Supplementary material to article by K. Baumann et al. "The Skin Reservoir Model: A Tool for Evaluating Microdialysis Sampling of Large Biomarkers from Human Skin"



Fig. S4. Relative skin recovery in response to different perfusate compositions. Two perfusates (with or without addition of lactic acid to a final concentration of 4 mM) were used for sampling and the relative skin recoveries of (a) CXCL1/GROa, (b) CXCL7/NAP-2, (c) CXCL10/IP-10, (d) EGF, (e) GM-CSF, (f) IFN- γ , (g) IL-1a, (h) IL-6, (i) IL-8, (j) IL-17, (k) IL-22, (l) IL-23, (m) TNF- α , (n) TSLP and (o) VEGF were measured. Depicted are individual samples with triplicate probes for each condition (n=3 probes) and bars denoting mean±standard deviation. Cytokine concentrations in dialysates were determined by enzyme-linked immunoassay (ELISA) and background cytokine levels were subtracted before calculating the relative skin recoveries. *Open symbols* indicate values below lower limit of quantification (LLOQ). All setups were tested in 1 donor per cytokine except for IL-6 and TNF- α , which use assessed in skin from 2 donors (CXCL1/GROa: D31, CXCL7/NAP-2: D26, CXCL10/IP-10: D32, EGF: D23, GM-CSF: D31, IFN- γ : D11, IL-1a: D5, IL-6: D17 and D26, IL-8: D30, IL-17: D25, IL-22: D11, IL-23: D17 and D25, TSLP: D23, and VEGF: D32). *p*-values depicted are based on *t*-tests with Welch's correction comparing skin recovery of cytokines in response to the 2 perfusate compositions.

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