cleansers and moisturizers, are a grossly underap-

**FIGURE 4.** Relationship between pH, barrier (dys)function, and the microbiome. Adapted from Lynde CW et al. *J Clin Aesthet Dermatol.* 2014;7(3):40-48 and *Proksch J Dermatol.* 2018;45:1044-1052.



preciated and overlooked cornerstone in patient education that may complement prescription therapy.

Quality moisturizers are composed of: humectants (eg, glycerin, hyaluronic acid) to attract water; occlusives that seal in moisture (eg, petrolatum); and emollients that soften and smoothen the skin.<sup>70</sup> Effective implementation can maintain a healthy epidermal barrier, and even rescue a deficient one (Figure 4).<sup>70,115,116</sup> For AV patients, moisturizers should also be non-comedogenic to avoid instigating additional lesions.<sup>20,21,117</sup> Individuals with AV may also benefit from using moisturizers with ceramides to supplement their prescription regimens.<sup>23,70,117-118</sup> Given the increased skin sensitivity seen in rosacea, moisturizers should avoid fragrances, surfactants and other potential instigators of allergic or irritant contact dermatitis, and have an acidic or physiologic pH.<sup>70</sup>

An ideal cleanser should remove debris, cosmetics, and transient bacteria without perturbing EBF or CM.<sup>70</sup> For both AV and rosacea, cleansers should be close to physiologic skin pH (~4-6) to avoid excessive burning, stinging, and dryness and to preserve the metabolic functions, including lipid processing (Figure 4).<sup>70</sup> Lipid-free cleansers and synthetic detergents allow for superior preservation of the skin’s natural lipids thereby decreasing irritancy.<sup>70</sup>

#### Therapeutic Probiotics

In addition to these essential qualities of skincare regimens, the authors note the clinical potential to augment current skincare products by utilizing pro-/pre-/postbiotics to further mend EBD and restore microbiome diversity and richness.

Probiotics are foods (eg, yogurts, fermented products such as kefir and kombucha) and topical/oral supplements that contain live microorganisms such as *Nitrosomonas eutropha*, *Lactobacillus spp.*, *Lactococcus spp.*, *Streptococcus spp.*, and

TABLE 3.

Epidermal Barrier Outcomes Following Topical and Systemic Management of Rosacea							
Study	RoA	Study Type & Size (n)	Acne Location, Severity	Participant Age	Study Length	Treatment Regimen	Results
Kang et al <sup>123</sup>	Topical	RCT (70)	Face, mild-moderate	≥12	8 weeks	Fecal <i>E. faecalis</i> anti-bacterial isolate lotion	Significant ( $P<.05$ ) reduction in inflammatory lesions (pustules) vs vehicle
Muizzuddin et al <sup>124</sup>	Topical	Open-label (10)	Face & Trunk, --	Range 18-50	4 days	<i>Lactobacillus spp.</i> 5% lotion spot application QD	Significant ( $P<.05$ ) reduction in acne lesion size and erythema compared with untreated lesions
AOBiome <sup>125</sup>	Topical	RCT, clinical trial phase 2b (358) <sup>*</sup>	--, Mild-moderate	--	12 weeks	Topical <i>Nitrosomonas eutropha</i> -containing spray	Significantly ( $P=.03$ ) more participants had 2-grade IGA reduction and greater reduction in inflammatory lesions ( $P=.028$ ) than vehicle
Jung et al <sup>126</sup>	Oral	RCT (45)	Face, mild-moderate	Range 18-35	12 weeks	BPO 5% and facial cleanser and: A) Minocycline QD or B) <i>Lactobacillus spp.</i> & <i>Bifidobacterium bifidum</i> BID or C) Both A&B	C had significantly greater reduction in total lesion count than group A (week 8: 67% vs 52%, $P<.001$ ; week 12: 82% vs 67%, $P<.001$ ) and B (week 8: 67% vs 56%, $P=.006$ ; week 12: 82% vs 67%, $P<.001$ ) C also had significantly less NIL and IL count at week 12 than A (NIL: 77% vs 70%, $P=.03$ ; IL: 77% vs 70%, $P<.001$ ) and B (NIL: 77% vs 65%, $P=.001$ ; IL: 77% vs 70% $P<.001$ )
Fabbrocini et al <sup>127</sup>	Oral	RCT (20)	Trunk, --	Mean (SD): 33.7±3.3	12 weeks	<i>Lactobacillus rhamnosus</i> liquid	Significant decrease 32% ( $P<.001$ ) in IGF-1 and 65% ( $P<.001$ ) increase in FOXO1 from biopsied acne lesions Participants taking probiotic were more likely to be rated as having (markedly) improved based on IGA (aOR 28.4, 95%CI 2.2-411.1, $P<.05$ )

-- Data not available from source document. \*ClinicalTrials.gov ID: NCT02832063. <sup>a</sup>OR, adjusted odds ratio; BID, twice daily; CI, confidence interval; IGA, investigator's global assessment; IL, inflammatory lesion; NIL, non-inflammatory lesion; QD, once daily; RCT, randomized controlled trial; RoA, route of administration; *spp.*, species.

*Bifidobacteria spp.*<sup>119-120</sup> Theoretically, introduction of these species may mitigate and outcompete pathogenic organisms,<sup>121</sup> thereby improving CM homeostasis.

Several clinical trials have found success in using probiotics as monotherapy or in conjunction with traditional AV therapy (Table 3).<sup>119,122-127</sup> Studies are investigating "grafting" autologous microbes cultured from non-lesional skin in individuals with dermatoses onto affected areas.<sup>128</sup> In the future, we may be able to extract the contents of an AV-prone follicle, apply a cocktail of *C. acnes* (IA1)-specific bacteriophage to deplete pathogenic strains and reintroduce commensal *C. acnes* phylotypes II and III.<sup>42,62</sup> This dual-pronged approach may provide a personalized, targeted means to improve disease severity while promoting a healthy CM.<sup>129</sup>

#### More than Moisture: Water, Prebiotics, and Postbiotics

Prebiotics are topical/oral supplements or foods that selectively support and stimulate the growth and/or activity of the microbiome.

Different microorganisms have been found to tolerate decreased SCH, such as *S. aureus*, whose abundance correlates temporally with AD flares,<sup>130</sup> and therefore may outcompete commensal microbes under more "arid" conditions.<sup>12,131</sup> Therefore adequate moisturization and water-retention may not only promote EBF but also microbiome diversity. In this way water, by cultivating

and stimulating CM growth and/or activity, may be considered a prebiotic.<sup>132</sup>

Postbiotics (or bacterial byproducts, metabolites, and excreted compounds from lysed organisms, such as *Xanthomonas*, and *Enterococcus faecalis*) may also have potential to affect dermatoses. An 8-week double-blind, randomized, vehicle-controlled trial of 70 mild-moderate AV patients found that topically-applied anti-bacterial isolate from fecal *Enterococcus faecalis* significantly ( $P<.05$ ) reduced more pustules than the vehicle.<sup>123</sup> Postbiotics from *Vitreoscilla filiformis*, a gram-negative filamentous, non-pathogenic bacterium that is naturally found in thermal springs and spa water<sup>133</sup> has been found to have antioxidant abilities mediated via mitochondrial superoxide dismutase in vitro<sup>134</sup> and anti-inflammatory activity via interleukin-10 (IL-10) mediated regulatory T cells within murine models of AD.<sup>135</sup> Preliminary data also suggest that addition of *V. filiformis* lysate to emollients may provide a method to improve EBF and CM diversity.<sup>136-138</sup>

#### DISCUSSION AND FUTURE DIRECTIONS

While the flow of causality between EBD and dysbiosis is not yet clear for AV and rosacea, they do appear to play a material role in the pathogenesis and development of clinical symptoms. Repairing the barrier and restoring the microbiome are essential, and quality skin care may help patients achieve this goal. It is vital that dermatologists are aware of the growing

role that EBD and dysbiosis may play in AV and rosacea, and how best to select prescription and OTC agents to address these deficiencies.

Currently, there are limited clinical studies investigating the use of adjunctive skincare products, especially in AV and rosacea.

Studies investigating the role of pro-/pre-/postbiotics have traditionally focused on AD (and psoriasis).

Future studies, and specifically clinical trials, should assess the implementation of skincare products in AV and rosacea, with particular attention afforded to pro-/pre-/postbiotics and how they may complement current prescription regimens to longitudinally augment clinical improvement and minimize adverse effects and local skin irritation.

## CONCLUSION

AV and rosacea are multifaceted inflammatory dermatoses that both have varying degrees of intrinsic EBD, which may be affected by selected therapy. Similarly, AV and rosacea have unique CM abnormalities that can be further perturbed by indiscriminate use of systemic broad-spectrum antibiotics. The interplay between EBF and the CM is critical for the research and development of new therapies and methods to approach these diseases. Appropriately developed prescription and OTC agents have the potential to selectively influence the microbiome in a beneficial direction and maintain, if not restore, EBF. By understanding this intricate interplay, dermatologists will be better able to educate their patients on the importance of appropriate skin care and potentially improve their quality of life.

## DISCLOSURES

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## REFERENCES

- Foster KR, Schluter J, Coyte KZ, Rakoff-Nahoum S. The evolution of the host microbiome as an ecosystem on a leash. *Nature*. 2017;548(7665):43-51. doi:10.1038/nature23292. PMID: 28770836; PMCID: PMC5749636.
- NIH HMP Working Group, Peterson J, Garges S, et al. The NIH human microbiome project. *Genome Res*. 2009;19(12):2317-2323. doi:10.1101/gr.096651.109
- Prescott SL, Larcombe DL, Logan AC, et al. The skin microbiome: impact of modern environments on skin ecology, barrier integrity, and systemic immune programming. *World Allergy Organ J*. 2017;10(1):29. doi:10.1186/s40413-017-0160-5
- Kinross JM, Darzi AW, Nicholson JK. Gut microbiome-host interactions in health and disease. *Genome Med*. 2011;3(3):14. doi:10.1186/gm228
- Lee YB, Byun EJ, Kim HS. Potential role of the microbiome in acne: a comprehensive review. *J Clin Med*. 2019;8(7):987. doi:10.3390/jcm8070987
- Wilson M. The Human Microbiota in Health And Disease: An Ecological And Community-Based Approach. *Boca Raton: Garland Science*. 2018.
- Cogen AL, Nizet V, Gallo RL. Skin microbiota: a source of disease or defence? *Br J Dermatol*. 2008;158(3):442-455. doi:10.1111/j.1365-2133.2008.08437.x
- Sanford JA, Gallo RL. Functions of the skin microbiota in health and disease. *Semin Immunol*. 2013;25(5):370-377. doi:10.1016/j.smim.2013.09.005
- Gallo RL, Nakatsuji T. Microbial symbiosis with the innate immune defense system of the skin. *J Invest Dermatol*. 2011;131(10):1974-1980. doi:10.1038/jid.2011.182
- Wiesner J, Vilcinskas A. Antimicrobial peptides: the ancient arm of the human immune system. *Virulence*. 2010;1(5):440-464. doi:10.4161/viru.1.5.12983
- Strugar TL, Kuo A, Seité S, et al. Connecting the dots: from skin barrier dysfunction to allergic sensitization, and the role of moisturizers in repairing the skin barrier. *J Drugs Dermatol*. 2019;18(6):581.
- Baldwin HE, Bhatia ND, Friedman A, et al. The role of cutaneous microbiota harmony in maintaining a functional skin barrier. *J Drugs Dermatol*. 2017;16(1):12-18.
- Baldwin H, Aguh C, Andriessen A, et al. Atopic dermatitis and the role of the skin microbiome in choosing prevention, treatment, and maintenance options. *J Drugs Dermatol*. 2020;19(10):935-940. doi:10.36849/JDD.2020.5393
- Barnard E, Shi B, Kang D, et al. The balance of metagenomic elements shapes the skin microbiome in acne and health. [published correction appears in *Sci Rep*. 2020 Apr 2;10(1):6037]. *Sci Rep*. 2016;6:39491. doi:10.1038/srep39491
- Naik S, Bouladoux N, Wilhelm C, et al. Compartmentalized control of skin immunity by resident commensals. *Science*. 2012;337(6098):1115-1119. doi:10.1126/science.1225152
- Rocha MA, Bagatin E. Skin barrier and microbiome in acne. *Arch Dermatol Res*. 2018;310(3):181-185. doi:10.1007/s00403-017-1795-3
- Misery L, Loser K, Ständer S. Sensitive skin. *J Eur Acad Dermatol Venereol*. 2016;30(Suppl 1):2-8. doi:10.1111/jdv.13532
- Thiboutot D, Del Rosso JQ. Acne vulgaris and the epidermal barrier: is acne vulgaris associated with inherent epidermal abnormalities that cause impairment of barrier functions? Do any topical acne therapies alter the structural and/or functional integrity of the epidermal barrier? *J Clin Aesthet Dermatol*. 2013;6(2):18-24.
- Tan J, Alexis A, Baldwin H, et al. The personalised acne care pathway-recommendations to guide longitudinal management from the personalising acne: consensus of experts. *JAAD Int*. 2021;5:101-111. doi:10.1016/j.jdin.2021.09.006
- Chularojanamontri L, Tuchinda P, Kulthanan K, Pongparit K. Moisturizers for acne: what are their constituents? *J Clin Aesthet Dermatol*. 2014;7(5):36-44.
- Alexis AF, Woolery-Lloyd H, Williams K, et al. Racial/ethnic variations in acne: implications for treatment and skin care recommendations for acne patients with skin of color. *J Drugs Dermatol*. 2021;20(7):716-725. doi:10.36849/JDD.6169
- Yamamoto A, Takenouchi K, Ito M. Impaired water barrier function in acne vulgaris. *Arch Dermatol Res*. 1995;287(2):214-218. doi:10.1007/BF01262335
- Pappas A, Kendall AC, Brownbridge LC, et al. Seasonal changes in epidermal ceramides are linked to impaired barrier function in acne patients. *Exp Dermatol*. 2018;27(8):833-836. doi:10.1111/exd.13499
- Meyer K, Pappas A, Dunn K, et al. Evaluation of seasonal changes in facial skin with and without acne. *J Drugs Dermatol*. 2015;14(6):593-601.
- Pappas A, Dunn K, Cula G.O., et al. Barrier and microbiome changes in the facial skin of children as they approach puberty. Presented at: 26th European Academy of Dermatology and Venerology(EADV) Congress: P006513-17 September 2017
- Borodzicz S, Rudnicka L, Mirowska-Guzel D, Cudnoch-Jedrzejewska A. The role of epidermal sphingolipids in dermatologic diseases. *Lipids Health Dis*. 2016;15:13. doi:10.1186/s12944-016-0178-7
- Dréno B, Pécastaings S, Corvec S, et al. Cutibacterium acnes(propionibacterium acnes) and acne vulgaris: a brief look at the latest updates. *J Eur Acad Dermatol Venereol*. 2018;32(Suppl 2):5-14. doi:10.1111/jdv.15043
- Dréno B, Martin R, Moyal D, et al. Skin microbiome and acne vulgaris: staphylococcus, a new actor in acne. *Exp Dermatol*. 2017;26(9):798-803. doi:10.1111/exd.13296
- Fitz-Gibbon S, Tomida S, Chiu BH, et al. Propionibacterium acnes strain populations in the human skin microbiome associated with acne. *J Invest Dermatol*. 2013;133(9):2152-2160. doi:10.1038/jid.2013.21
- Gollnick H. Current concepts of the pathogenesis of acne: implications for drug treatment. *Drugs*. 2003;63(15):1579-1596. doi:10.2165/00003495-200363150-00005

31. Kim J. Review of the innate immune response in acne vulgaris: activation of Toll-like receptor 2 in acne triggers inflammatory cytokine responses. *Dermatology*. 2005;211(3):193-198. doi:10.1159/000087011
32. Kurokawa I, Danby FV, Ju Q, et al. New developments in our understanding of acne pathogenesis and treatment. *Exp Dermatol*. 2009;18(10):821-832. doi:10.1111/j.1600-0625.2009.00890.x
33. Cundell AM. Microbial ecology of the human skin. *Microb Ecol*. 2018;76(1):113-120. doi:10.1007/s00248-016-0789-6
34. Sørensen K, Aksglaede L, Petersen JH, Juul A. Recent changes in pubertal timing in healthy Danish boys: associations with body mass index. *J Clin Endocrinol Metab*. 2010;95(1):263-270. doi:10.1210/jc.2009-1478
35. Friedlander SF, Eichenfield LF, Fowler JF Jr, et al. Acne epidemiology and pathophysiology. *Semin Cutan Med Surg*. 2010;29(2 Suppl 1):2-4. doi:10.1016/j.sder.2010.04.002
36. Cunliffe WJ, Gould DJ. Prevalence of facial acne vulgaris in late adolescence and in adults. *Br Med J*. 1979;1(6171):1109-1110. doi:10.1136/bmj.1.6171.1109
37. Jugé R, Rouaud-Tinguely P, Breugnot J, et al. Shift in skin microbiota of western European women across aging. *J Appl Microbiol*. 2018;125(3):907-916. doi:10.1111/jam.13929
38. Shibagaki N, Suda W, Clavaud C, et al. Aging-related changes in the diversity of women's skin microbiomes associated with oral bacteria. *Sci Rep*. 2017;7(1):10567. Published 2017 Sep 5. doi:10.1038/s41598-017-10834-9
39. Dagnelie MA, Corvec S, Saint-Jean M, et al. Cutibacterium acnes phylotypes diversity loss: a trigger for skin inflammatory process. *J Eur Acad Dermatol Venereol*. 2019;33(12):2340-2348. doi:10.1111/jdvs15795
40. Beylot C, Auffret N, Poli F, et al. Propionibacterium acnes: an update on its role in the pathogenesis of acne. *J Eur Acad Dermatol Venereol*. 2014;28(3):271-278. doi:10.1111/jdvs12224
41. Oyewole AO, Birch-Machin MA. Sebum, inflammasomes and the skin: current concepts and future perspective. *Exp Dermatol*. 2015;24(9):651-654. doi:10.1111/exd.12774
42. Paugam C, Corvec S, Saint-Jean M, et al. Propionibacterium acnes phylotypes and acne severity: an observational prospective study. *J Eur Acad Dermatol Venereol*. 2017;31(9):e398-e399. doi:10.1111/jdvs14206
43. Dagnelie MA, Corvec S, Saint-Jean M, et al. Decrease in diversity of propionibacterium acnes phylotypes in patients with severe acne on the back. *Acta Derm Venereol*. 2018;98(2):262-267. doi:10.2340/00015555-2847
44. YuY, Champer J, Agak GV, et al. Different propionibacterium acnes phylotypes induce distinct immune responses and express unique surface and secreted proteomes. *J Invest Dermatol*. 2016;136(11):2221-2228. doi:10.1016/j.jid.2016.06.615
45. Higaki S, Kitagawa T, Kagoura M, et al. Correlation between Propionibacterium acnes biotypes, lipase activity and rash degree in acne patients. *J Dermatol*. 2000;27(8):519-522. doi:10.1111/j.1346-8138.2000.tb02219.x
46. Nazipi S, Stødkilde-Jørgensen K, Scavenius C, Brüggemann H. The skin bacterium propionibacterium acnes employs two variants of hyaluronate lyase with distinct properties. *Microorganisms*. 2017;5(3):57. doi:10.3390/microorganisms5030057
47. Nakatsuji T, Tang DC, Zhang L, Gallo RL, Huang CM. Propionibacterium acnes CAMP factor and host acid sphingomyelinase contribute to bacterial virulence: potential targets for inflammatory acne treatment. *PLoS One*. 2011;6(4):e14797. doi:10.1371/journal.pone.0014797
48. Valanne S, McDowell A, Ramage G, et al. CAMP factor homologues in Propionibacterium acnes: a new protein family differentially expressed by types I and II. *Microbiology (Reading)*. 2005;151(Pt 5):1369-1379. doi:10.1099/mic.0.27788-0
49. Lheure C, Grange PA, Ollagnier G, et al. TLR-2 Recognizes Propionibacterium acnes camp factor 1 from highly inflammatory strains. *PLoS One*. 2016;11(11):e0167237. doi:10.1371/journal.pone.0167237
50. Bek-Thomsen M, Lomholt HB, Scavenius C, et al. Proteome analysis of human sebaceous follicle infundibula extracted from healthy and acne-affected skin. *PLoS One*. 2014;9(9):e107908. doi:10.1371/journal.pone.0107908
51. Sørensen M, Mak TN, Hurwitz R, et al. Mutagenesis of Propionibacterium acnes and analysis of two CAMP factor knock-out mutants. *J Microbiol Methods*. 2010;83(2):211-216. doi:10.1016/j.mimet.2010.09.008
52. Liu PF, Nakatsuji T, Zhu W, et al. Passive immunoprotection targeting a secreted CAMP factor of Propionibacterium acnes as a novel immunotherapeutic for acne vulgaris. *Vaccine*. 2011;29(17):3230-3238. doi:10.1016/j.vaccine.2011.02.036
53. Shu M, Kuo S, Wang Y, et al. Porphyrin metabolites in human skin commensal Propionibacterium acnes bacteria: potential application to monitor human radiation risk. *Curr Med Chem*. 2013;20(4):562-568. doi:10.2174/0929867311320040007
54. Johnson T, Kang D, Barnard E, Li H. Strain-level differences in porphyrin production and regulation in propionibacterium acnes elucidate disease associations. *mSphere*. 2016;1(1):e00023-15. doi:10.1128/mSphere.00023-15
55. Brüggemann H, Lomholt HB, Tettelin H, Kilian M. CRISPR/cas loci of type II Propionibacterium acnes confer immunity against acquisition of mobile elements present in type I P. acnes. *PLoS One*. 2012;7(3):e34171. doi:10.1371/journal.pone.0034171
56. McDowell A, Barnard E, Nagy I, et al. An expanded multilocus sequence typing scheme for propionibacterium acnes: investigation of 'pathogenic', 'commensal' and antibiotic resistant strains. *PLoS One*. 2012;7(7):e41480. doi:10.1371/journal.pone.0041480
57. Lomholt HB, Kilian M. Clonality and anatomic distribution on the skin of antibiotic resistant and sensitive Propionibacterium acnes. *Acta Derm Venereol*. 2014;94(5):534-538. doi:10.2340/00015555-1794
58. Thompson KG, Rainer BM, Antonescu C, et al. Comparison of the skin microbiota in acne and rosacea. *Exp Dermatol*. 2021;30(10):1375-1380. doi:10.1111/exd.14098
59. Wang Y, Kuo S, Shu M, et al. Staphylococcus epidermidis in the human skin microbiome mediates fermentation to inhibit the growth of Propionibacterium acnes: implications of probiotics in acne vulgaris. *Appl Microbiol Biotechnol*. 2014;98(1):411-424. doi:10.1007/s00253-013-5394-8
60. Marinelli LJ, Fitz-Gibbon S, Hayes C, et al. Propionibacterium acnes bacteriophages display limited genetic diversity and broad killing activity against bacterial skin isolates. *mBio*. 2012;3(5):e00279-12. doi:10.1128/mBio.00279-12
61. Brown TL, Petrovski S, Dyson ZA, et al. The formulation of bacteriophage in a semi solid preparation for control of propionibacterium acnes growth. *PLoS One*. 2016;11(3):e0151184. doi:10.1371/journal.pone.0151184
62. Liu J, Yan R, Zhong Q, et al. The diversity and host interactions of Propionibacterium acnes bacteriophages on human skin. *ISME J*. 2015;9(9):2116. doi:10.1038/ismej.2015.144
63. Jorczyk-Matysiak E, Weber-Dąbrowska B, Zaczek M, et al. Prospects of phage application in the treatment of acne caused by propionibacterium acnes. *Front Microbiol*. 2017;8:164. doi:10.3389/fmicb.2017.00164
64. McCoy WH 4th, Otchere E, Rosa BA, Martin J, Mann CM, Mitreva M. Skin ecology during sebaceous drought – how skin microbes respond to isotretinoin. *J Invest Dermatol*. 2019;139(3):732-735. doi:10.1016/j.jid.2018.09.023
65. Ryan-Kewley AE, Williams DR, Hepburn N, Dixon RA. Non-antibiotic isotretinoin treatment differentially controls propionibacterium acnes on skin of acne patients. *Front Microbiol*. 2017;8:1381. doi:10.3389/fmicb.2017.01381
66. Kelhällä HL, Aho VTE, Fyhrquist N, et al. Isotretinoin and lymecycline treatments modify the skin microbiota in acne. *Exp Dermatol*. 2018;27(1):30-36. doi:10.1111/exd.13397
67. Crawford GH, Pelle MT, James WD. Rosacea: I. Etiology, pathogenesis, and subtype classification. *J Am Acad Dermatol*. 2004;51(3):327-344. doi:10.1016/j.jaad.2004.03.030
68. Draelos ZD. Facial hygiene and comprehensive management of rosacea. *Cutis*. 2004;73(3):183-187.
69. Corazza M, la Malfa W, Lombardi A, et al. Role of allergic contact dermatitis in rosacea. *Contact Dermatitis*. 1997;37(1):40-41. doi:10.1111/j.1600-0536.1997.tb00379.x
70. Baldwin H, Alexis AF, Andriessen A, et al. Evidence of barrier deficiency in rosacea and the importance of integrating otc skincare products into treatment regimens. *J Drugs Dermatol*. 2021;20(4):384-392. doi:10.36849/JDD.2021.5861
71. Darlenski R, Kazandjieva J, Tsankov N, Fluhr JW. Acute irritant threshold correlates with barrier function, skin hydration and contact hypersensitivity in atopic dermatitis and rosacea. *Exp Dermatol*. 2013;22(11):752-753. doi:10.1111/exd.12251
72. Darlenski R, Kazandjieva J, Fluhr JW, et al. Lactic acid sting test does not differentiate between facial and generalized skin functional impairment in sensitive skin in atopic dermatitis and rosacea. *J Dermatol Sci*. 2014;76(2):151-153. doi:10.1016/j.jdermsci.2014.08.014
73. Dirschka T, Tronnier H, Fölster-Holst R. Epithelial barrier function and atopic diathesis in rosacea and perioral dermatitis. *Br J Dermatol*. 2004;150(6):1136-1141. doi:10.1111/j.1365-2133.2004.05985.x
74. Lonne-Rahm SB, Fischer T, Berg M. Stinging and rosacea. *Acta Derm Venereol*. 1999;79(6):460-461. doi:10.1080/000155599750009915
75. Pelle MT, Crawford GH, James WD. Rosacea: II. Therapy. *J Am Acad Dermatol*. 2004;51(4):499-514. doi:10.1016/j.jaad.2004.03.033
76. Draelos ZD. Cosmetics in acne and rosacea. *Semin Cutan Med Surg*. 2001;20(3):209-214. doi:10.1053/sder.2001.27556
77. Del Rosso JQ. The use of moisturizers as an integral component of topical therapy for rosacea: clinical results based on the assessment of skin characteristics study. *Cutis*. 2009;84(2):72-76.
78. Subramanyan K. Role of mild cleansing in the management of patient skin. *Dermatol Ther*. 2004;17(Suppl 1):26-34. doi:10.1111/j.1396-0296.2004.04s1003.x
79. Draelos ZD, Green BA, Edison BL. An evaluation of a polyhydroxy acid skin care regimen in combination with azelaic acid 15% gel in rosacea patients. *J Cosmet Dermatol*. 2006;5(1):23-29. doi:10.1111/j.1473-2165.2006.000219.x
80. Ni Raghallaigh S, Powell FC. Epidermal hydration levels in patients with rosacea improve after minocycline therapy. *Br J Dermatol*. 2014;171(2):259-266. doi:10.1111/bjd.12770
81. Holmes AD. Potential role of microorganisms in the pathogenesis of rosacea. *J Am Acad Dermatol*. 2013;69(6):1025-1032. doi:10.1016/j.jaad.2013.08.006
82. Casas C, Paul C, Lahfa M, et al. Quantification of demodex folliculorum by pcr in rosacea and its relationship to skin innate immune activation. *Exp Dermatol*. 2012;21(12):906-910. doi:10.1111/exd.12030
83. Roihu T, Kariniemi AL. Demodex mites in acne rosacea. *J Cutan Pathol*. 1998;25(10):550-552. doi:10.1111/j.1600-0560.1998.tb01739.x
84. Forton F, Germaux MA, Brasseur T, et al. Demodicosis and rosacea: epidemiology and significance in daily dermatologic practice. *J Am Acad Dermatol*. 2005;52(1):74-87. doi:10.1016/j.jaad.2004.05.034
85. Bonamigo RR, Bakos L, Edelweiss M, Cartell A. Could matrix metalloproteinase-9 be a link between Demodex folliculorum and rosacea? *J Eur Acad Dermatol Venereol*. 2005;19(5):646-647. doi:10.1111/j.1468-3083.2005.01221.x
86. Forton FMN. The pathogenic role of demodex mites in rosacea: a potential therapeutic target already in erythematotelangiectatic rosacea? *Dermatol Ther (Heidelb)*. 2020;10(6):1229-1253. doi:10.1007/s13555-020-00458-9
87. Lacey N, Delaney S, Kavanagh K, Powell FC. Mite-related bacterial antigens stimulate inflammatory cells in rosacea. *Br J Dermatol*. 2007;157(3):474-481. doi:10.1111/j.1365-2133.2007.08028.x
88. O'Reilly N, Menezes N, Kavanagh K. Positive correlation between serum immunoreactivity to Demodex-associated Bacillus proteins and erythematotelangiectatic rosacea. *Br J Dermatol*. 2012;167(5):1032-1036. doi:10.1111/j.1365-2133.2012.11114.x
89. Li J, O'Reilly N, Sheha H, et al. Correlation between ocular Demodex infestation

- and serum immunoreactivity to Bacillus proteins in patients with Facial rosacea. *Ophthalmology*. 2010;117(5):870-877.e1. doi:10.1016/j.ophtha.2009.09.057
90. O'Reilly N, Bergin D, Reeves EP, McElvaney NG, Kavanagh K. Demodex-associated bacterial proteins induce neutrophil activation. *Br J Dermatol*. 2012;166(4):753-760. doi:10.1111/j.1365-2133.2011.10746.x
  91. Cogen AL, Yamasaki K, Muto J, et al. Staphylococcus epidermidis antimicrobial delta-toxin (phenol-soluble modulin-gamma) cooperates with host antimicrobial peptides to kill group A Streptococcus. *PLoS One*. 2010;5(1):e8557. doi:10.1371/journal.pone.0008557
  92. Bastos MC, Ceotto H, Coelho ML, Nascimento JS. Staphylococcal antimicrobial peptides: relevant properties and potential biotechnological applications. *Curr Pharm Biotechnol*. 2009;10(1):38-61. doi:10.2174/138920109787048580
  93. Dowd SE, Sun Y, Secor PR, et al. Survey of bacterial diversity in chronic wounds using pyrosequencing, DGGE, and full ribosome shotgun sequencing. *BMC Microbiol*. 2008;8:43. doi:10.1186/1471-2180-8-43
  94. Uçkay I, Pittet D, Vaudaux P, et al. Foreign body infections due to Staphylococcus epidermidis. *Ann Med*. 2009;41(2):109-119. doi:10.1080/07853890802337045
  95. Whitfeld M, Gunasingam N, Leow LJ, et al. Staphylococcus epidermidis: a possible role in the pustules of rosacea. *J Am Acad Dermatol*. 2011;64(1):49-52. doi:10.1016/j.jaad.2009.12.036
  96. Dahl MV, Ross AJ, Schlievert PM. Temperature regulates bacterial protein production: possible role in rosacea. *J Am Acad Dermatol*. 2004;50(2):266-272. doi:10.1016/j.jaad.2003.05.005
  97. Woo YR, Lee SH, Cho SH, Lee JD, Kim HS. Characterization and analysis of the skin microbiota in rosacea: impact of systemic antibiotics. *J Clin Med*. 2020;9(1):185. doi:10.3390/jcm9010185
  98. Wang R, Farhat M, Na J, Li R, Wu Y. Bacterial and fungal microbiome characterization in patients with rosacea and healthy controls. *Br J Dermatol*. 2020;183(6):1112-1114. doi:10.1111/bjd.19315
  99. Searle T, Ali FR, Carolides S, Al-Naiimi F. Rosacea and the gastrointestinal system. *Australas J Dermatol*. 2020;61(4):307-311. doi:10.1111/ajd.13401
  100. Egeberg A, Weinstock LB, Thyssen EP, et al. Rosacea and gastrointestinal disorders: a population-based cohort study. *Br J Dermatol*. 2017;176(1):100-106. doi:10.1111/bjd.14930
  101. Kim M, Choi KH, Hwang SW, et al. Inflammatory bowel disease is associated with an increased risk of inflammatory skin diseases: A population-based cross-sectional study. *J Am Acad Dermatol*. 2017;76(1):40-48. doi:10.1016/j.jaad.2016.08.022
  102. Parodi A, Paolino S, Greco A, et al. Small intestinal bacterial overgrowth in rosacea: clinical effectiveness of its eradication. *Clin Gastroenterol Hepatol*. 2008;6(7):759-764. doi:10.1016/j.cgh.2008.02.054
  103. Nam JH, Yun Y, Kim HS, et al. Rosacea and its association with enteral microbiota in Korean females. *Exp Dermatol*. 2018;27(1):37-42. doi:10.1111/exd.13398
  104. Marson J, Berto S, Mouser P, Baldwin H. Association between rosacea, environmental factors, and facial cutaneous dysbiosis: A pilot study from the largest national festival of twins. *SKIN*. 2021;5(5):487-495. <https://doi.org/10.25251/skin.5.5.6>
  105. Zaidi AK, Spaunhurst K, Sprockett D, et al. Characterization of the facial microbiome in twins discordant for rosacea. *Exp Dermatol*. 2018;27(3):295-298. doi:10.1111/exd.13491
  106. Zaenglein AL, Pathy AL, Schlosser BJ, et al. Guidelines of care for the management of acne vulgaris. [Published correction appears in *J Am Acad Dermatol*. 2020 Jun;82(6):1576]. *J Am Acad Dermatol*. 2016;74(5):945-73.e33. doi:10.1016/j.jaad.2015.12.037
  107. Weber SU, Thiele JJ, Han N, et al. Topical alpha-tocotrienol supplementation inhibits lipid peroxidation but fails to mitigate increased transepidermal water loss after benzoyl peroxide treatment of human skin. *Free Radic Biol Med*. 2003;34(2):170-176. doi:10.1016/s0891-5849(02)01875-5
  108. Zhou L, Chen L, Liu X, et al. The influence of benzoyl peroxide on skin microbiota and the epidermal barrier for acne vulgaris. *Dermatol Ther*. 2022;35(3):e15288. doi:10.1111/dth.15288
  109. Lam M, Hu A, Fleming P, Lynde CW. The Impact of Acne Treatment on Skin Bacterial Microbiota: A Systematic Review. *J Cutan Med Surg*. 2022;26(1):93-97. doi:10.1177/12034754211037994
  110. Chien AL, Tsai J, Leung S, et al. Association of systemic antibiotic treatment of acne with skin microbiota characteristics. *JAMA Dermatol*. 2019;155(4):425-434. doi:10.1001/jamadermatol.2018.5221
  111. Marson JW, Baldwin HE. An overview of acne therapy, part 2: Hormonal therapy and isotretinoin. *Dermatol Clin*. 2019;37(2):195-203. doi:10.1016/j.det.2018.12.002
  112. Marson JW, Baldwin HE. Rosacea: a wholistic review and update from pathogenesis to diagnosis and therapy. *Int J Dermatol*. 2020;59(6):e175-e182. doi:10.1111/ijd.14757
  113. Del Rosso JQ, Tangheiti E, Webster G, et al. Update on the management of rosacea from the american acne & rosacea society (AARS). *J Clin Aesthet Dermatol*. 2019;12(6):17-24.
  114. Korting HC, Ponce-Pöschl E, Klövekorn W, Schmötzer G, Arens-Corell M, Braun-Falco O. The influence of the regular use of a soap or an acidic syndet bar on pre-acne. *Infection*. 1995;23(2):89-93. doi:10.1007/BF01833872
  115. van Zureen EJ, Fedorowicz Z, Tan J, et al. Interventions for rosacea based on the phenotype approach: an updated systematic review including GRADE assessments. *Br J Dermatol*. 2019;181(1):65-79. doi:10.1111/bjd.17590
  116. Addor FA. Skin barrier in rosacea. *An Bras Dermatol*. 2016;91(1):59-63. doi:10.1590/abd1806-4841.20163541
  117. Lain E, Andriessen AE. Choosing the Right Partner: Complementing prescription acne medication with over-the-counter cleansers and moisturizers. *J Drugs Dermatol*. 2020;19(11):1069-1075. doi:10.36849/JDD.2020.5536
  118. Zeichner JA, Del Rosso JQ. Multivesicular emulsion ceramide-containing moisturizers: an evaluation of their role in the management of common skin disorders. *J Clin Aesthet Dermatol*. 2016;9(12):26-32.
  119. Marson JW, Baldwin HE. New concepts, concerns, and creations in acne. *Dermatol Clin*. 2019;37(1):1-9. doi:10.1016/j.det.2018.07.002
  120. Dessinoti C, Dreno B. Acne treatments: future trajectories. *Clin Exp Dermatol*. 2020;45(8):955-961. doi:10.1111/ced.14239
  121. Claesen J, Spagnolo JB, Ramos SF, et al. A Cutibacterium acnes antibiotic modulates human skin microbiota composition in hair follicles. *Sci Transl Med*. 2020;12(570):eaay5445. doi:10.1126/scitranslmed.aay5445
  122. Yu Y, Dunaway S, Champer J, et al. Changing our microbiome: probiotics in dermatology. *Br J Dermatol*. 2020;182(1):39-46. doi:10.1111/bjd.18088
  123. Kang BS, Seo JG, Lee GS, et al. Antimicrobial activity of enterococci from Enterococcus faecalis SL-5 against Propionibacterium acnes, the causative agent in acne vulgaris, and its therapeutic effect. *J Microbiol*. 2009;47(1):101-109. doi:10.1007/s12275-008-0179-y
  124. Muizzuddin N, Maher W, Sullivan M, et al. Physiological effect of a probiotic on skin. *J Cosmet Sci*. 2012;63(6):385-395.
  125. AOBiome Therapeutics. AOBiome Therapeutics reports positive efficacy results from phase 2b clinical trial of ammonia oxidizing bacteria (AOB) for the treatment of acne vulgaris. Available at: <https://www.aobiome.com/pressreleases/aobiome-therapeutics-reports-positive-efficacy-results-from-phase-2b-clinical-trial-of-ammonia-oxidizing-bacteria-aob-for-the-treatment-of-acne-vulgaris/#primary>. Published October 19, 2017. Accessed December 7, 2021.
  126. Jung GW, Tse JE, Guiha I, Rao J. Prospective, randomized, open-label trial comparing the safety, efficacy, and tolerability of an acne treatment regimen with and without a probiotic supplement and minocycline in subjects with mild to moderate acne. *J Cutan Med Surg*. 2013;17(2):114-122. doi:10.2310/7750.2012.12026
  127. Fabbrocini G, Bertona M, Picazo Ó, et al. Supplementation with Lactobacillus rhamnosus SP1 normalises skin expression of genes implicated in insulin signalling and improves adult acne. *Benef Microbes*. 2016;7(5):625-630. doi:10.3920/BM2016.0089
  128. Nakatsuji T, Gallo RL, Shafiq F, et al. Use of autologous bacteriotherapy to treat staphylococcus aureus in patients with atopic dermatitis: A randomized double-blind clinical trial. *JAMA Dermatol*. 2021;157(8):978-982. doi:10.1001/jamadermatol.2021.1311
  129. O'Neill AM, Gallo RL. Host-microbiome interactions and recent progress into understanding the biology of acne vulgaris. *Microbiome*. 2018;6(1):177. doi:10.1186/s40168-018-0558-5
  130. Kong HH, Oh J, Deming C, et al. Temporal shifts in the skin microbiome associated with disease flares and treatment in children with atopic dermatitis. *Genome Res*. 2012;22(5):850-859. doi:10.1101/gr.131029.111
  131. Stevenson A, Cray JA, Williams JP, et al. Is there a common water-activity limit for the three domains of life? *ISME J*. 2015;9(6):1333-1351. doi:10.1038/ismej.2014.219
  132. Al-Ghazzawi FH, Tester RF. Impact of prebiotics and probiotics on skin health. *Benef Microbes*. 2014;5(2):99-107. doi:10.3920/BM2013.0040
  133. Nelson DC, Wirsen CO, Jannasch HW. Characterization of large, autotrophic beggiatoa spp. abundant at hydrothermal vents of the guaymas basin. *Appl Environ Microbiol*. 1989;55(11):2909-2917. doi:10.1128/aem.55.11.2909-2917.1989
  134. Mahé YF, Martin R, Aubert L, et al. Induction of the skin endogenous protective mitochondrial MnSOD by Vitreoscilla filiformis extract. *Int J Cosmet Sci*. 2006;28(4):277-287. doi:10.1111/j.1467-2494.2006.00333.x
  135. Volz T, Skabytska Y, Guenova E, et al. Nonpathogenic bacteria alleviating atopic dermatitis inflammation induce IL-10-producing dendritic cells and regulatory Tr1 cells. *J Invest Dermatol*. 2014;134(1):96-104. doi:10.1038/ijd.2013.291
  136. Gueniche A, Knautt B, Schuck E, et al. Effects of nonpathogenic gram-negative bacterium Vitreoscilla filiformis lysate on atopic dermatitis: a prospective, randomized, double-blind, placebo-controlled clinical study. *Br J Dermatol*. 2008;159(6):1357-1363. doi:10.1111/j.1365-2133.2008.08836.x
  137. ISAD 2014, 8th Georg Rajka symposium on atopic dermatitis, Nottingham, 21-23 may 2014. *Br J Dermatol*. 2014;170(6):e1-e55. doi:10.1111/bjd.13064
  138. Seité S, Zelenkova H, Martin R, et al. Using a specific emollient to manage skin microbiome dysbiosis. *J Am Acad Dermatol*. 2016;74(5):AB89. doi:10.1016/j.jaad.2016.02.348

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