## Supplemental Figure 3



**Supplementary Figure 3.** Number of differentially expressed genes (DEGs) induced by Fe deficiency and Gene ontology (GO) enrichment analysis of DEGs. **(A)** Up-regulated DEGs in AC and *Cnr* roots. **(B)** Down-regulated DEGs in AC and *Cnr* roots. **(C-D)** GO enrichment analysis of DEGs in AC and *Cnr* epimutant.