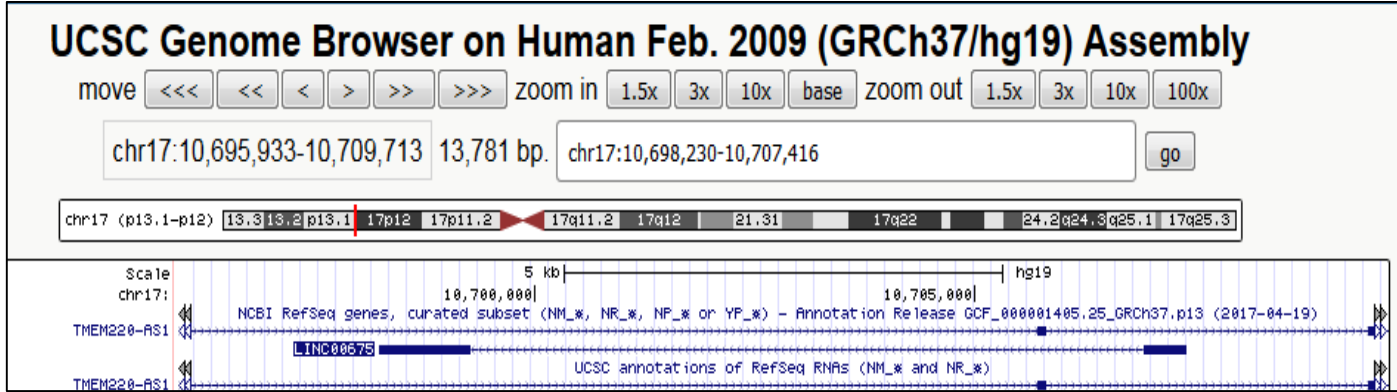
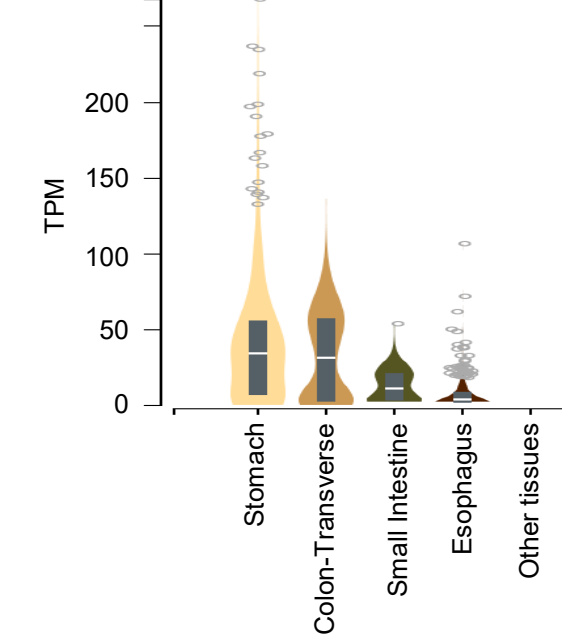


A



B



C

Gene	Median TPM (N=406) in Colon Transverse in GTEx Portal
FORCP	31.3
MALAT1	219.7
NORAD	95.03

D

Cell line	FPKM for FORCP (<i>LINC00675</i>)
HCT116	0.054
RKO	0.01
SW48	0.01

E

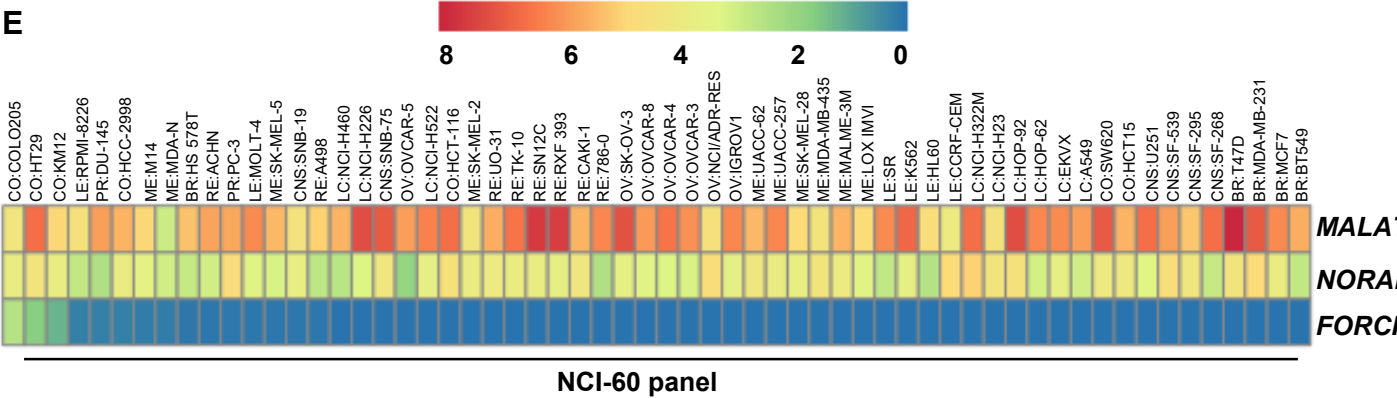


Figure 1-figure supplement 1.

(A) Snapshot of UCSC genome browser shows that *FORCP* (*LINC00675*) locus overlaps with the *TMEM220-AS1* intron transcribed from the opposite strand. (B) *FORCP* expression in RNA-seq across normal human tissues (data from GTExPortal). TPM refers to transcripts per kilobase million. (C) Expression of *FORCP*, *MALAT1* and *NORAD* in RNA-seq from normal human tissues (data from GTEx Portal). (D) Expression of *FORCP* in poorly differentiated CRC lines HCT116, RKO and SW48 is shown. See Supplementary file 2 for more details. FPKM refers to fragments per kilobase of exon model per million reads mapped. (E) Heat map showing the expression pattern of *FORCP* and the abundant lncRNAs *MALAT1* and *NORAD* in the NCI-60 panel of cell lines. See Supplementary file 1 for more details.