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

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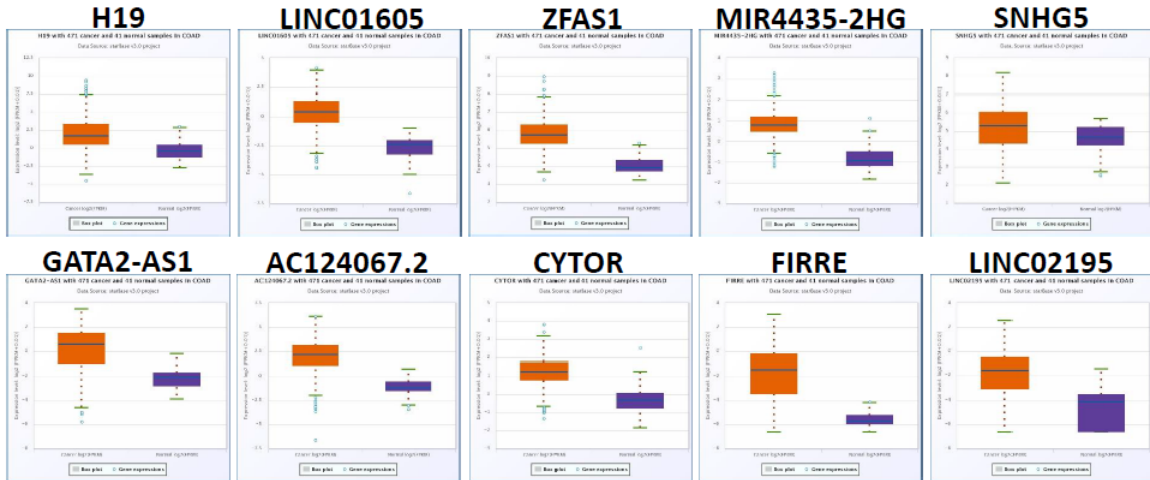
<http://dx.doi.org/10.1038/s41598-021-93531-y>

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Type	Article
URL	This version is available at: http://usir.salford.ac.uk/id/eprint/61219/
Published Date	2021

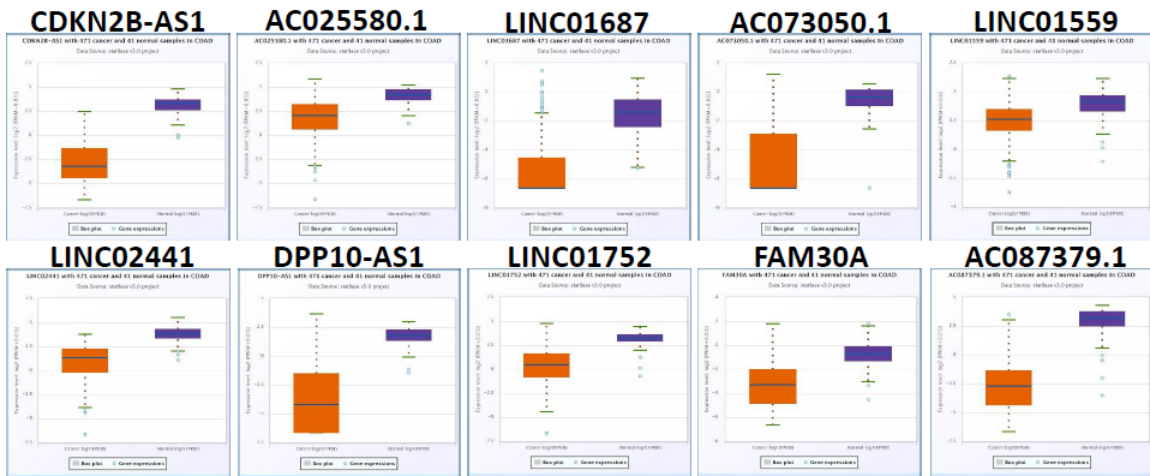
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Upregulated



Downregulated



Supplementary figure 2. Expression of top 10 upregulated and top 10 downregulated lncRNAs 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.