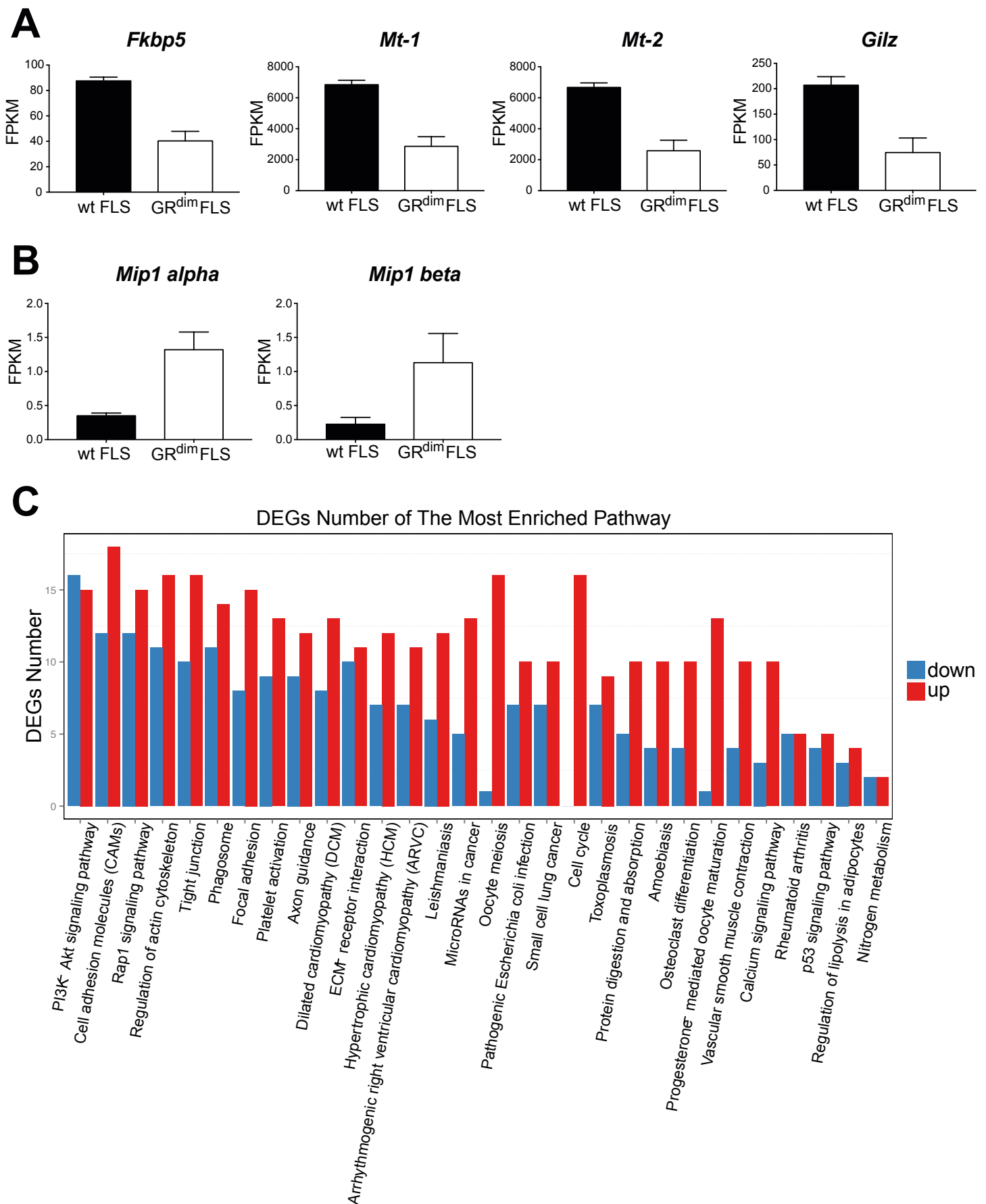




**Supplement Figure 9 GR<sup>dim</sup> FLS show a lack of induction of classical GC-target genes and fail to suppress Mip1- $\alpha$  and - $\beta$  and show enhanced gene expression involved in multiple pathways.**

(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<sup>dim</sup> FLS (B) FPKM (Fragment Per Kilobase Million) of chemokines *Mip1*- $\alpha$  and - $\beta$  in wt and GR<sup>dim</sup> FLS. (C) Enriched KEGG pathways in wt versus GR<sup>dim</sup> FLS treated with IL-1 $\beta$  and Dexamethasone.