



Fig. S1. Expression of IL13 correlates with several inflammation and epidermal barrier associated genes in AD lesional skin. Each scatter plot displays the correlation with IL13 expression of the indicated gene in lesional skin samples from three individual transcriptomic studies that were pooled together ($n = 100$). All CPM (counts per million) data were floored to 0.01 and log₂-transformed before performing differential gene expression analysis by paired t-test (patient eliminated as factor), and correlation analysis by linear regression. Multiple testing was accounted for by estimating false discovery rates (FDR).