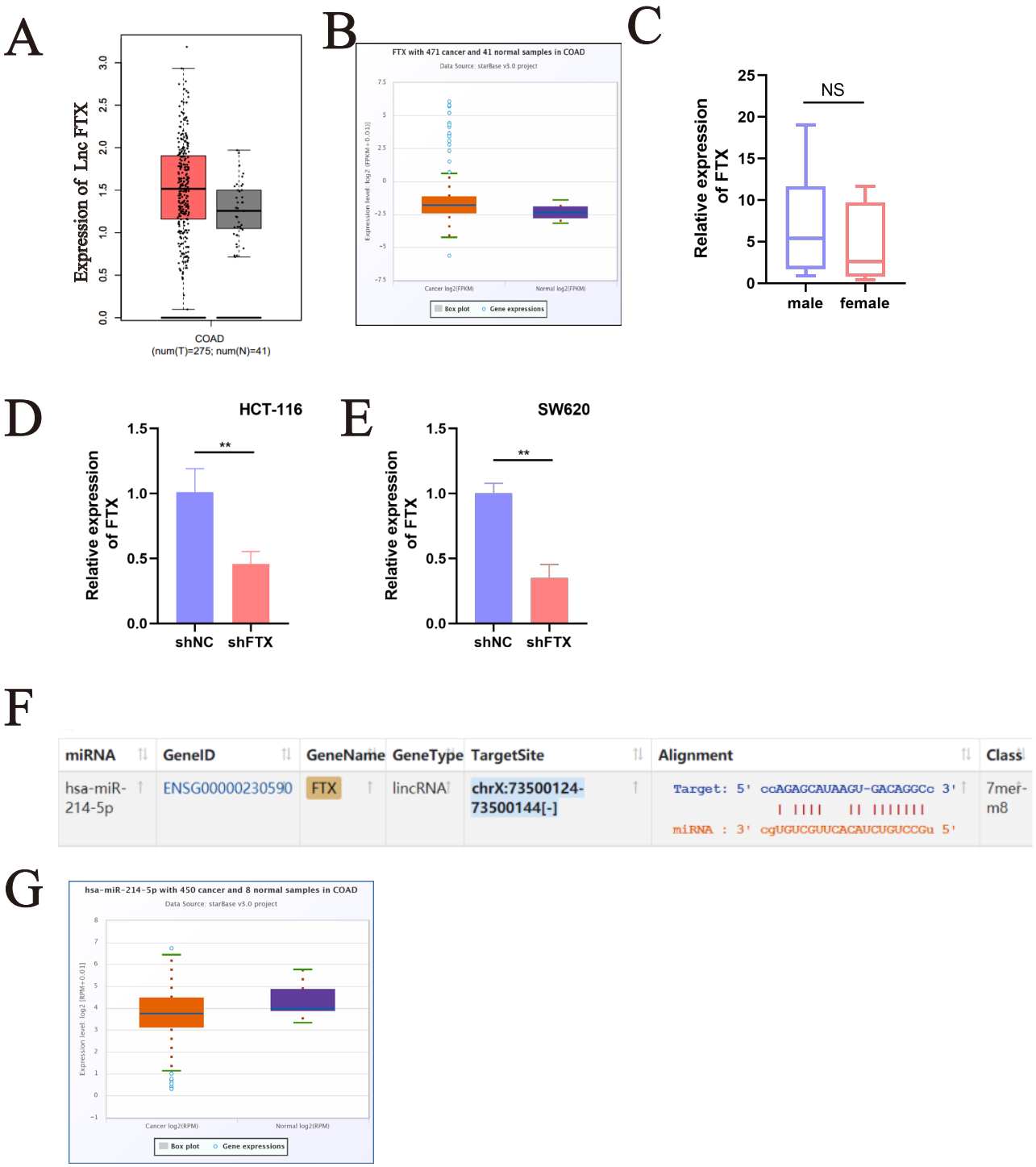

Figure S1 Expression of lncRNA FTX inversely correlates the expression of miR-214-5p in clinical samples and knockdown efficiency of lncRNA FTX in CRC cell lines. (A,B) LncRNA *FTX* expression in colorectal cancer tissues based on the Gene Expression Profiling Interactive Analysis and TCGA databases. (C) Comparison of lncRNA *FTX* expression between male and female CRC patients. (D,E) Expression of lncRNA *FTX* in HCT-116 and SW620 cells following transfection with shNC or sh-*FTX* was determined by quantitative real-time polymerase chain reaction. (F) Predicted target sites in miR-214-5p and lncRNA *FTX* based on the starBase database. (G) Expression of miR-214-5p in CRC tissues based on the TCGA database. **, P<0.01; NS, nonsignificant. CRC, colorectal cancer.

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