



Supplement Figure 9 GR^{dim} FLS show a lack of induction of classical GC-target genes and fail to suppress Mip1- α and - β and show enhanced gene expression involved in multiple pathways.

(A) FPKM (Fragment Per Kilobase Million) of classical GC-target genes *Fkbp5*, *Mt* metallothionein-1 and -2 and Glucocorticoid-induced leucine zipper (*Gilz*) in wt and GR^{dim} FLS (B) FPKM (Fragment Per Kilobase Million) of chemokines *Mip1*- α and - β in wt and GR^{dim} FLS. (C) Enriched KEGG pathways in wt versus GR^{dim} FLS treated with $\text{IL-1}\beta$ and Dexamethasone.