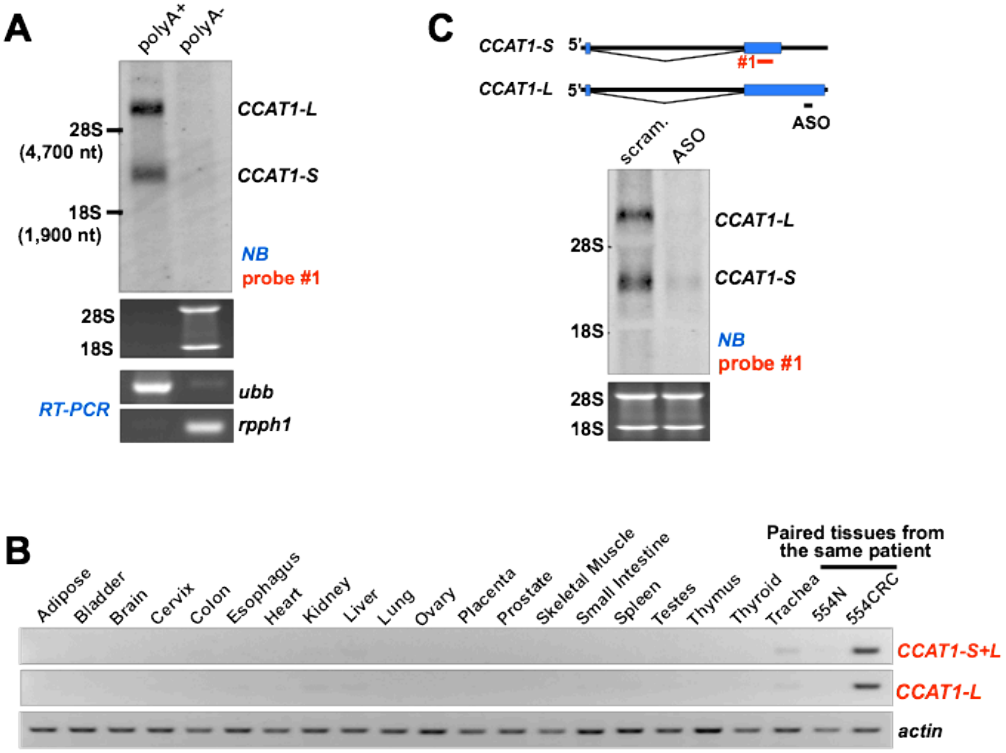

Supplementary information, Figure S2 Characterization of *CCAT1-L* (related to Figure 1). (A) *CCAT1* RNAs are polyadenylated (poly(A)+). NB validated both *CCAT1* isoforms are enriched in oligo d(T) selection. Middle, 28S and 18S rRNAs were used as markers for non-polyadenylated (poly(A)-) RNAs on native Agarose gel. Bottom, *ubb* and *rpph1* were used as markers for poly(A)+ and poly(A)- RNAs in semi-quantitative RT-PCR. (B) *CCAT1-L* is specifically expressed in human CRC as revealed by semi-quantitative RT-PCR. Primers used were described in Figure 1B. The same amount of cDNA from each sample was used to amplify *actin* mRNA as a loading control. (C) Knockdown of *CCAT1-L* led to the simultaneous disruption of *CCAT1-S*. NB revealed the relative abundance of *CCAT1-L* and

CCAT1-S after the treatment of HT29 cells with an ASO (shown in Figure 2A) that targets only *CCAT1-L.* 18S and 18S rRNAs were used as loading controls.