



Supplementary information, Figure S1 Genome organization of 8q24 region lncRNAs (related to Figure 1). (A) 3' RACE result of CCAT1-L. The Sanger sequencing of the 3' end RACE result of CCAT1-L is shown. (B) Genome organization of 8q24 region lncRNAs and their tissue-/tumor-type specific expression patterns are also indicated. (C) CCAT1s are much more abundantly expressed in human CRC patient samples than other 8q24 lncRNAs. Data were retrieved from RNA-sequencing datasets of the paired

CRC/control mucosa samples in this study. (D) CCAT1-L is much more abundantly expressed in examined CRC-derived cell lines than other 8q24 lncRNAs. RT-qPCR detected the abundant expression of CCAT1-L, but not other 8q24 lncRNAs in HCT116, HT29 and SW48 cell lines.