



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 \pm SD) with respect to the controls (unpaired two-tailed t-test, $n = 3$).