Supplemental Figure 5. Inverse correlation between pri-miR-221 and REDD1 expression during glucocorticoid treatment. B6D2 mice were treated with FA topically as described in Materials and Methods, epidermis was harvested at indicated time points. A. Western blot analysis. B. Q-PCR analysis of primary miR-221. Data are expressed as the fold change (mean ± SD) with respect to the controls (unpaired two-tailed t-test, n = 3).