

Description of Additional Supplementary Files

File name: Supplementary Data 1

Description: All differentially expressed genes based on whole gonad RNA-seq of E12.5 gonads. Genes are organised based on the cluster as appears in the heatmap in Fig 3B.

File name: Supplementary Data 2

Description: Contains the matrices of all mammalian transcription factors (2346) used to screen changes in the mutant versus WT Enh13. There are also 16 gonad-specific TFs used to screen the WT and mutant Enh13 as presented in Figure 4A and Supplementary Figure 10.