

## **Electrical Impedance Spectroscopy Quantifies Skin Barrier Function in Organotypic In Vitro Epidermis Models**

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## Abbreviations

ANOVA – analysis of variance

AHR – aryl hydrocarbon receptor

$C_A$ ,  $C_B$  – cellular membrane capacitance

$C_{Cell}$  – cellular capacitance

$C_{EI}$  – electrode capacitance

CLDN1 – claudin 1

CLDN4 – claudin 4

CRISPR/Cas9 – clustered regularly interspaced short palindromic repeats / CRISPR-associated protein 9

EIS – electrical impedance spectroscopy

$EIS^{diff}$  – keratinocyte differentiation-attributable electrical impedance

$EIS^{SC}$  – *stratum corneum*-attributable electrical impedance

H&E – hematoxylin and eosin

HEE – human epidermal equivalents

IL – interleukin

PBS – phosphate-buffered saline

$R_A$ ,  $R_B$  – cellular membrane resistance

$R_{Cell}$  – cellular resistance

$R_{Cyt}$  – cytoplasmatic resistance

$R_{\text{Medium}}$  – culture medium resistance

$R_p$  – paracellular resistance

TEER – transepithelial electrical resistance

TEWL – transepidermal water loss

TFAP2A – transcription factor activating enhancer binding protein 2 alpha

IVL – involucrin

FLG – filaggrin

KRT16 – keratin 16

SKALP – skin-derived antileukoprotease

## ABSTRACT

3 D human epidermal equivalents (HEEs) are a state-of-the-art organotypic culture model in pre-clinical investigative dermatology and regulatory toxicology. Here, we investigated the utility of electrical impedance spectroscopy (EIS) for non-invasive measurement of HEE epidermal barrier function. Our setup comprised a custom-made lid fit with 12 electrode pairs aligned on the standard 24-transwell cell culture system. Serial EIS measurements for seven consecutive days did not impact epidermal morphology and readouts showed comparable trends to HEEs measured only once. We determined two frequency ranges in the resulting impedance spectra: a lower frequency range termed  $EIS^{diff}$  correlated with keratinocyte terminal differentiation independent of epidermal thickness and a higher frequency range termed  $EIS^{SC}$  correlated with *stratum corneum* thickness. HEEs generated from CRISPR/Cas9 engineered keratinocytes that lack key differentiation genes *FLG*, *TFAP2A*, *AHR* or *CLDN1* confirmed that keratinocyte terminal differentiation is the major parameter defining  $EIS^{diff}$ . Exposure to pro-inflammatory psoriasis- or atopic dermatitis-associated cytokine cocktails lowered the expression of keratinocyte differentiation markers and reduced  $EIS^{diff}$ . This cytokine-associated decrease in  $EIS^{diff}$  was normalized after stimulation with therapeutic molecules. In conclusion, EIS provides a non-invasive system to consecutively and quantitatively assess HEE barrier function and to sensitively and objectively measure barrier development, defects and repair.

## INTRODUCTION

Intact physical barriers are of highest importance for our body to define a biophysically enclosed environment. The skin, our largest barrier organ, serves a dual role: it forms an outside–in barrier, protecting the insides of our body from mechanical damage and environmental triggers, and it protects the epidermis and subjacent tissues from dehydration as an inside–out barrier. Barrier functionality is achieved most prominently by a highly organized physical barrier, constituted of tight junctions in the *stratum granulosum* and corneodesmosomes in the *stratum corneum* (Natsuga 2014). *Stratum corneum* corneocytes also are coated with a heavily crosslinked cornified envelope (Évora *et al.* 2021) and the intercellular space between is filled with lipids, generating a hydrophobic environment (van Smeden *et al.* 2014). Next to this physical barrier, the additional chemical, microbial and immunological barriers completes the multifaceted barrier function of mammalian skin (Eyerich *et al.* 2018; Niehues *et al.* 2018).

The importance of the skin barrier is apparent from its malfunction in common skin diseases, like psoriasis and atopic dermatitis. The disease–associated pro–inflammatory milieu also negatively affects keratinocyte differentiation and impairs tight junction and corneodesmosome function (Al Kindi *et al.* 2021; Orsmond *et al.* 2021; Yoshida *et al.* 2022). Next to these multifactorial diseases, monogenic diseases caused by mutations in skin barrier–related genes illustrate the devastating effects of impaired skin barrier function on our health and wellbeing (Hadj-Rabia *et al.*, 2004; Supplemental Table 1 in van den Bogaard *et al.*, 2019). Aside from these intrinsic factors, environmental factors including exhaust fumes or detergents influence the skin barrier function of healthy individuals and patients (Celebi Sözüner *et al.* 2020). Determining the functional consequences of such genetic and environmental risk factors on the skin barrier will aid in our

understanding of disease pathogenesis and may help in the possible future prevention of disease onset or exacerbation.

To investigate skin barrier function, *in vitro* organotypic skin and human epidermal equivalents (HEEs) have become a mainstay approach. By mimicking epidermal barrier morphology and function, HEEs offer advantages over *in vitro* monolayer cultures that lack epidermal stratification and *stratum corneum* formation. In addition, HEEs are considered relevant alternatives to *in vivo* animal models that prompt ethical questions and require depilation to measure biophysical barrier function. HEEs are used from fundamental research to preclinical drug testing to regulatory toxicology in a broad range of applications (El Ghalbzouri *et al.* 2008; Niehues *et al.* 2018).

To assess skin barrier function in HEEs, various technologies can be used ranging from mathematical penetration modelling (Pecoraro *et al.* 2019; Roberts *et al.* 2021) and computational simulation of lipid organization (Shamaprasad *et al.* 2022) to ultrastructural imaging (Riethmüller 2018) and measuring gene and/or protein expression. In a recent consensus paper, we and others have discussed the requirements and methodologies for barrier studies in organotypic skin models (van den Bogaard *et al.*, 2021). In summary, functional barrier assessment using Franz cell diffusion and permeation flux studies provide most accurate estimates of the outside–in barrier (Arkl *et al.* 2018; Neupane *et al.* 2020; Tárnoki-Zách *et al.* 2021). On the other hand, water evaporation (e.g. trans–epidermal water loss (TEWL)) is considered most relevant to describe inside–out barrier function (Alexander *et al.*, 2018; van den Bogaard *et al.*, 2021). Unfortunately, these methods are often labor intensive and rely on highly specific expertise and equipment (Franz cell diffusion assay), are poorly standardized (transepithelial electrical resistance (TEER), TEWL) or require destructive endpoint measurements (permeation studies) (Table S1). Furthermore, the

mechanistic correlation of such measurements to skin barrier function often remains unclear (Yoshida *et al.* 2022).

Electrical impedance spectroscopy (EIS) has been developed and implemented for the assessment of skin barrier function *in vivo* and appears to correlate well with disease severity of atopic dermatitis lesions (Rinaldi *et al.* 2021). For assessment of *in vitro* barrier function, EIS has been implemented for gut, airway and neuroepithelial *in vitro* cultures (Fernandes *et al.* 2022; Wegener *et al.* 2000) and *ex vivo* pig ear skin models (Morin *et al.* 2020). Explorative studies have applied EIS in *in vitro* epidermis models (Groeber *et al.* 2015; Mannweiler *et al.* 2021) and link EIS to viable epidermis and *stratum corneum* barrier properties (Mannweiler *et al.* 2021). Yet, a comprehensive study which extensively assesses EIS applicability and its relation to skin barrier properties in a broad range of experimental models and disease conditions is missing. Here, we demonstrate and validate the use of EIS as a reproducible, non-invasive and quantitative high-throughput system for HEE barrier assessment and assess its correlation to epidermal barrier physiology.

## RESULTS

### **Development of an EIS device for *in vitro* HEE application**

For quantitative and reproducible *in vitro* skin barrier analysis we sought to develop and validate an EIS device for use in HEEs. The system comprises a smart lid with fixed gold-plated electrodes that are customized to fit the individual wells of a Nunc carrier plate with cell culture inserts. The setup enables standardized and automated measurements with a run time of 2 min per well for a maximum of 12 wells (within a 24 well-plate format). To perform the measurements, the smart lid with fixed electrodes is placed onto the HEE carrying culture plate and the connected measurement device (Figure 1a). The electrodes apply a very low alternating voltage  $V$  in a frequency range from 10 Hz to 100 kHz through the culture while measuring the amplitude and phase shift of the resulting alternating current  $I$ . The EIS device returns the impedance  $Z$ , which reflects the opposition to an alternating current over time:

$$\text{Impedance } Z = \frac{V(t)}{I(t)}$$

Impedance and phase spectra are reported in the form of a Bode plot (Figure 1b, c).

For quantitative analysis of EIS spectra, an electrical equivalent circuit model of the examined culture system is required. In conventional 2 D monolayer cultures, there are two main routes of the current: a paracellular, which is determined by the ionic conductance of cell junctions serving as a resistance  $R_P$ , and a transcellular, which consists of the resistance and capacitance of apical and basolateral membrane ( $R_A$ ,  $R_B$  and  $C_A$ ,  $C_B$ ) next to the resistance of the cytoplasm  $R_{Cyt}$  (Benson *et al.* 2013; Srinivasan *et al.* 2015) (Figure 1d). In a simplified model, both membranes can be reduced to  $R_{Cell}$  and  $C_{Cell}$  respectively. In 2 D monolayers, the high cellular resistance  $R_{Cell}$  and the low cytoplasmatic resistance  $R_{Cyt}$  results in paracellular flux being determined by the cellular

capacitance  $C_{\text{Cell}}$  (Benson *et al.* 2013). In 3 D HEEs, multiple individual cell layers result in a parallel series of  $n$  resistor–capacitor electrical circuits (Groeber *et al.* 2015). While we speculate  $R_{\text{P}}$ ,  $R_{\text{Cyt}}$ ,  $R_{\text{Cell}}$  and  $C_{\text{Cell}}$  to be changing depending on cell–cell contacts, differentiation status and the cell shape in the corresponding layer, we assume the dominance of  $C_{\text{Cell}}$  over  $R_{\text{Cell}}$  and  $R_{\text{Cyt}}$  to be persistent in 3 D (Figure 1e). Lastly, the resistance of the medium  $R_{\text{Medium}}$  and the electrodes, acting as pure capacitors with a capacitance  $C_{\text{El}}$ , conclude the electrical circuit.

These electrical circuit elements also determine the generated impedance spectrum (Figure 1f) (Benson *et al.* 2013). Both,  $R_{\text{Medium}}$  and  $C_{\text{El}}$  are fixed parameters whose characteristics are determined by the chosen setup and device. The variable parameters paracellular resistance  $R_{\text{P}}$  and cellular capacitance  $C_{\text{Cell}}$  are determined by the measured cells and the chosen culture system (monolayer vs 3 D organoid). They influence the height and frequency span of a mid–range plateau and the onset of its decline (Yeste *et al.* 2018) (Figure 1g, h). To link EIS to epidermal barrier properties, we determined two frequency ranges in the HEE impedance spectrum approximately indicating  $R_{\text{P}}$  and  $C_{\text{Cell}}$  contribution,  $\text{EIS}^{\text{diff}}$  (127–2212 Hz) and  $\text{EIS}^{\text{SC}}$  (28072–100000 Hz) respectively, which we analyzed by calculating their area under the curve (Benson *et al.* 2013) (Figure 1i).

### **Serial measurements show increased electrical resistance without affecting HEE development**

To determine whether EIS can be used to monitor the development of HEEs non–invasively, serial measurements were performed over consecutive days of immortalized N/TERT–2G keratinocyte air–liquid interface culture. When comparing serial with end point measurements performed directly before harvesting, EIS spectra showed similar trends (Figure 2a, b) and differences in

EIS<sup>diff</sup> and EIS<sup>SC</sup> were not significant (Figure 2c, d). Morphological analysis by hematoxylin and eosin (H&E) staining did not indicate destructive effects of EIS in endpoint or in serial measurements (Figure 2e, top panel). In addition, neither keratinocytes proliferation capacity (Ki67 staining), expression of differentiation proteins (involucrin (IVL) and filaggrin (FLG)), nor the expression of stress-related markers (keratin 16 (KRT16) and skin-derived antileukoprotease (SKALP)) were changed by EIS measurements (Figure 2e). Of note, expression levels of SKALP and the proliferation marker KRT16 are known to be higher in neonatal-derived immortal N/TERT-2G than in primary adult keratinocytes (Smits et al. 2017; Tjabringa et al. 2008). To further evaluate EIS' reliability, HEEs were subjected to EIS measurements six times within one hour (Figure 2f), clearly indicating a very high repeatability. When checking for potentially delayed cytotoxic effects, HEEs harvested 24 hours after repeated EIS measurements showed no morphological signs of cytotoxicity and continuing maturation, as seen by the formation of additional epidermal layers (Figure 2g).

### **Different electrical impedance spectra can be linked to keratinocyte differentiation and *stratum corneum* thickness**

The EIS device measures impedance over a broad range of frequencies which we sought to correlate with epidermal barrier properties. For this, we determined EIS<sup>diff</sup> and EIS<sup>SC</sup> of HEEs during barrier development (Figure 3a-c) and performed correlation analysis with principle epidermal barrier compartments (Figure 3d-g). We observed the thickness of viable epidermal layers to be increasing from day 1-10 before decreasing at day 12 and 14 (Figure S1a, b), which correlated with EIS<sup>diff</sup> but not with EIS<sup>SC</sup> (Figure 3d, e). At the same time *stratum corneum* thickness did not correlate with EIS<sup>diff</sup> but strongly correlated with EIS<sup>SC</sup> explaining 62 % of its

variance (Figure 3f–g). Since keratinocyte terminal differentiation plays a major role in the formation of the skin barrier, we also investigated the protein expression of essential terminal differentiation proteins in relation to  $EIS^{diff}$  (Figure 3j–l).  $EIS^{diff}$  correlated with the quantified expression of keratinocyte differentiation markers FLG and IVL which increased in early days of HEE development before decreasing after maturation at day 14 (Figure 3 h–j). Expression of the tight junction proteins, claudin 1 (CLDN1) and claudin 4 (CLDN4) could not be linked to  $EIS^{diff}$  (Figure 3h, k–l). When investigating the contributions of FLG, IVL, CLDN1 and CLDN4 altogether, CLDN1 and CLDN4 were not observed contributing to  $EIS^{diff}$  during HEE development and did not add additional explanatory value to the correlation model (Figure S1c, d). The expression of FLG, IVL and their collaborative interaction on the other hand together could explain 76 % of  $EIS^{diff}$  (Figure S1e, f).

### **Cytokine stimulation proves that $EIS^{diff}$ is independent of epidermal thickness**

After examining EIS in epidermal homeostasis we aimed to study the relevance of EIS in the context of disturbed homeostasis and to deepen our investigation into the correlation between  $EIS^{diff}$ , epidermal thickness and terminal differentiation. Therefore, HEEs from normal human keratinocytes (NHEKs) were stimulated with single cytokines (interleukin–(IL–) 17A or IL–22) or cytokine mixes (IL–17A + IL–22 and IL–4 + IL–13) to mimic a pro–inflammatory milieu that is known to affect keratinocyte proliferation (IL–4, IL–13, IL–17A), the cell volume (IL–22), and terminal differentiation (all cytokines) (Niehues *et al.* 2021) (Figure 4a). We hypothesized that if  $EIS^{diff}$  would merely quantify epidermal thickness, cytokines known to increase epidermal thickness would increase  $EIS^{diff}$ . Nevertheless, while IL–4 + IL–13 stimulation of HEEs significantly increased epidermal thickness, a reduction of  $EIS^{diff}$  was observed (Figure 4b, d).

Stimulation with IL-17A did not change epidermal thickness but resulted in increased  $EIS^{diff}$  and  $EIS^{SC}$  (Figure 4b-d). In contrast, IL-22 did not induce any changes in  $EIS^{diff}$ , while significantly increasing the epidermal thickness (Figure 4b-d). Correlation analysis furthermore showed no correlation between epidermal thickness and  $EIS^{diff}$  (Figure 4e). To reassess the relationship between  $EIS^{diff}$  and terminal differentiation, we analyzed the expression levels of key terminal differentiation proteins FLG and IVL which are known to be reduced in human skin related to barrier defects, and known to be affected upon stimulation with IL-4 + IL-13 cytokines *in vitro* (Furue 2020). Indeed, decreased expression levels of FLG and IVL as well as unchanged levels of CLDN4 in HEEs treated with IL-4 + IL-13 cytokines (Figure 4g) corresponded to reduced  $EIS^{diff}$  (Figure 4b). On the other hand, IL-17A treatment, which is known to strengthen tight junction function (Brewer *et al.* 2019), significantly increased  $EIS^{diff}$ , while epidermal thickness and FLG and IVL expression appeared unchanged (Figure 4b, d, g). This again indicates  $EIS^{diff}$  to quantify the complex dynamics of terminal differentiation and skin barrier formation rather than mirroring epidermal thickness. Quantifying FLG and IVL protein expression was not sufficient to model  $EIS^{diff}$  behavior (data not shown), indicating that the complex effect of cytokines on epidermal barrier function cannot be explained by the expression of FLG and IVL alone.  $EIS^{SC}$ , however, was also found here to significantly correlate with *stratum corneum* thickness (Figure 4f).

### **Knockout out of epidermal differentiation and cell-cell adhesion genes links $EIS^{diff}$ to HEE differentiation.**

To further test our hypothesis that keratinocyte terminal differentiation significantly defines  $EIS^{diff}$ , we knocked out key epidermal differentiation proteins by clustered regularly interspaced short palindromic repeats (CRISPR)/ CRISPR-associated protein 9 (Cas9)-mediated genome editing

through non-homologous end-joining. We created keratinocyte cell lines lacking terminal differentiation protein *FLG* (Smits *et al.* 2023a), tight junction protein *CLDN1* (Arnold *et al.* 2023, accepted with minor revisions) or transcription factors known to coordinate terminal differentiation namely aryl hydrocarbon receptor (AHR) (Smits *et al.* 2023b, accepted) and transcription factor activating enhancer binding protein 2 alpha (TFAP2A) (Smits *et al.* 2023b, accepted). All knockout lines showed a reduction in EIS<sup>diff</sup> and EIS<sup>SC</sup> (Figure 5a–c). Notably, *CLDN1* knockout caused reduced EIS<sup>diff</sup> but showed increased EIS<sup>SC</sup> in concordance with observed parakeratosis, increased *stratum corneum* compaction (Figure 5d). *FLG* expression was clearly decreased in *AHR*, *TFAP2A* and *CLDN1* knockout lines and completely absent in the *FLG* knockout line (Figure 5d). Expression of *IVL* was decreased in all knockout cell lines, except the *FLG* knockout. *CLDN1* (only fully absent in *CLDN1* knockout) and *CLDN4* expression appeared unchanged related to the epidermal cell layers which were affected by all genotypes as compared to control (Figure 5d). Hence, we conclude that EIS<sup>diff</sup> is strongly determined by the degree of epidermal terminal differentiation.

### **EIS detects therapeutic response in pro-inflammatory IL-4 + IL-13 epidermis model**

Besides detecting or monitoring epidermal defects, reversing these defects is a key component in the treatment of inflammatory skin diseases and an important parameter in the development of potential novel therapeutics. Therefore, we investigated if EIS can measure the reversal of barrier defects for future implementation in pre-clinical drug screening. For this we chose pharmacological molecules known to activate AHR, a key regulator of epidermal differentiation (Figure 5d) and novel target for topical anti-inflammatory treatment (Bissonnette *et al.* 2021; Van Den Bogaard *et al.* 2013). Hereto IL-4 and IL-13 stimulated HEEs were additionally treated with

an array of AHR-activating ligands (L1–5) known to have a therapeutic effect, next to structurally-related non-activating compounds (M1–2) (Table S2). AHR-activating compounds resulted in restored IL-4 + IL-13-impaired  $EIS^{diff}$  and  $EIS^{SC}$  impedance spectra, indicating capability of EIS to measure the AHR-dependent repair of skin barrier defects (Figure 6a–c). Compounds that do not activate AHR signaling (M1–2) did not restore the cytokine-mediated reduction in EIS (Figure 6a–c). In fact, these compounds known to block endogenous AHR signaling and further decreased  $EIS^{SC}$  significantly with similar trends in  $EIS^{diff}$ .

Since we now confidently determined that  $EIS^{diff}$  measurements correlate with keratinocyte differentiation, we assumed that expression levels of differentiation proteins FLG and IVL would also correlate with the rescue in  $EIS^{diff}$  by AHR ligands. This we could demonstrate clearly for IVL, as AHR agonists were partially able to restore the dampened IVL expression by IL-4 and IL-13. AHR-binding but non-activating compounds (M1–2) did not restore IVL expression (Figure S2a). FLG protein levels followed a similar expression pattern as IVL albeit differences were less pronounced. Again, the expression of tight junction protein CLDN4 remained unchanged throughout treatments (Supplemental Figure 2) confirming earlier results.

## DISCUSSION

In this study, we investigated the applicability of EIS to assess skin barrier function in 3 D HEE *in vitro* organotypic epidermis models. EIS proved an easy to handle and non-invasive system to obtain real-time quantitative readouts correlating to functional barrier properties.

While fit here to the Nunc carrier plate system, the device can be customized to various transwell cell culture platforms. Running costs are low and the measurements are performed in a semi-automated fashion. The fixed electrode setup additionally standardizes the measurements and produced results are highly repeatable when taking into account that electrical impedance readouts dependent on cell passage number, culture medium and temperature (Srinivasan *et al.* 2015). Reported readout trends are reproducible while absolute values have been observed to differ between replicated experiments. Sample readouts therefore need to be correlated with controls within the same experiment (see guideline box). While endpoint measurements cater to less variance, measurements can be performed and repeated in high quantities at virtually any time during HEE culture without harming tissue integrity. With current functional barrier assessments structurally relying on invasive endpoint measurements (permeation studies, Franz cell diffusion assay) and / or being laborious and sensitive to handling (Franz cell diffusion assay, TEWL, TEER), EIS provides a non-invasive, semi-automated and reproducible alternative.

EIS measures a broad range of frequencies in contrast to single frequency TEER measurements (Faway *et al.* 2021; Srinivasan *et al.* 2015) therefore being able to capture different functional skin barrier parameters. For quantitative analysis, experimentally obtained impedance spectra are usually fit to the corresponding electrical circuit model to isolate individual electrical parameters (Magar *et al.* 2021). To aid biologic interpretation, we here chose to correlate the obtained impedance spectra with known biological barrier properties. Theoretical considerations in

combination with our own data highlighted two frequency ranges which we described through calculating their area under the curve.

Frequencies on a plateau around 100–2000 Hz were termed  $EIS^{diff}$  as they quantified differentiation in viable keratinocytes as assessed by the expression of differentiation markers FLG and IVL, which together predicted 76 % of  $EIS^{diff}$  during HEE development. While  $EIS^{diff}$  exhibits clear independence of viable epidermis and *stratum corneum* thickness, a correlation with tight junction function remains uncertain. On the one hand,  $EIS^{diff}$  was observed to be independent from CLDN1 and CLDN4 protein expression during HEE development and cytokine stimulations.  $\Delta CLDN1$  HEEs exhibits a strongly reduced  $EIS^{diff}$ , but this can be explained by a concomitant reduced expression of keratinocyte differentiation markers as CLDN1 knockout is known to alter pro-FLG processing (Sugawara et al. 2013). On the other hand,  $EIS^{diff}$  frequencies overlap with  $R_p$  impedance contribution, with  $R_p$  being the electrical circuit model element to describe the resistance of tight junctions (Srinivasan et al. 2015). Furthermore, during IL-17A cytokine stimulation  $EIS^{diff}$  increased independent of differentiation protein expression, which could be a result of an IL-17A strengthening effect on tight junction function (Brewer et al. 2019). In conclusion,  $EIS^{diff}$  uniquely quantifies keratinocyte differentiation independent of CLDN1 and CLDN4 protein expression, however a contribution of tight junction function cannot be ruled out since sole protein expression does not entirely mirror tight junction functionality (Bäsler et al. 2016; Kirschner et al. 2012).

Frequencies of a higher frequency range between 20,000 Hz and 100,000 Hz were termed  $EIS^{SC}$  and overlap with  $C_{Cell}$ , a parameter describing ability of a cell to store an electrical charge.  $EIS^{SC}$  conclusively quantifies *stratum corneum* more than complete HEE thickness and seems to be a descriptive rather than functional assessor since the thickened but parakeratotic  $\Delta CLDN1$  *stratum*

*corneum* exhibited an increased EIS<sup>SC</sup>. We did not investigate EIS<sup>SC</sup> dependence on lipid organization and *stratum corneum* composition, which should be the focus of future studies.

This study used EIS to assess HEE skin barrier function during formation, to study the effects of single genes and to assess skin barrier function under inflammatory conditions and treatment. EIS was able to measure the defects induced by knockout of cardinal differentiation-driving transcription factors (AHR and TFAP2A) and differentiation effector genes (FLG). The observed defects were congruent with other barrier function assessments reporting an elevated TEWL in  $\Delta$ TFAP2A and  $\Delta$ FLG HEEs and in human FLG loss-of-function variants (Nemoto-Hasebe *et al.* 2009; Smits *et al.* 2023b, accepted; Smits *et al.* 2023a; Yang *et al.* 2020). Cytokine-induced pro-inflammatory conditions resulted in keratinocyte differentiation deficiencies and changes in *stratum corneum* thickness which could be captured and quantified by EIS. The IL-4 + IL-13 induced decrease in EIS<sup>diff</sup> and EIS<sup>SC</sup> *in vitro* also replicates the *in vivo* situation where the IL-4 and IL-13 driven skin disease atopic dermatitis is accompanied by elevated TEWL and decreased EIS values measured on *in vivo* patient skin (Chamlin *et al.* 2002; Flohr *et al.* 2010; Rinaldi *et al.* 2021; Sugarman *et al.* 2003). *In vivo*, EIS can also detect therapeutic improvements of atopic dermatitis associated with improvements in clinical scoring and reduced expression of inflammatory biomarkers (Breternitz *et al.* 2008; Rinaldi *et al.* 2021) similar to the detected therapeutic improvements in our *in vitro* atopic dermatitis model.

To conclude, we propose EIS to be a valuable tool to non-invasively study epidermal barrier function in organotypic skin models. The dual viable epidermis/*stratum corneum* barrier assessment and the quantification of keratinocyte differentiation is to our knowledge singular across all barrier evaluation techniques. The proposed semi-quantitative EIS analysis is easy to replicate and uniquely correlates impedance readouts with biological barrier properties. We

suggest EIS to be especially suited for longitudinal studies of barrier development, keratinocyte differentiation and barrier-disrupting skin diseases including pre-clinical therapeutic studies. In addition, EIS can be used in multi-cell type models to investigate the interplay between epidermis, extrinsic and intrinsic factors, potentially in combination with patient-derived cells, immune cells and/or bacteria. The possibility to correlate *in vitro* and *in vivo* EIS measurements facilitates a unique translational approach from bedside to bench and back.

### **~~GUIDELINE BOX~~ TECHNICAL RECOMMENDATIONS**

To ensure optimal, reproducible EIS measurements without compromising culture integrity, we propose several guidelines for implementing EIS in the laboratory:

- Measurements depend on temperature and ion content of the surrounding fluid. To ensure maximum comparability between conditions, use an isotonic buffer solution (e.g. phosphate-buffered saline (PBS)) and allow samples to adjust to room temperature for at least 30 min before measurements.
- To minimize variation when performing serial measurements at various days of the cell culture, the time of topical exposure to PBS should be kept minimal and PBS should be carefully removed after measurements to maintain the air-liquid interface as much as possible for proper barrier formation and function.
- Before commencing measurements, blank measurements on PBS only or empty filters should be performed, as this provides information on intrinsic capacitance of the electrodes and the resistance of PBS and filters. When analyzing the results, blanks should be subtracted from measured sample values.
- Previous publications have normalized EIS based on the surface area of the used cell

culture system using various methods (Chen et al. 2015; Haorah et al. 2008; Liu et al. 2020). Considering the lack of consensus, we report uncorrected EIS values and the surface area of HEEs (0.47 cm<sup>2</sup>) to aid comparisons.

- Control conditions should be taken along for each individual experiment and measurement time point to interpret relative changes in preference to absolute values.
- EIS<sup>diff</sup> (127–2212 Hz) and EIS<sup>SC</sup> (28,072–100,000 Hz) are determined through calculating the area under the curve at respective frequencies.

## **MATERIALS & METHODS**

### **Cell culture**

Human primary keratinocytes were isolated from surplus human skin obtained through plastic surgery according to the principles and guidelines of the principles of Helsinki. From the skin, biopsies were taken and keratinocytes were isolated as described previously (Tjabringa *et al.* 2008). N/TERT–2G keratinocytes were a kind gift of James Rheinwald, Brigham’s Woman hospital (Dickson *et al.* 2000) and were cultured as monolayers in CnT–prime (CELLnTEC, Bern, Switzerland, CnT–PR) until confluent before use in HEE cultures (Smits *et al.* 2017). Knockout N/TERT–2G cell lines were generated through CRISPR/Cas9 and validated previously (FLG (Smits *et al.* 2023a), CLDN1 (Arnold *et al.* 2023, accepted with minor revisions), TFAP2A (Smits *et al.* 2023b, accepted), AHR (Smits *et al.* 2023b, accepted)).

### **Generation of HEEs**

HEE cultures were performed as previously described (Smits *et al.* 2023a). In short, cell culture inserts in a 24 wells carrier plate (Nunc, Thermo Fisher Scientific, 141002) were coated using

100 µg/mL rat tail collagen (Sigma–Aldrich, C3867) for 1 hour at 4 °C. After phosphate–buffered saline (PBS) washing the filters, 150000 cells were seeded and submerged in CnT–prime medium (CELLnTEC, CnT–PR) at the lowest insert stand. After 48 hours, the medium was switched to differentiation medium (40 % Dulbecco’s modified Eagle’s Medium (Sigma–Aldrich, D6546) and 60 % 3 D barrier medium (CELLnTEC, CnT–PR–3D)) and 24 hours afterwards the HEEs were lifted to the highest stand, air–exposed and medium was refreshed every other day. For stimulation experiments, IL–4, IL–13, IL–17A or IL–22 (50 ng/mL per cytokine, Peprotech, Rocky Hill, NJ, USA, 200-04/200-13/200-17/200-22) supplemented with 0.05 % bovine serum albumin (Sigma–Aldrich, A2153) were added to the medium of HEEs of primary keratinocyte from day 5 of air exposure until day 8. AHR ligands (Table S2) were supplemented in the culture medium as previously described (Rikken et al. 2023).

### **EIS measurements**

For the EIS measurements the Locsense Artemis (Locsense, Enschede, the Netherlands) device was used and equipped with a custom–made incubator compatible smart lid. The Artemis consists of a detector element that is connected to the smart lid with electrodes aligning to the two middle rows of a 24–well plate. A laptop equipped with the Locsense Artemis monitoring software (version 2.0) displays the readouts. During the measurements each well contains two electrodes: one disc–shaped 4.2 mm diameter electrode situated in the center of the transwell insert and a rod–shaped 1.9 mm diameter electrode passing sideways of the transwell insert. Before measurements, HEEs were acclimated to room temperature and cultures were lowered to the middle position in the transwell plate while 1600 µL PBS at room temperature was added below and 500 µL PBS on top of the filter. Thereafter, the smart lid was placed on the wells ensuring both electrodes being

submerged. Following device self-calibration, impedance was measured over a frequency range from 10 Hz to 100,000 Hz in 30 logarithmic intervals. Measurement output contains impedance as well as phase values. Phase values can be interpreted as is while a PBS only blank measurement was subtracted from the corresponding electrode of the impedance output. For further specific considerations during measurements, see guideline box. For  $EIS^{diff}$  (127–2212 Hz) and  $EIS^{SC}$  (28,072–100,000 Hz) the area under the curve was calculated using the respective frequency ranges.

### **Immunohistochemistry**

For histological processing, 4 mm biopsies were fixated in 4 % formalin solution for 4 hours and embedded in paraffin. Afterwards, 6  $\mu$ m sections were deparaffinized and either stained with hematoxylin (Klinipath, 4085.9005) and eosin (Klinipath, 4082.9002) or by antibodies listed in Table S3 followed by avidin-biotin complex (Vectastain, AK-5000). Epidermal thickness specifies the average of three measurements on H&E stained sample pictures while *stratum corneum* thickness was determined by subtracting epidermal from total construct thickness. Protein expression was quantified with ImageJ following sections C–E of (Crowe and Yue 2019) by freehand selecting the viable epidermis and measuring the “area” i.e. number of staining-positive square pixels.

### **Statistics**

Data sets were analyzed using the GraphPad Prism 10 software version 10.1.1. All barplots are shown as mean  $\pm$  standard error of the mean and significance testing was performed using one-way analysis of variance (ANOVA) in combination with Dunnett correction for multiple testing

and unpaired t-testing (exclusively in figure 2c, d). Differences under p value < 0.05 were considered statistically significant, ns p value > 0.05, \* p value < 0.05, \*\* p value < 0.01, \*\*\* p value < 0.001.

### **Correlation analysis of EIS to protein expression and HEE morphology**

Correlation analysis was conducted using simple (epidermis thickness and *stratum corneum* thickness) and multiple linear regression modelling (protein expression) in Graphpad Prism and the R programming language (version 4.2.3) (R Core Team 2023) with the psychometric package (Fletcher 2023). All correlation analysis were conducted with individual replicates, figures depict replicate averages for readability.

### **DATA AVAILABILITY STATEMENT**

Datasets related to this article are available from the corresponding author upon request.

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## **CONFLICT OF INTEREST**

SR is CEO and founder of Locsense B.V., which contributed in-kind to this work. The results presented in the study are not influenced nor determined by the views or wishes of Locsense, nor did Locsense provide any financial support for this study that may conflicted with the results interpretation or presentation of data. The contribution of Locsense was limited to the development of the smart lid to fit the cell culture system and to providing technical support. Discussions with Locsense on the data representation and electrical circuit interpretation aided in correlating the data output to biological interpretations. The remaining authors declare no conflicts of interest.

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## **AUTHOR CONTRIBUTIONS**

Conceptualization: NB, FP, LM, JS, and EB; Data Curation: NB, FP, and LM; Formal Analysis: NB and FP; Funding Acquisition: JS, EB; Investigation: NB, FP, LM, HN, IVW, MB, DRO, PJ JS; Methodology: NB, FP, LM, JS; Project Administration: EB; Software: SR; Supervision: JS and EB; Validation: NB and FP; Visualization: NB and FP; Writing – Original Draft Preparation: NB and FP; Writing – Review and Editing: all authors.

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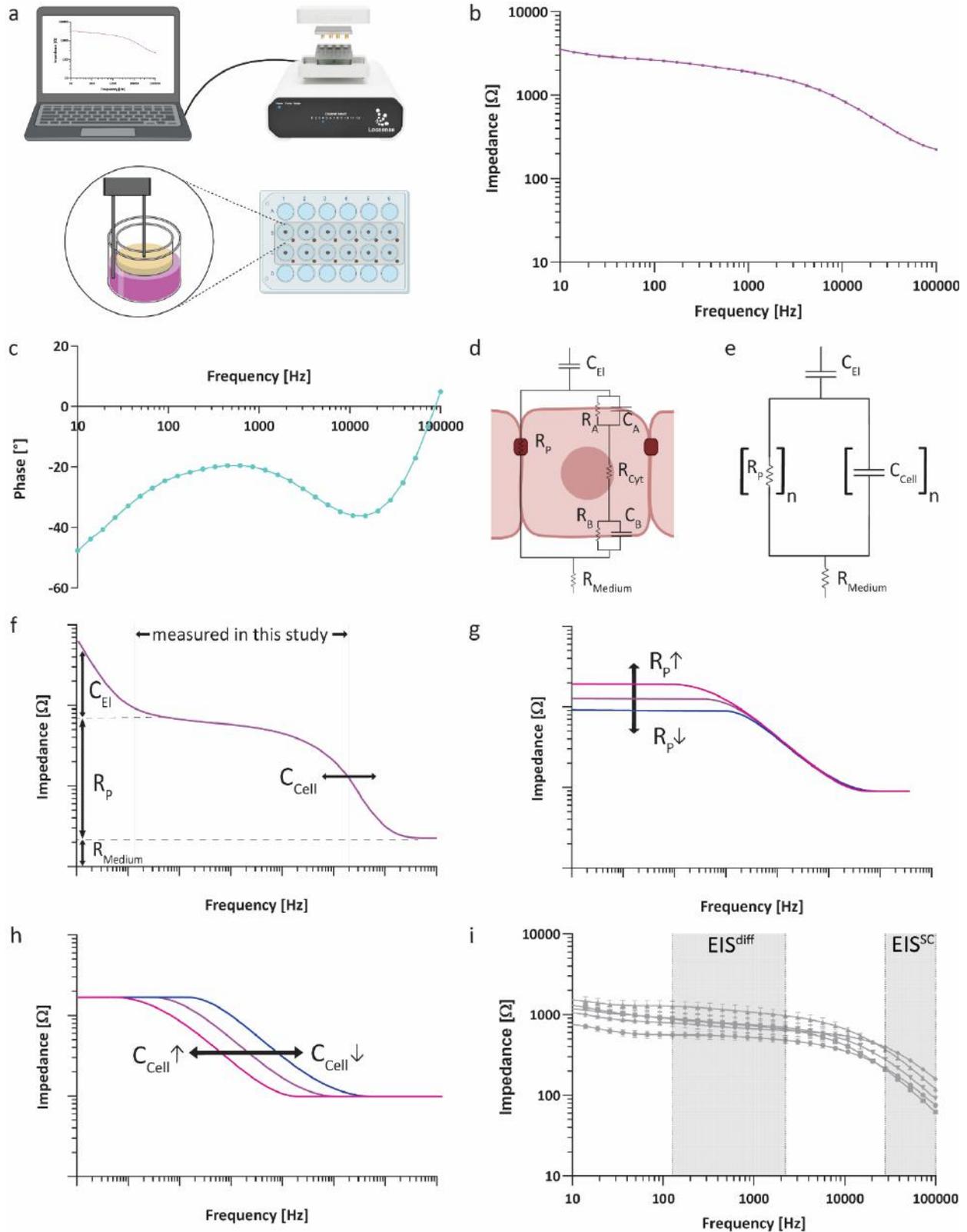
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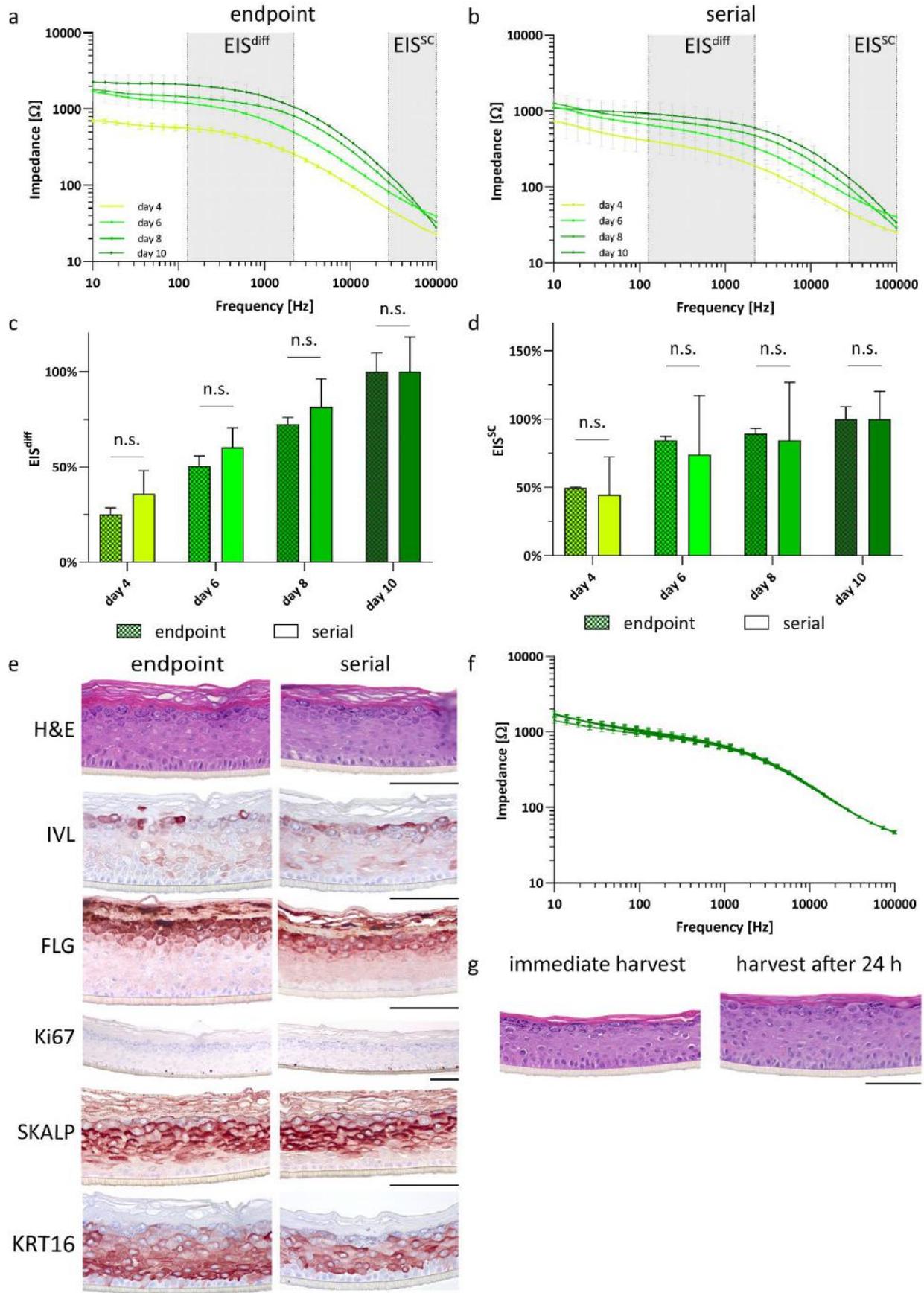
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## FIGURE LEGENDS

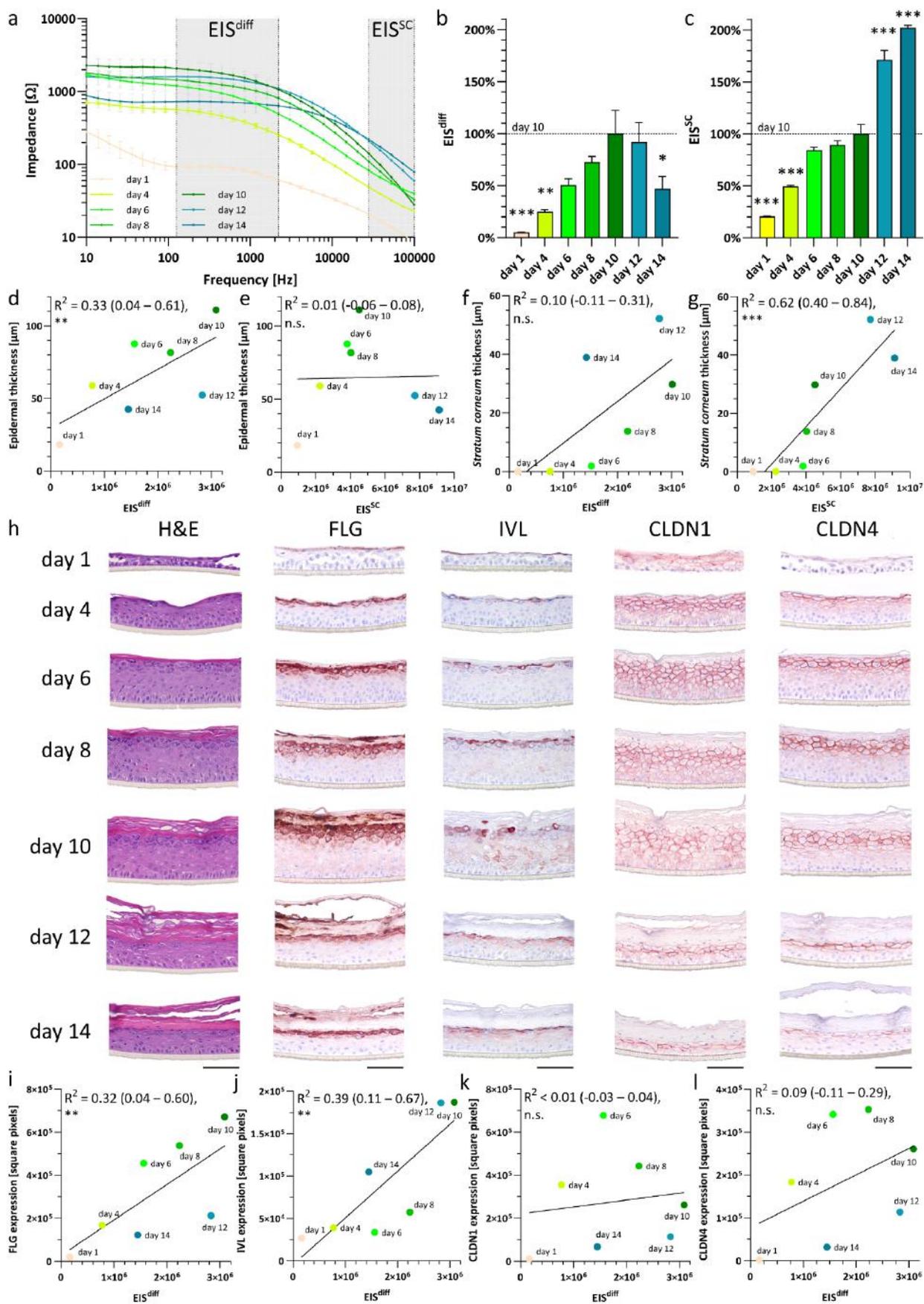


**Figure 1: Design and function principle of a custom-made EIS device fitting the HEE culture system.** (a) Schematic overview of the EIS setup on HEE cultures. (b) Impedance and (c) phase spectrum of a fully-developed HEE culture after 8 days of air exposure. (d) Extended electrical equivalent circuit of an epidermal monolayer culture made up of the capacitance of the electrodes  $C_{EI}$ , paracellular resistance  $R_P$ , transcellular resistance of cytoplasm  $R_{Cyt}$ , apical and basolateral membrane  $R_A$ ,  $R_B$  as well as their capacitance  $C_A$ ,  $C_B$  and the resistance of the medium  $R_{Medium}$  (adapted from (Yeste et al. 2018)). (e) Simplified electrical equivalent circuit of a HEE with the capacitance of both membranes taken together as  $C_{Cell}$  which together with  $R_P$  extends to a series of  $n$  parallel circuits in multi-layered 3 D culture systems (adapted from (Srinivasan et al. 2015) and (Groeber et al. 2015)). (f) Schematic overview indicating the contribution of individual electrical circuit parameters to the impedance spectrum (adapted from (Benson et al. 2013)). (g, h) Simulated impedance spectra illustrating the influence of changes in (g) paracellular flux ( $R_P$ ) and (h) transcellular flux ( $C_{Cell}$ ) (adapted from (Yeste et al. 2018)). (i) EIS impedance spectrum displaying  $EIS^{diff}$  (127–2212 Hz) and  $EIS^{SC}$  (28072–100000 Hz).

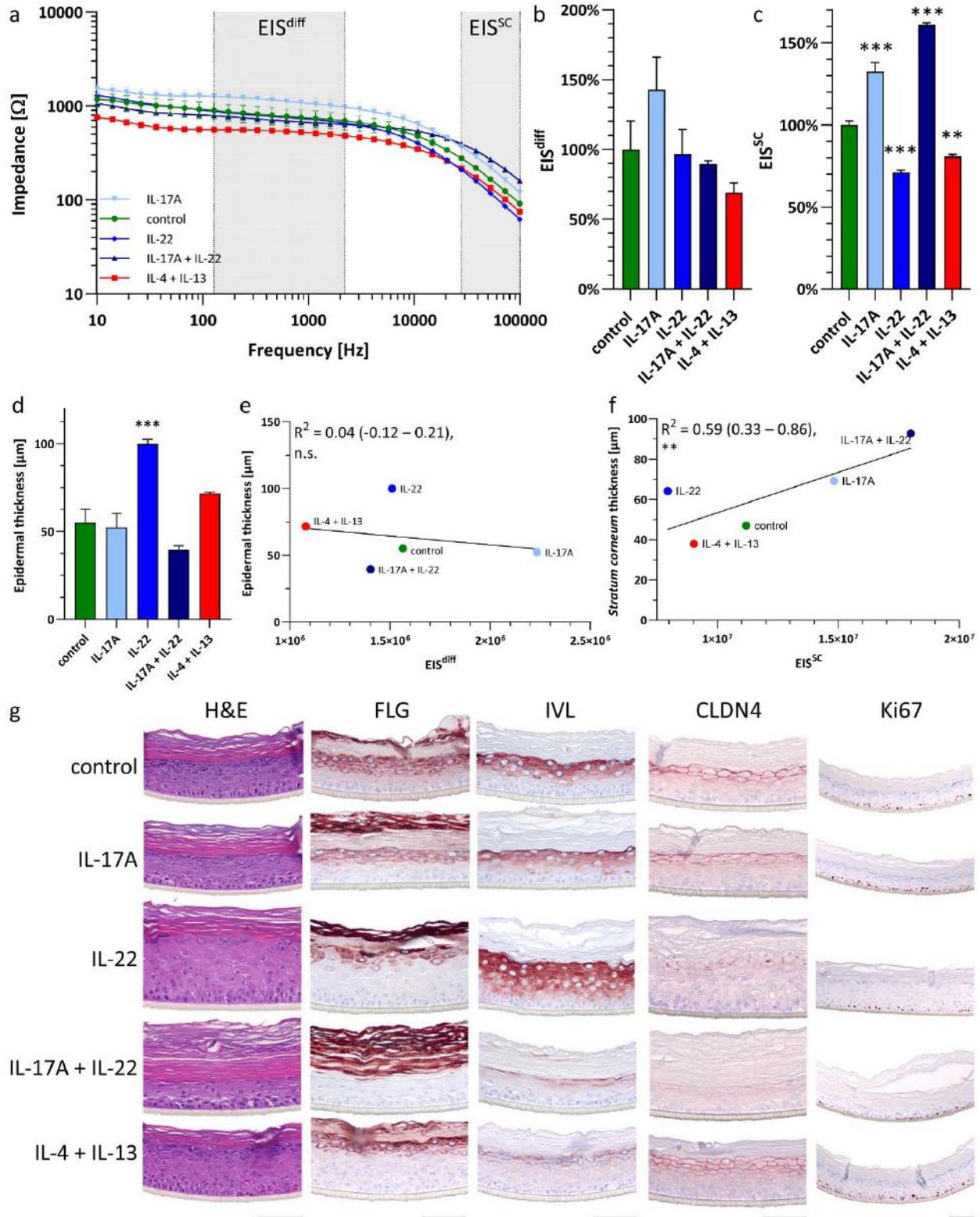


**Figure 2: Relative EIS measurements are reproducible and do not impair HEE development.**

(a, b) EIS impedance spectra during HEE development with constructs being harvested (a) directly after measurements (endpoint measurements) and (b) at day 10 of air exposure (serial measurements). Each timepoint averages three biological replicates. (c) Comparison of EIS<sup>diff</sup> and (d) EIS<sup>SC</sup> between endpoint and serial measurements. (e) Histological comparison of HEEs undergoing endpoint or serial EIS measurements based on general morphology (H&E staining), differentiation status (FLG, IVL expression), proliferation (Ki67) and stress response (SKALP, KRT16). Pictures represent three biological replicates at day 10 of air exposure and taken at either 20x (Ki67) or 40x magnification. Size bars indicate 100  $\mu\text{m}$ . (f) Impedance spectrum of HEEs ( $n = 3$ ) measured 6 times within 1 hour at day 6 of air exposure. (g) Histological comparison of HEEs measured 6 times in 1 hour, either harvested directly or 24 hours after EIS measurements. Pictures represent three biological replicates and were taken at 40x magnification. Size bar indicates 100  $\mu\text{m}$ .

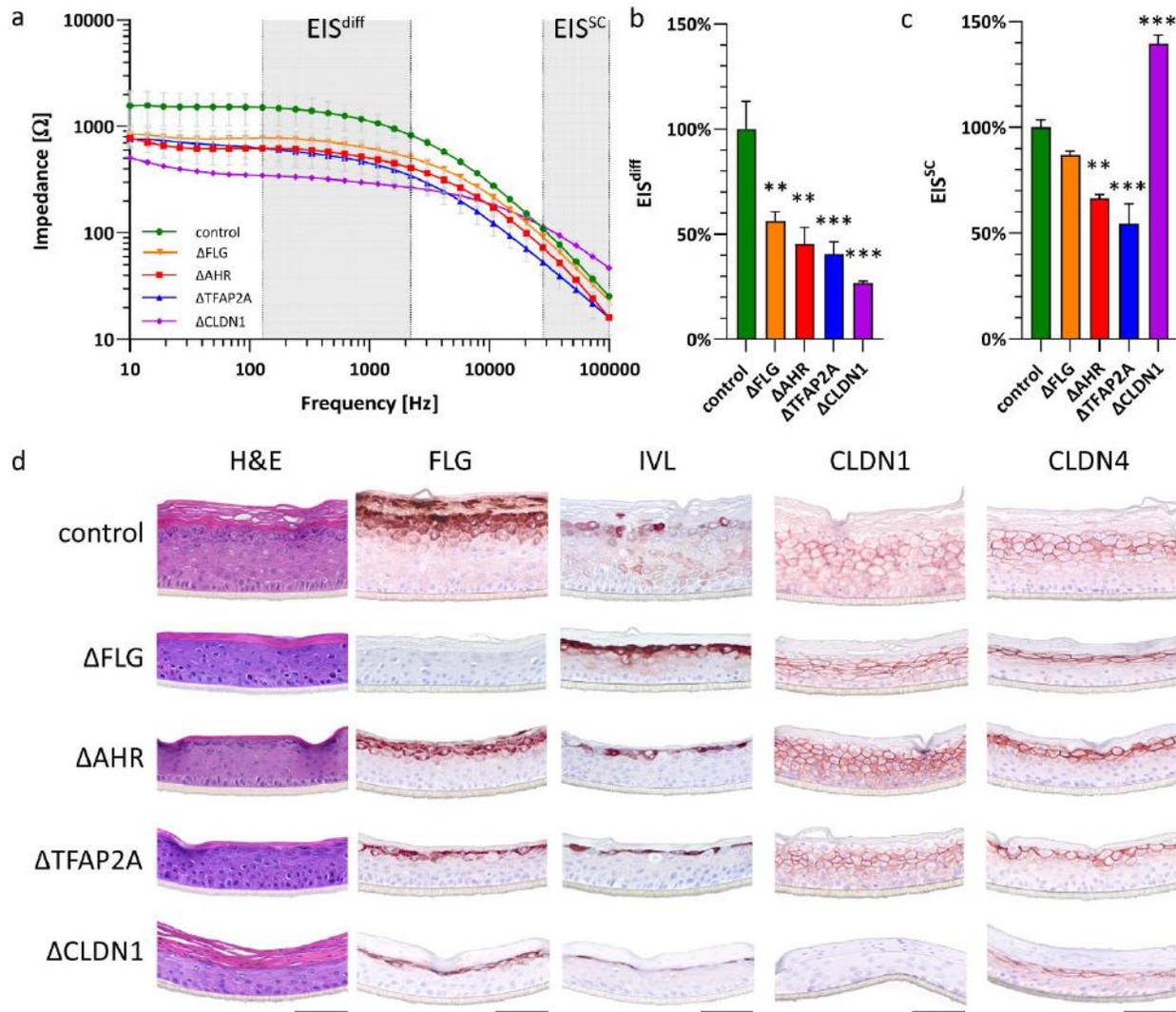


**Figure 3: During HEE development, EIS<sup>diff</sup> correlates with keratinocyte differentiation and epidermal thickness while EIS<sup>SC</sup> correlates with *stratum corneum* thickness.** (a) Endpoint-measured impedance spectra, (b) EIS<sup>diff</sup> and (c) EIS<sup>SC</sup> during HEE development. Each timepoint represents three biological replicates and EIS<sup>diff</sup> and EIS<sup>SC</sup> were compared to 10 day air-exposed cultures. (d – g) Correlation of epidermal thickness with (d) EIS<sup>diff</sup> and (e) EIS<sup>SC</sup> and *stratum corneum* thickness with (f) EIS<sup>diff</sup> and (g) EIS<sup>SC</sup>. Each timepoint averages three biological replicates, R<sup>2</sup> values and significances indicate the correlation of individual replicates. (h) Staining of general morphology (H&E), keratinocyte differentiation (FLG, IVL) and cell-cell adhesions (CLDN1, CLDN4) during HEE development. Pictures represent three biological replicates and were taken at 40x magnification. Size bars indicate 100 μm. (i – l) Correlation of (i) FLG, (j) IVL, (k) CLDN1 and (l) CLDN4 protein expression with EIS<sup>diff</sup>. Each timepoint averages three biological replicates, R<sup>2</sup> values and significances indicate the correlation of individual replicates.

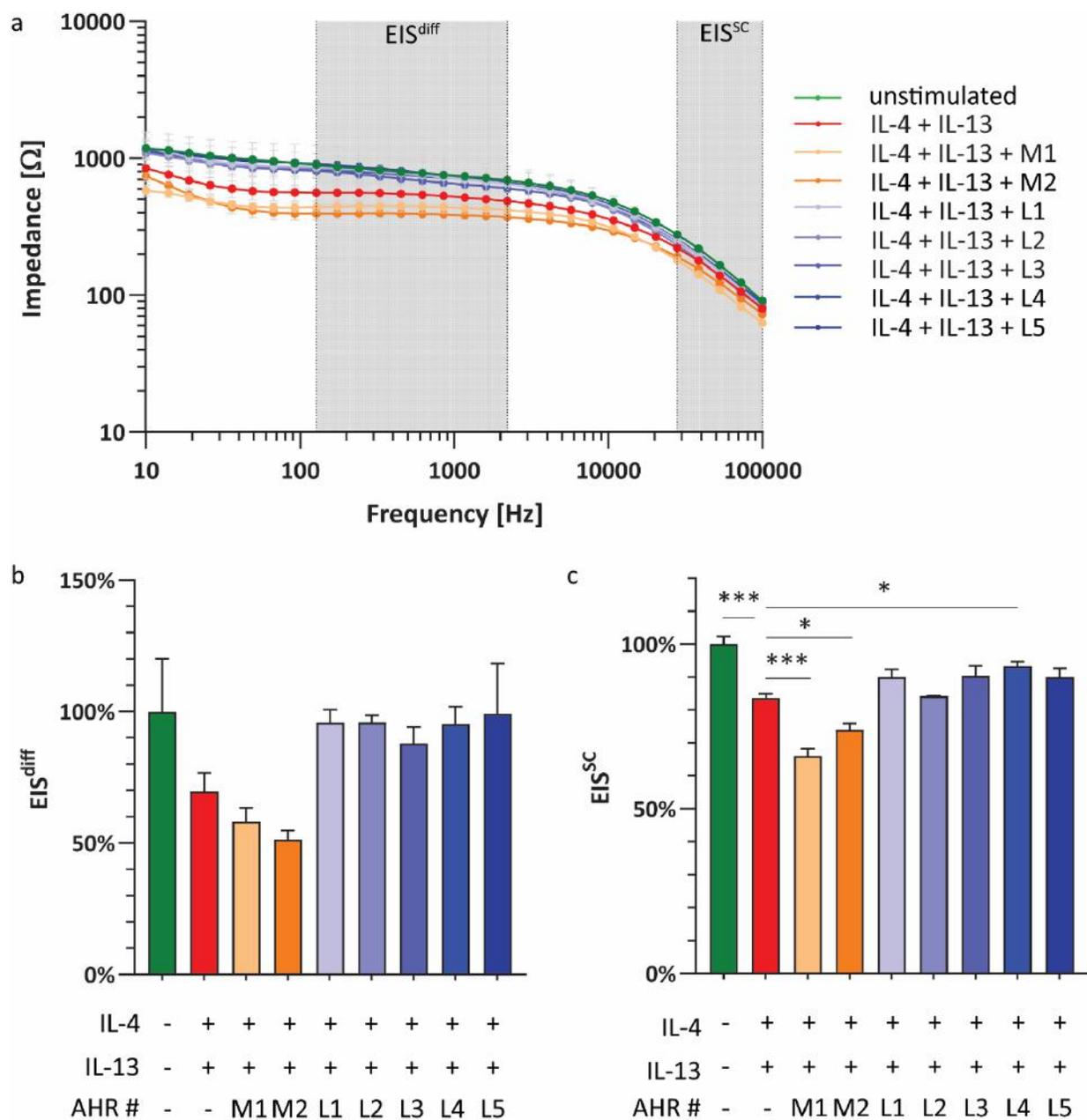


**Figure 4: Stimulation with cytokines prove  $EIS^{diff}$ -determined barrier function to be independent of epidermal thickness.** (a) Endpoint-measured impedance spectra, (b)  $EIS^{diff}$  and

(c) EIS<sup>SC</sup> of cytokine–stimulated HEEs at day 8 of air exposure. Each condition represents three biological replicates and EIS<sup>diff</sup> and EIS<sup>SC</sup> were compared to control. (d) Epidermal thickness of cytokine–stimulated HEEs as compared to control. Correlation of (e) epidermal thickness to EIS<sup>diff</sup> and (f) *stratum corneum* thickness to EIS<sup>SC</sup>. Each condition averages three biological replicates, R<sup>2</sup> values and significances indicate the correlation of individual replicates. (g) HEEs stained for differentiation (FLG, IVL) and cell–cell adhesion (CLDN4) and proliferation (Ki67) proteins. Pictures represent three biological replicates and were taken at 20x (Ki67) or 40x magnification. Size bars indicate 100 μm.



**Figure 5: Knockout of genes involved in keratinocyte differentiation and cell–cell adhesion decreases  $EIS^{diff}$ .** (a) Endpoint–measured impedance spectrum, (b)  $EIS^{diff}$  and (c)  $EIS^{SC}$  of HEEs with knockout of target gene at day 10 of air exposure. Each condition represents three biological replicates and  $EIS^{diff}$  and  $EIS^{SC}$  are compared to control. Each (d) HEEs stained for differentiation (FLG, IVL) and cell–cell adhesion (CLDN1, CLDN4) proteins. Pictures represent three biological replicates and were taken at 40x magnification. Size bars indicate 100  $\mu m$ .



**Figure 6: EIS detects therapeutic AHR response in a pro-inflammatory epidermis model.** (a)

Endpoint-measured impedance spectrum, (b)  $EIS^{diff}$  and (c)  $EIS^{SC}$  of HEEs stimulated with IL-4 and IL-13 cytokines alone and in combination with AHR activating therapeutic compounds at day 8 of air exposure. Each condition represents three biological replicates and  $EIS^{diff}$  and  $EIS^{SC}$  of control conditions and AHR-binding compounds are compared to IL-4 + IL-13 stimulation.

**SUPPLEMENTARY MATERIAL**

**Supplemental Table 1. Overview of methodologies used for measurement of *in vitro* epidermal barrier function**

Technique	Principle of analysis	of Assessed barrier component	Advantages	Disadvantages	References
<b>Permeation studies</b>	Paracellular diffusion of dyes/tracers of various molecular weights	tight junction pore of size, of corneodesmosome integrity, lipid barrier	<ul style="list-style-type: none"> <li>• relatively cheap</li> <li>• easy to implement</li> </ul>	<ul style="list-style-type: none"> <li>• destructive end–point parameter</li> <li>• conventionally non–quantitative and not high–throughput</li> </ul>	(Arlk et al. 2018; van den Bogaard et al. 2021)
<b>Franz cell diffusion assay</b>	Assessment of drug permeation from donor chamber through HEE into acceptor chamber	of outside–in barrier function including <i>stratum corneum</i> and tight junction barrier function	<ul style="list-style-type: none"> <li>• gold standard for assessing outside–in barrier</li> <li>• quantitative</li> </ul>	<ul style="list-style-type: none"> <li>• destructive end–point parameter</li> <li>• high level of expertise required</li> <li>• model system must stay intact during sampling period after harvest</li> <li>• highly influenced by temperature, sampling frequency and stirring conditions</li> </ul>	(Luengo et al. 2021; Neupane et al. 2020)

<b>Transepidermal water loss (TEWL)</b>	Measurement of lipid barrier changes in air humidity on top of HEE	• non-destructive relation to <i>stratum corneum</i> thickness	• highly influenced by probe type and angle, contact pressure, temperature and atmospheric pressure • quantitative • high variability between different instruments • high background readings • labor intensive	(Alexander et al. 2018; Klotz et al. 2022)
<b>Transepithelial electrical resistance (TEER)</b>	2 electrodes apply low alternating current over the HEE at a single frequency	• non-destructive (“tight junction tightness”)	• highly influenced by electrode positioning, temperature and medium composition • quantitative • not designed for complex tissues • easy to implement • labor intensive	(Srinivasan et al. 2015)

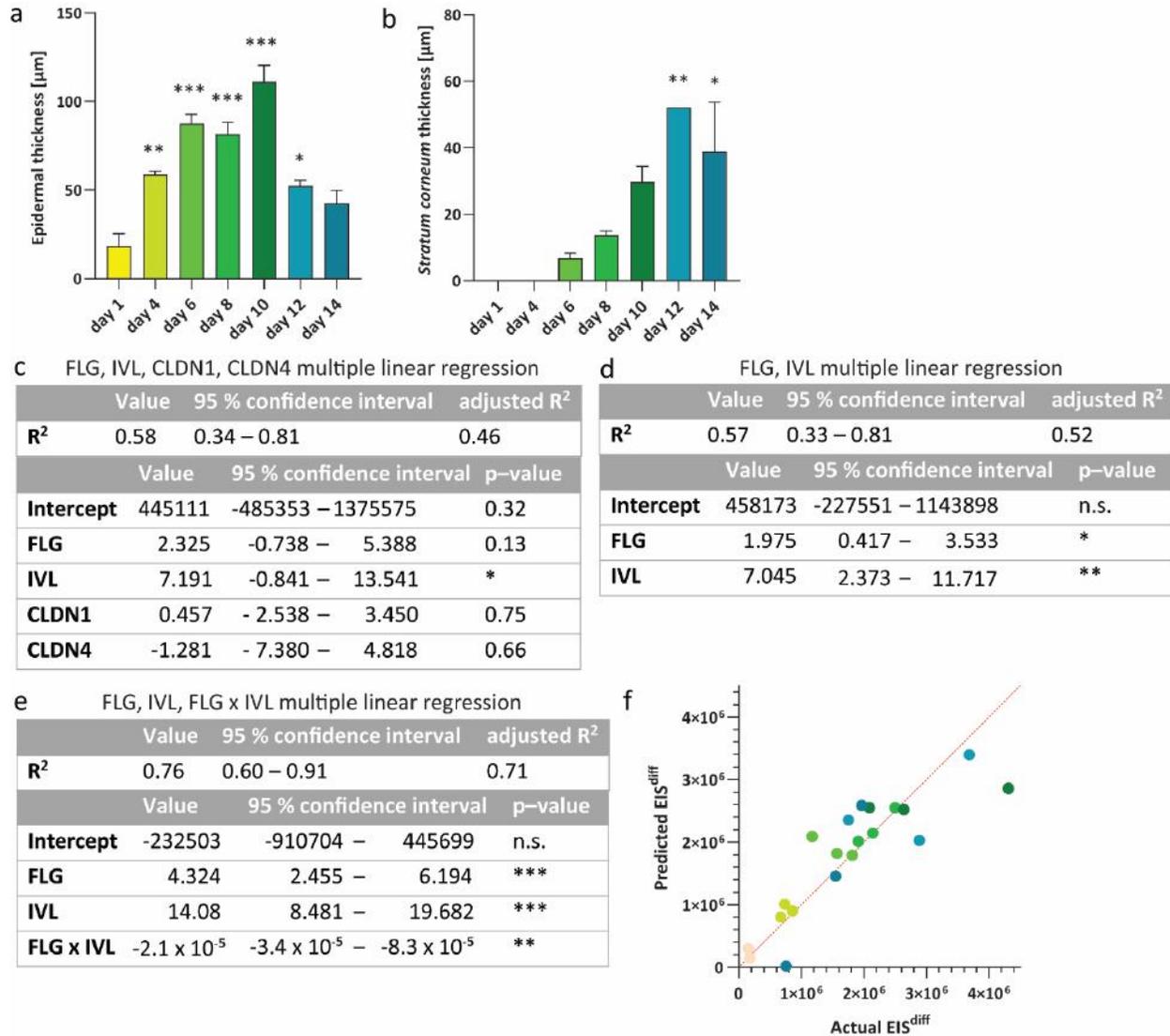
HEE: human epidermal equivalent; TEER: transepithelial electrical resistance; TEWL: transepidermal water loss

### Supplemental Table 2: AHR-binding compounds

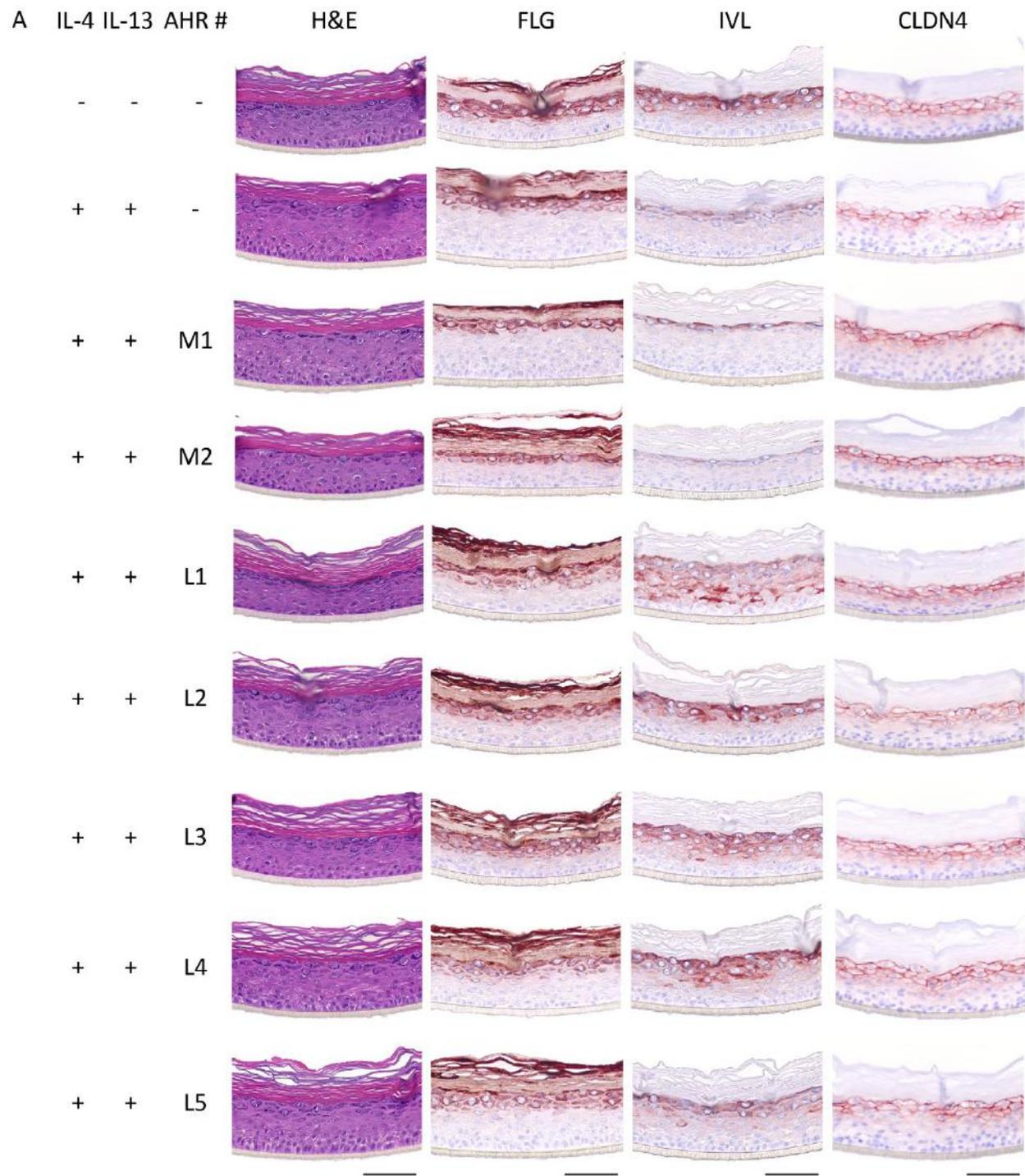
Abbreviation	Compound	Concentration	Reference
<b>M1</b>	SGA360	10 nM	(Rikken et al. 2023)
<b>M2</b>	Teriflunomide	10 $\mu$ M	(O'Donnell et al. 2010)
<b>L1</b>	coal tar	2 $\mu$ g/mL	(Van Den Bogaard et al., 2013)
<b>L2</b>	SGA388	10 nM	(Rikken et al. 2023)
<b>L3</b>	Leflunomide	10 $\mu$ M	(O'Donnell et al. 2010)
<b>L4</b>	IMA-7101	1 nM	(Rikken et al. 2022)
<b>L5</b>	TCDD	10 nM	(Poland and Glover 1976; Safe 1986)

### Supplemental Table 3: Antibodies used for immunohistochemical analysis

Antigen	Species	Dilution	Company
<b>FLG</b>	Mouse	1:100	Thermo Fisher
<b>IVL</b>	Mouse	1:20	Mon 150 (own antibody)
<b>CLDN1</b>	Rabbit	1:200	Invitrogen
<b>CLDN4</b>	Mouse	1:200	Invitrogen
<b>Ki67</b>	Rabbit	1:200	Abcam
<b>SKALP</b>	Rabbit	1:4000	Own antibody
<b>KRT16</b>	Mouse	1:200	Santa-Cruz
<b>Goat anti-rabbit</b>	Goat	1:200	Vector laboratories
<b>Horse anti-mouse</b>	Mouse	1:200	Vector laboratories



**Supplemental Figure 1: FLG, IVL and FLG x IVL interaction significantly influence EIS<sup>diff</sup> during HEE development.** (a–b) (a) Epidermal thickness and (b) *stratum corneum* thickness of during HEE development compared to day 1. Each condition represents three biological replicates. (c–e) Multiple linear regression using (c) FLG, IVL, CLDN1 and CLDN4, (d) FLG and IVL and (e) FLG, IVL and FLG x IVL expression to predict EIS<sup>diff</sup>. Analysis is based on all timepoints and replicates from figure 3. (f) Correlation of actual and FLG, IVL, FLG x IVL regressed EIS<sup>diff</sup>.



**Supplemental Figure 2: AHR activation mediates therapeutic response in a pro-inflammatory epidermis model.** (a) HEEs stained for differentiation (FLG, IVL) and cell-cell

adhesion (CLDN4) proteins. Pictures represent three biological replicates of HEEs at day 8 of air exposure and were taken with 40x magnification. Size bars indicate 100  $\mu\text{m}$ .