

Exon Ontology: Functional Genomics At Exon Level Resolution

Supplemental Figure S1

RT-PCR analysis of the 81 selected exons that are differentially spliced when comparing mesenchymal- and epithelial-like cells, using total RNAs from 4 normal mesenchymal cell types (Fibro), 4 normal epithelial cell types (Epi), 4 breast cancer mesenchymal-like cell types (Claudin low), and 4 breast cancer epithelial-like cell types (Luminal).





