



Fig. S2. Real-time-PCR experiments reveal differences in expression between non-involved psoriatic and healthy epidermis samples in response to T-cell lymphokine treatment. (A) Mean gene expression changes upon T-cell lymphokine treatment are presented. Expression values of the healthy treated samples (black bar) are compared to the untreated healthy ones (grey bar), while lymphokine-treated psoriatic samples (striped bar) are compared to the gene expression values measured in the untreated psoriatic ones (white bar). (B) Basal gene expression levels of untreated healthy (grey bar) and psoriatic non-involved epidermis samples (white bar) were also compared. All data were compared to the expression of the untreated healthy samples. In both panels, results are presented as mean of 4 samples. Error bars indicate standard error of mean.