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Integrated whole transcriptome and small RNA analysis revealed multiple regulatory networks in colorectal cancer

Shaath, H, Toor, SM, Nada, MA, Elkord, E and Alajez, NM

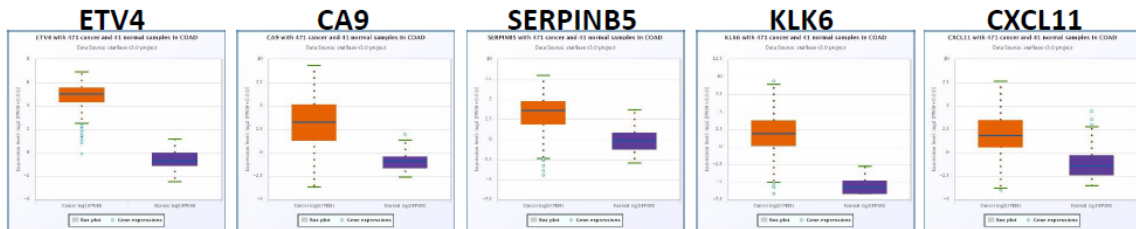
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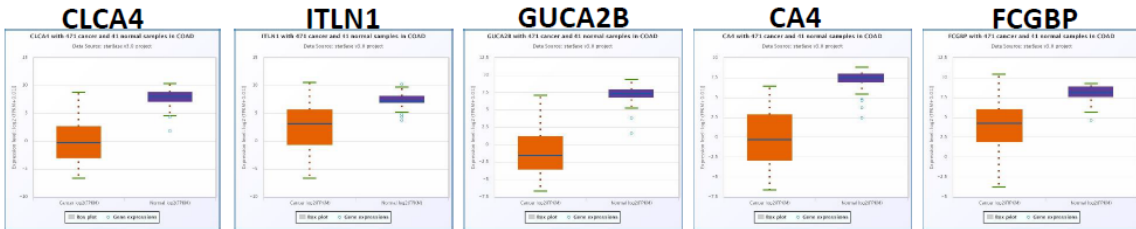
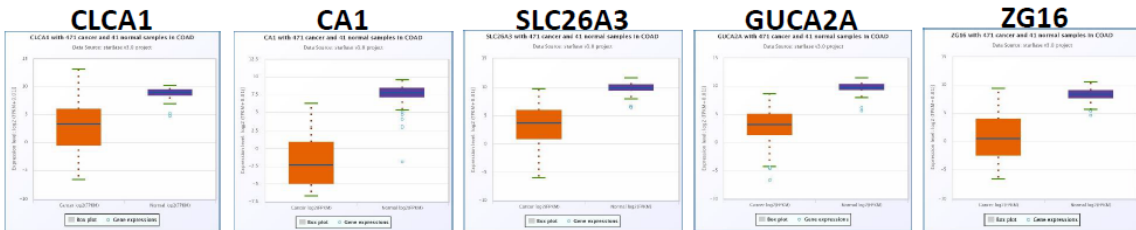
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Upregulated



Downregulated



Supplementary figure 1. Expression of top 10 upregulated and top 10 downregulated mRNAs in the TCGA COAD database. Expression of the indicated gene in COAD (n=471) compared to normal tissue (n=41) is shown as box plot. Data were retrieved from the starBase V3.0 database.