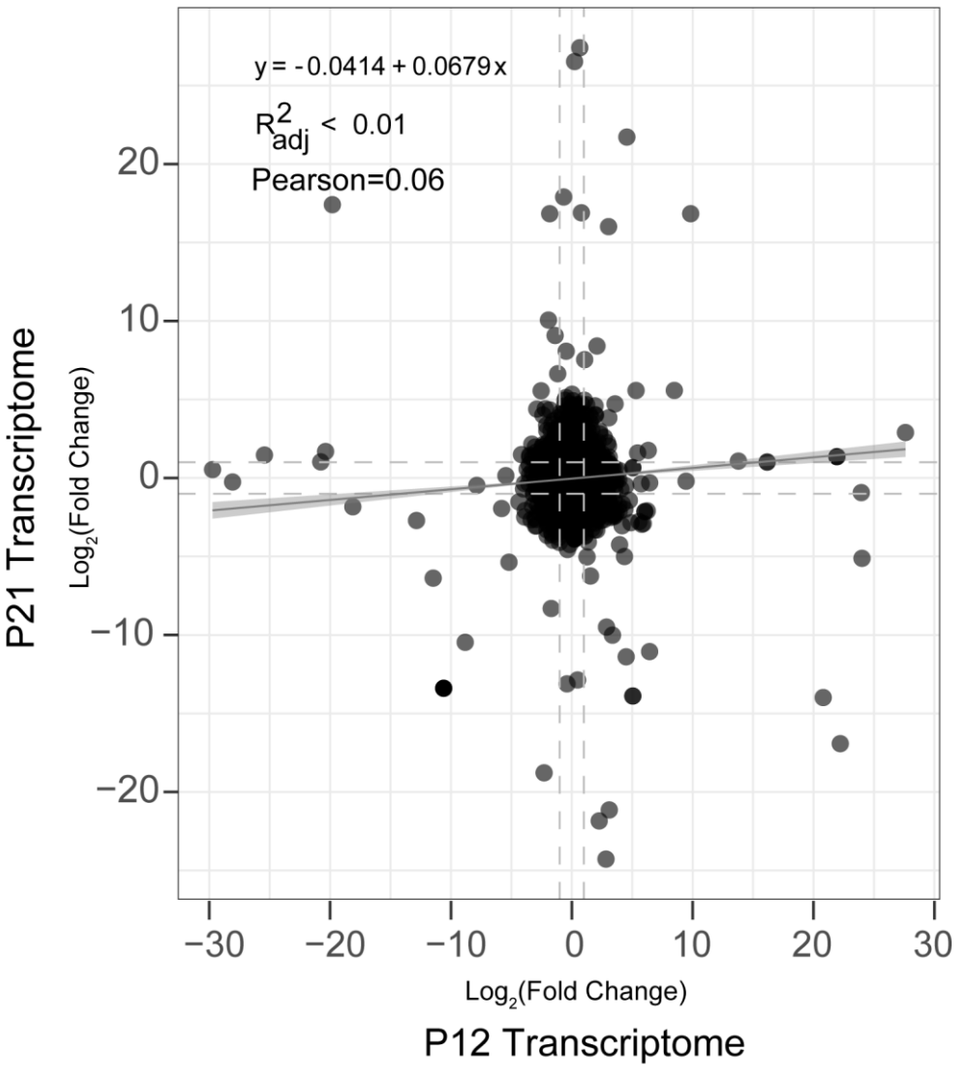


Figure S1. Correlation analysis of the altered transcriptome at P12 and P15. The data were downloaded from National Center for Biotechnology Information Gene Expression Omnibus (https://ncbi.nlm.nih.gov/geo/): GSE153874). Those genes quantified in all samples (P12_WT, P12 Rs1 KO, P21_WT, and P21_Rs1 KO) were identified as quantified genes. In total, 26,005 quantified genes (remove duplicate values) were used for correlation analysis.