### **Supplementary Tables and Figures**

Preventing bacterial adhesion to skin by altering their physicochemical cell surface properties specifically.

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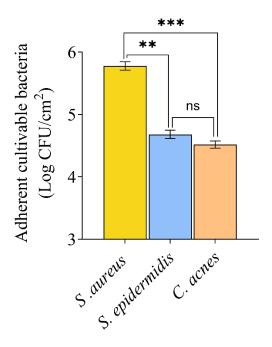
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#Equivalent contribution

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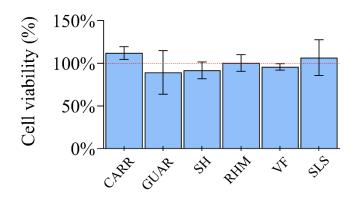


# Supplementary Figure 1: Native adhesion of wild-type skin bacteria after 2 h of incubation on the 3D skin model.

Results are expressed in *Log10* of the number of bacteria adhered per cm<sup>2</sup> of reconstructed skin of the 3D skin model. Error bars show standard error of the mean (SEM) of independent treatments (n > 3). 3 wild-type *S. aureus* strains and *C. acnes* trains and 2 wild-type *S. epidermidis* strains. Strains used are from clinical origins. Statistical significance was calculated by a multiple comparison of strains using the Tukey's test following one-way ANOVA. (\*) for p < 0.05, (\*\*) for p < 0.01 and (\*\*\*).

Supplementary Table 1: wild-type strains used in Supplementary Figure 1

Bacteria	Id. Strain	Origin	Ref.			
	AD08 <sub>cc1</sub>	Human Atopic dermatitis	Fleury et al, 2017 <sup>1</sup>			
Staphylococcus aureus	BC1406	Human Atopic dermatitis	This Study			
	BC1407	Human Atopic dermatitis	This Study			
Staphylococcus epidermidis	BC1190	Human normal skin	Landemaine et al, 2023 <sup>2</sup>			
Stuphylococcus epiderinidis	45A6	Human Atopic dermatitis	Landemaine et al, 2023 <sup>2</sup>			
	BC1089	Human normal skin	This Study			
Cutibacterium acnes	BC1083	Human oily skin	This Study			
	BC1077	Human normal skin	This Study			



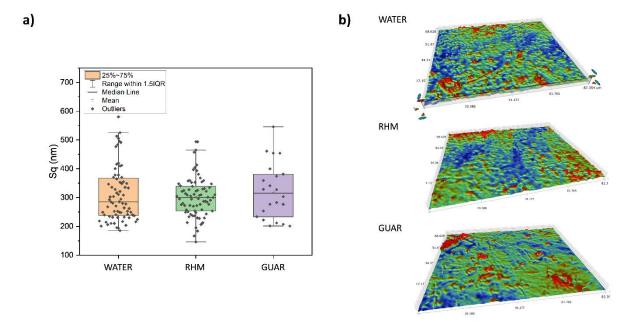
# Supplementary Figure 2: Preservation of 3d skin cell viability after topical compound exposure

MTT assay demonstrating the maintenance of 3D skin model viability following a 2-hour topical treatment of 3D skin cell to the tested compounds. Results are expressed as percentage of viability relative to the water-treated control. SLS: Sodium Laureth Sulfate; CARR: Carrageenan; GUAR: Guar Hydroxypropyltrimonium Chloride; SH: Sodium Hyaluronate; RHM: Rhamnolipids; VF: *Vitreoscilla filiformis* extract. Error bars represent standard deviation (SD) of two independent experiments ( $n \ge 2$ ).

Supplementary Table 2: Contact angle measurement after application of tested compounds using the 3D skin model epidermal surface.

		Cosmetic raw materials										
	Water	CARR	GUAR	SH	RHM	VF	SLS					
Angle contact	34.6 ± 5.9	24.4 ± 2.6	10.4 ± 1.4	19.8 ± 5.7	31.5 ± 1.3	26.1 ± 1.2	8 ± 0.0					

<sup>\*</sup> Angle contacts are expressed in degrees. Values are defined from at least three independent experiments ± standard error of the mean (SEM).



## Supplementary Figure 3: Roughness characterization of the treated 3D skin model using optical profilometry.

a) Box plots representing Sq (root mean square height) values obtained from optical profilometry measurements. Each data point represents a single profilometry image. b) Representative profilometry images (80 x 69  $\mu$ m<sup>2</sup>) for each condition. GUAR: Guar Hydroxypropyltrimonium Chloride; RHM: Rhamnolipids. Error bars represent standard deviation (SD) of a minimum of 22 images per condition, with three biological replicates for RHM and water controls and two biological replicates for GUAR.

Supplementary Table 3: Surface polarity and Lewis-acid balance of treated skin bacteria with tested compounds.

	Physicochemical	Compounds tested																	
Bacteria	parameters*	С	AR	R	G	UA	R		SH		R	HM		,	۷F		8	SLS	
S. aureus	Hydrophobic (%) Lewis acid $\gamma^+$ Lewis base $\gamma^-$	23.4	±	4.0	3.6	±	6.4 1.6 4.2	10.0	±	2.4	39.0	±	1.1	8.5	±	3.0	41.6	±	7.3
C. acnes	Hydrophobic (%) Lewis acid γ <sup>+</sup> Lewis base γ <sup>-</sup>	18.1	±	3.1	13.5	±	10.9 0.2 3.1	8.8	±	1.5	32.1	±	3.8	6.8	±	0.9	20.2	±	1.3

<sup>\*</sup>Parameters are expressed in percent for hydrophobicity and in arbitrary units for Lewis acid/base. Values are defined from at least three independent MATS experiments ± standard error of the mean (SEM).

Supplementary Table 4. Differential RNA expression analysis (≥ 1.5-fold; padj < 0.1) of S. aureus between compounds-treated conditions and water-treated condition.

		log2FoldChange									
Locus tag	Description	CARR	GUAR	SH	RHM	VF					
SAOUHSC_00198	AMP-dependent synthetase and ligase				1.70						
SAOUHSC_00199	Hypothetical unknow protein belongs to the 3-				1.84						
	oxoacid CoA-transferase family										
SAOUHSC_00420	Hypothetical unknow protein belongs to the				1.59						
	sodium neurotransmitter symporter (SNF) (TC										
	2.A.22) family										
SAOUHSC_00465	Biofilm formation stimulator VEG	-2.25									
SAOUHSC_00767	Ribosomal S30AE domain containing protein	-1.71									
SAOUHSC_00786	Hypothetical unknow protein		-1.96								
SAOUHSC_00914	pyruvate carboxyltransferase		1.52								
SAOUHSC_00975	DoxX family transmembrane protein	-2.21									
SAOUHSC_01036	Hypothetical unknow protein belongs to the		-1.70								
	UPF0356 family										
SAOUHSC_01144	Cell division protein	-1.53									
SAOUHSC_01349	Hypothetical unknow protein belongs to the	-1.56									
	HesB/IscA family iron-sulfur cluster assembly										
	accessory protein										
SAOUHSC_01704	Hypothetical unknow protein	-1.66									
SAOUHSC_01764	peptidase A24A, prepilin type IV	-1.67									
SAOUHSC_01782	MutT/nudix hydrolases		-1.56								
SAOUHSC_01902	Hypothetical unknow protein	-1.68									
SAOUHSC_02118	gatC – glutamyl-tRNA(Gln) amidotransferase, C	-1.56	-1.53								
	subunit										
SAOUHSC_02176	Hypothetical unknow protein		-1.90								
SAOUHSC_02286	leuB – 3-isopropylmalate dehydrogenase				1.60						
SAOUHSC_02288	leuD – 3-isopropylmalate dehydratase, small				1.64						
	subunit										
SAOUHSC_02320	Hypothetical unknow protein		3.15								
SAOUHSC_02523	Hypothetical unknow protein	-1.64									
SAOUHSC_02853	Hypothetical unknow protein belongs to the UPF0346 family	-2.09									
SAOUHSC_03017	N-acethyltransferase		-2.24								
SAOUHSC_03045	Cold shock protein				1.82						
SAOUHSC T00012	tRNAasp		-2.98								

#### **Supplementary References**

- Fleury, O. M. *et al.* Clumping Factor B Promotes Adherence of Staphylococcus aureus to Corneocytes in Atopic Dermatitis. *Infect Immun* **85** (2017). <a href="https://doi.org:10.1128/iai.00994-16">https://doi.org:10.1128/iai.00994-16</a>
- 2 Landemaine, L. *et al.* Staphylococcus epidermidis isolates from atopic or healthy skin have opposite effect on skin cells: potential implication of the AHR pathway modulation. *Frontiers in Immunology* **14** (2023). <a href="https://doi.org:10.3389/fimmu.2023.1098160">https://doi.org:10.3389/fimmu.2023.1098160</a>