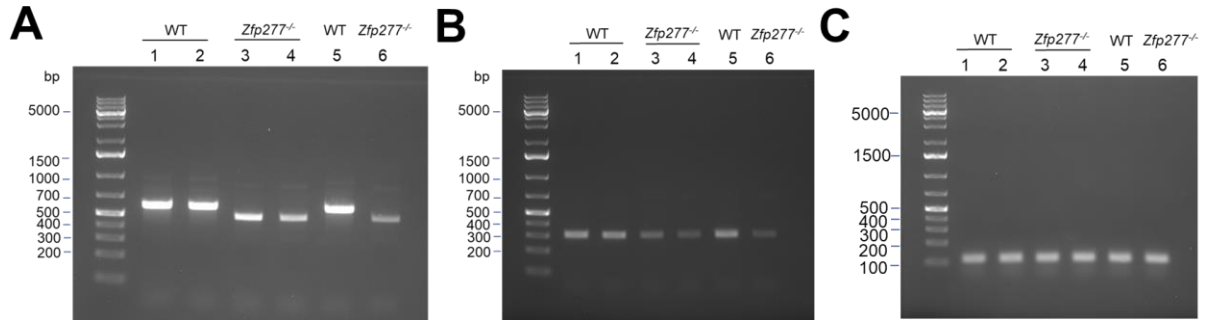
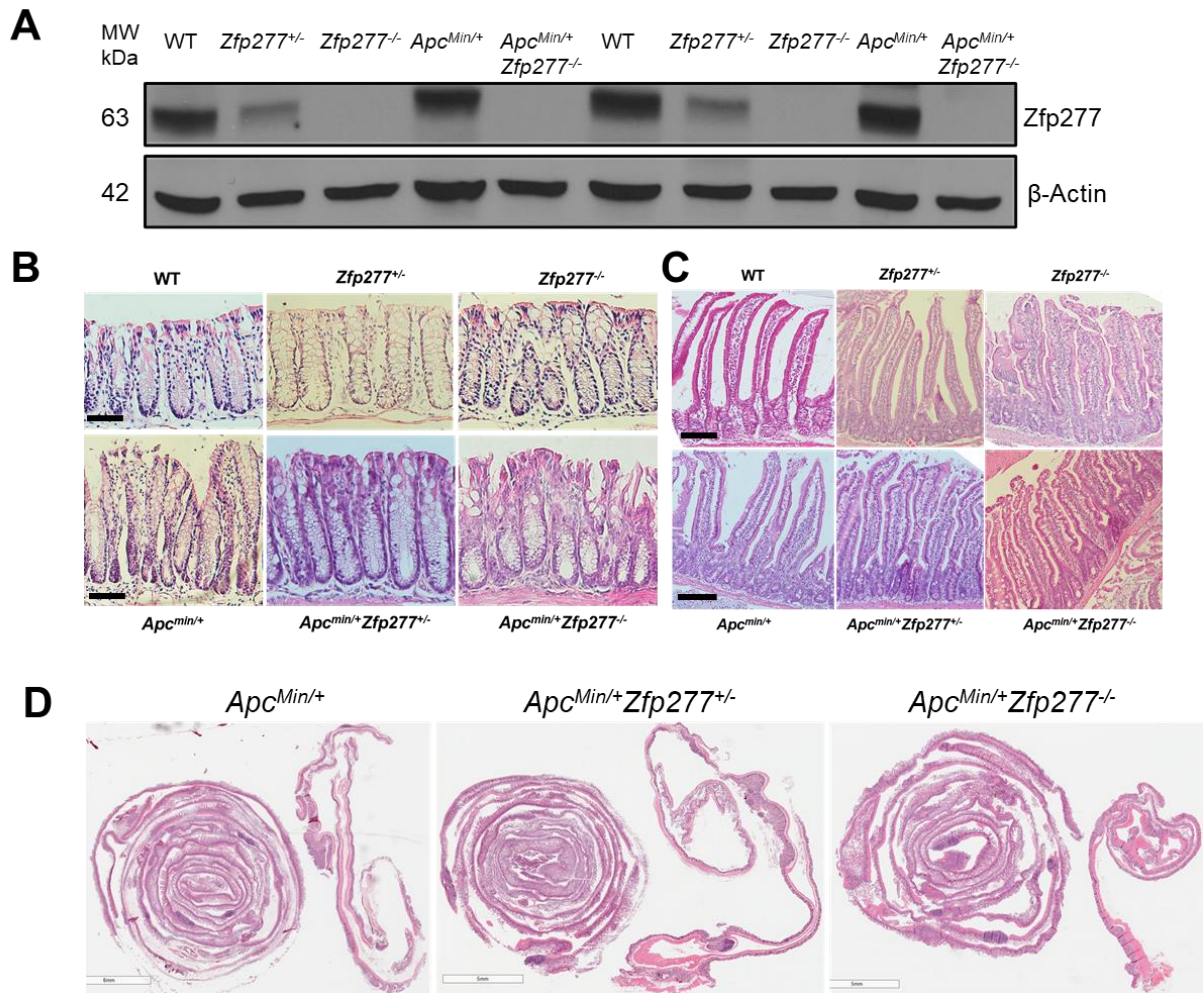


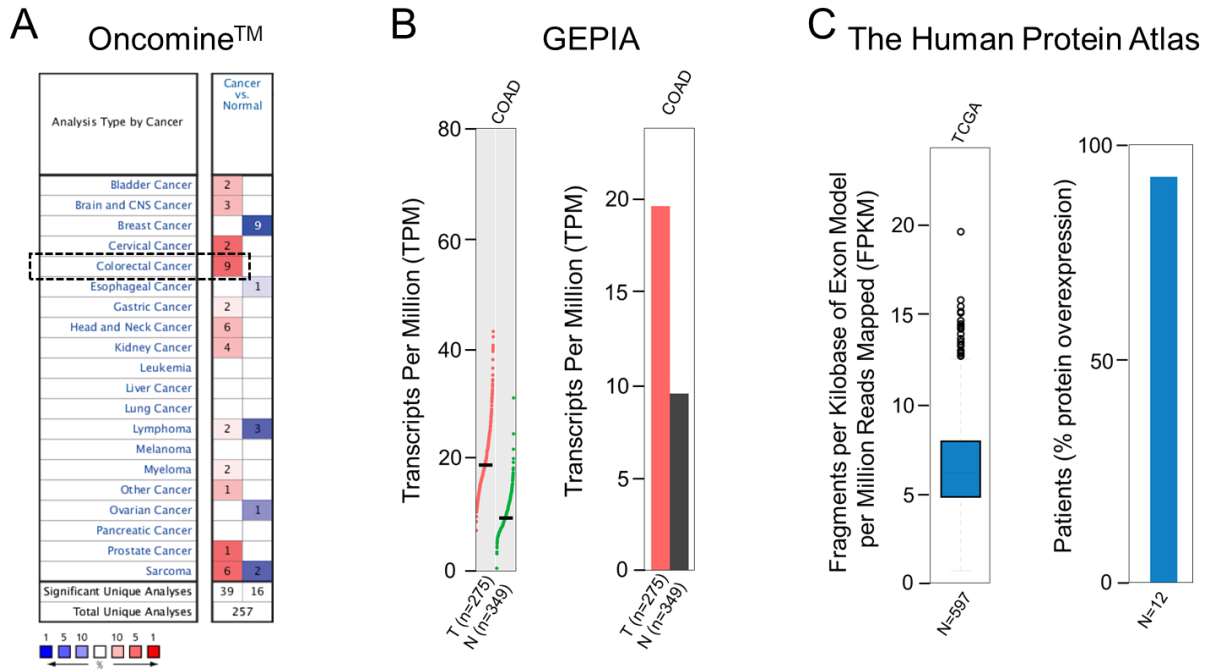
**Supplemental Figure 1. Zfp277, Ki67 and  $\beta$ -catenin expression in enteroids cultivated from WT and Zfp277<sup>-/-</sup> mice. (A) Representative light micrographs of enteroids, IF images of Zfp277, DAPI and merged confocal image of DAPI and Zfp277. (B and C) Representative confocal micrographs of enteroids stained with anti-Ki67 (B) and anti- $\beta$ -catenin antibodies (C) and merged images. 10X.**



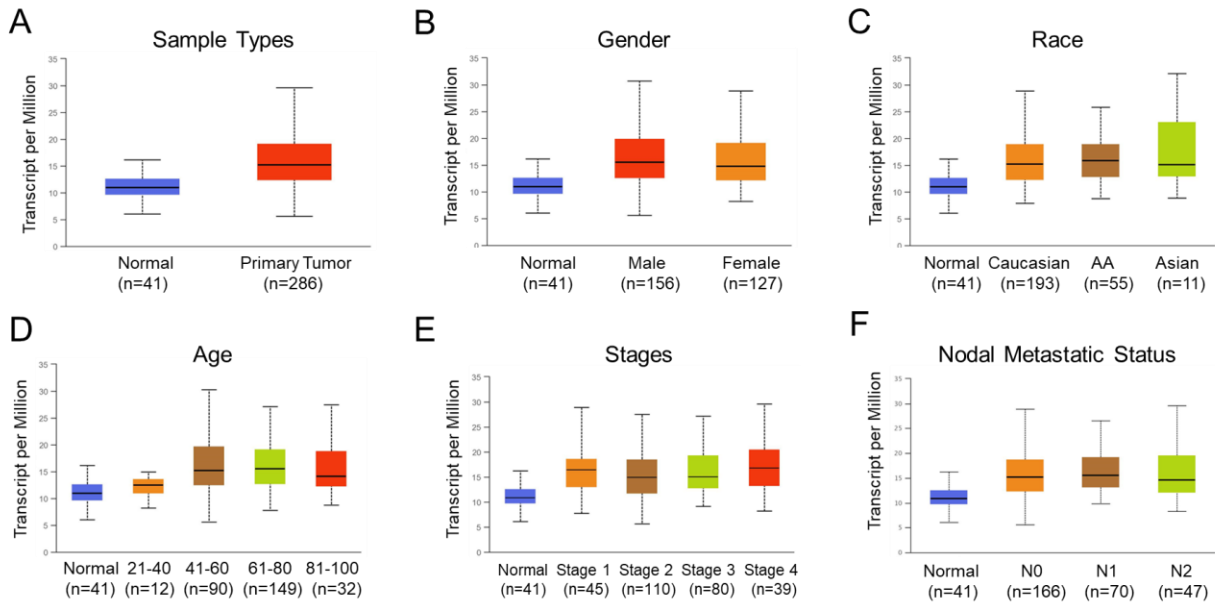
**Supplemental Figure 2. *Zfp277* transcript variant 1 and 2 levels in murine colon mucosa and small intestine enteroids. (A)** Variant 1 expression. Lanes 1 and 2, *Zfp277* mRNA RT-PCR product using cDNA prepared from colonic mucosa from two different WT mice. Lanes 3 and 4, *Zfp277* mRNA RT-PCR product using cDNA prepared from colon mucosa from two different *Zfp277*-deficient mice. Lane 5, *Zfp277* mRNA RT-PCR product using cDNA prepared from enteroids from WT mice. Lane 6, *Zfp277* mRNA RT-PCR product using cDNA prepared from enteroids from *Zfp277*-deficient mice. **(B)** Variant 2 expression. Lanes are the same as in A. **(C).** *Gapdh* RT-PCR. Lanes are the same as in A and B.



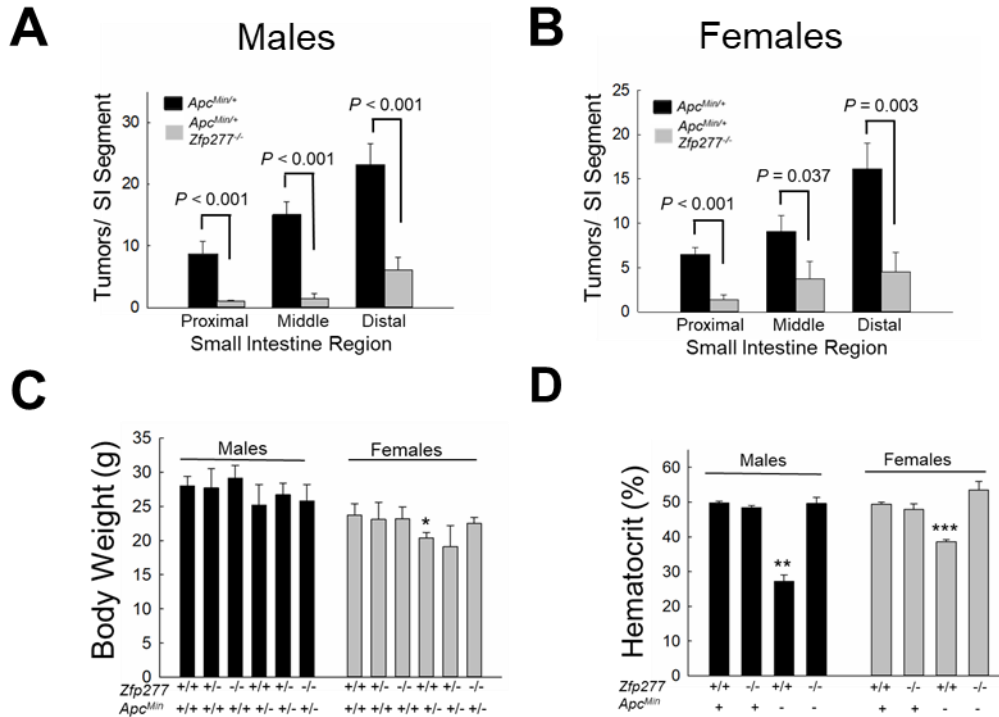
**Supplemental Figure 3. Characterization of *Zfp277* expression in transgenic mice. (A)** Representative *Zfp277* immunoblots using liver tissue extracts from WT, *Zfp277*<sup>+/-</sup>, *Zfp277*<sup>-/-</sup>, *Apc*<sup>Min/+</sup>, *Apc*<sup>Min/+</sup>*Zfp277*<sup>+/-</sup>, and *Apc*<sup>Min/+</sup>*Zfp277*<sup>-/-</sup> mice. **(B)** Representative H&E staining of colon sections from mice with the indicated genotypes. **(C)** Representative H&E staining of small intestine sections from mice with the indicated genotypes. **(D)** H&E staining of Swiss rolls of small intestines (left) and colons (right) from mice with the indicated genotypes (proximal section at center). Size bars, 100  $\mu$ M.



**Supplemental Figure 4. ZNF277 mRNA and protein overexpression in colon cancer reported in public cancer databases. (A)** Levels of *ZNF277* transcripts in colon cancer reported in the OncoPrint™ (A) and GEPIA (B) databases. (C) Levels of *ZNF277* protein in colon cancer reported in the Human Protein Atlas database.

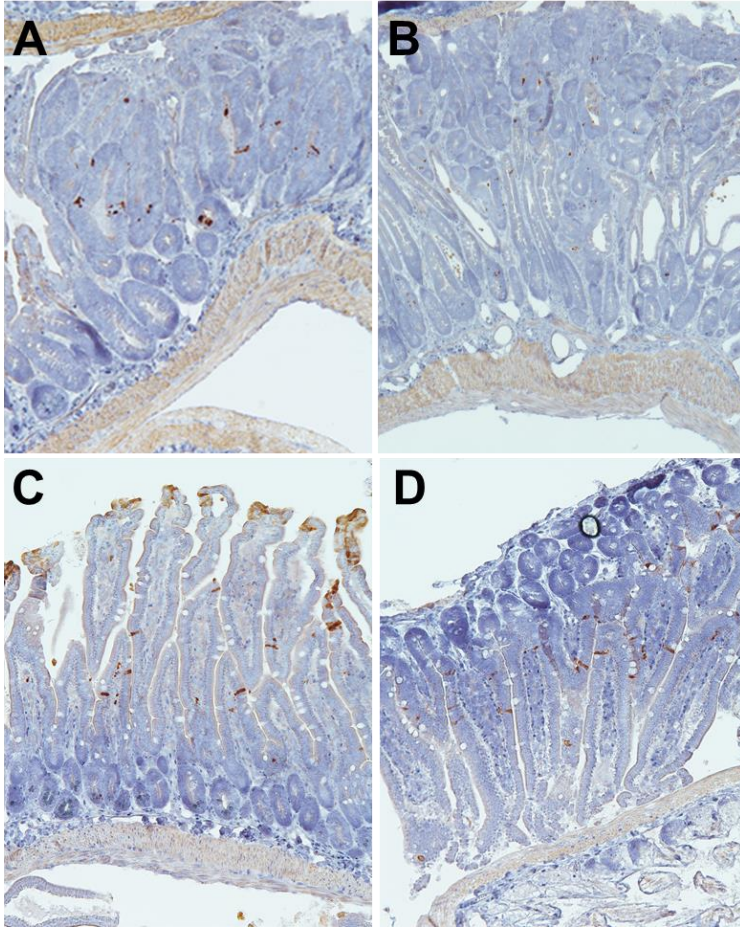


**Supplemental Figure 5. Box-whisker plots showing *ZNF277* mRNA overexpression in all stages of colon cancer compared with normal colon reported in the UALCAN server analysis. *ZNF277* transcript levels are increased in primary tumors (A), and in tumors regardless of gender (B), race (C), Age (D), colon cancer stages (E) and lymph node metastasis status (F). All  $P < 0.0001$ . Student's  $t$ -tests.**



**Supplemental Figure 6. Zfp277 deficiency attenuates small intestinal neoplasia, increases body weight and hematocrits in 15-week-old *Apc<sup>Min/+</sup>* mice. (A and B) Numbers of small intestinal tumors in the proximal, middle and distal thirds of the small intestine of *Apc<sup>Min/+</sup>* and *Apc<sup>Min/+</sup>Zfp277<sup>-/-</sup>* male (A) and female (B) mice. Male *Apc<sup>Min/+</sup>* (n=12); Male *Apc<sup>Min/+</sup>Zfp277<sup>-/-</sup>* (n=10); Female *Apc<sup>Min/+</sup>* (n=10); Female *Apc<sup>Min/+</sup>Zfp277<sup>-/-</sup>* (n=9). Values represent means  $\pm$  SEM. Data were analyzed using the one- or two-tailed *t*-test or the Mann-Whitney *U* test. (C) Body weights of mice with the indicated genotypes. (D) Hematocrits of mice with the indicated genotype. Values represent means  $\pm$  SD. One-tailed *t*-test. \*  $P < 0.05$  vs. female *Apc<sup>Min/+</sup>* mice; \*\*  $P < 0.001$  vs. *Zfp277<sup>-/-</sup>Apc<sup>Min/+</sup>* mice; \*\*\*  $P = 0.002$  vs. female *Apc<sup>Min/+</sup>* mice. Body weights were measured in at least six 15-week-old mice of each genotype. Hematocrits were measured in three mice of each genotype.**





**Supplemental Figure 7. *Zfp277* deficiency does not affect apoptosis. (A and B).** Images of cleaved caspase 3 IHC staining of small intestine adenomas from *Apc*<sup>Min/+</sup> (A) and *Apc*<sup>Min/+</sup>*Zfp277*<sup>-/-</sup> (B) mice. (C and D). Images of cleaved caspase 3 IHC staining of small intestines from *Apc*<sup>Min/+</sup> (C) and *Apc*<sup>Min/+</sup>*Zfp277*<sup>-/-</sup> (D) mice.

# Supplemental Table 1

## Antibodies used in this study

Antibody	Catalog Number	Lot Number	WB (h)	WB (m)	HRP IHC	IF	ChIP
ZNF277	INV-PA555577				✓	✓	
ZNF277	PTG-25094-1-AP	21820	✓	✓			
BM1	ABM-ab14389-25ug				✓	✓	
BM1	CST #6964S	3	✓				
P16ink4a/cdkn2a	RND-NB200-106SS				✓		
p16INK4A	92803T	1	✓				
p21WAF1	RND-NBP2-29463-20ug	1026-		✓			
p21WAF1	CST #2947	11	✓				
p27KIP1	CST 3686S	5	✓				
p14ARF	CST #2407S	5	✓				
p15ink4b/cdkn2b	RND-MAB6798-SP						
p57KIP2	CST-2557T	2	✓				
Keratin (KRT) 20	CST-13063S	1			✓	✓	
β-Catenin	PMG-610153	7187864	✓	✓			✓
Ki67	INV-MA514520				✓	✓	
Ki67	PMG-550609				✓	✓	
Ki67	HPA000451-100UL				✓	✓	
p19ARF/CDKN2A Antibody	NB200-106SS						
Cleaved Caspase-3	CST-9664S				✓	✓	
Goat Anti-rabbit IgG , Alexa Fluor 555	INV-A21429				✓	✓	
Goat Anti-mouse IgG, Alexa Fluor 555	INV-A21422				✓	✓	
Goat Anti-rabbit IgG, Alexa Fluor488	INV-A11034				✓	✓	
Goat Anti-mouse IgG,Alexa Fluor488	INV-A11029				✓	✓	
Donkey Anti-rabbit IgG, Alexa Fluor 555	INV-A31572				✓	✓	
Donkey Anti-goat IgG, Alexa Fluor 488	INV-A11055				✓	✓	
GAPDH	sc-47724	D2117	✓	✓			
β-actin	PTG-60008-1-Ig	10004413	✓	✓			
Lamin B1	CST-13435		✓				
RNA Polymerase II	Thermo Scientific						✓
rabbit IgG	Thermo Scientific						✓
Mouse IgG	Thermo Scientific						✓



## Supplemental Table 2

PCR Primer sequences

<b>Genes</b>	<b>Forward primer sequence 5' to 3'</b>	<b>Reverse primer sequence 5' to 3'</b>
<i>ZNF277</i>	ACAGCAGCAAGAACGAAATG	TGGCAATCCAATGTTGAAAG
<i>p21<sup>WAF1</sup></i>	GCGGTTGAATGAGAGGTTCC	AAGGAGAACACGGGATGAGG
<i>Zfp277</i>	TACTGTGCCAGGTAATGGCTG	TTCGGAAATCTGCAACGAGC
<i>Isoform 1</i>		
<i>Zfp277</i>	GTGCCAGGTAATGGCTGCTA	TGGCCATGTGGTTCAACAGT
<i>Isoform 2</i>		
<i>HOXD13</i>	CTTCGGCAACGGCTACTACA	CTTCTCCACGGGAAAGCCTC
<i>ZNF277</i>	AGATCACTAAAGCATGTTCCCC	TGGTTGGGGAAAGTGGAGAT
<i>ChIP 1</i>		
<i>ZNF277</i>	TGGCAAAGACTCCTCTGACA	AGTTTGACCTGAACCCACCT
<i>ChIP 2</i>		
<i>ZNF277</i>	GGAGGAGGAAAAGGAAGGGA	AGGAACAGGATAGCTCTCGC
<i>ChIP 3</i>		
<i>ZNF277</i>	TCACCCACGCCGATTTTCT	GGGACTGTGCATTGTTGTGA
<i>ChIP 4</i>		
<i>GAPDH</i>	CCCCATGGTGTCTGAGCG	CGACAGTCAGCCGCATCTT
<i>Gapdh</i>	GGTGAAGGTCGGTGTGAAC	TGATGGCAACAATCTCCACT
<i>(m)</i>		

## Supplemental Table 3

Differentially-expressed Zfp277 target genes identified by RNA-Seq using murine colonic mucosa. Increased expression in red and decreased expression in blue.

Zfp277 Gene Pathways identified by DAVID	Gene names
Anterior/posterior pattern specification	Hoxd9, Hhex, Hoxc8, Wnt3, Msx1, Sfrp1, Ifitm1, Sfrp2, Hoxd13, Gli2, Hoxd10, Hoxd11
Pathway in cancer	Flt3, Gli1, Gli2, Hhip, Rasgrp3, Adcy7, Bdkrb2, Cxcl12, F2r, Csf1r, Csf3r, Ccne1, Fgf18, Fzd6, Fzd9, Lama2, Lef1, Mmp2, Mmp9, Nos2, Pdgfra, Rxrg, Spi1, Wnt3, Wnt4, Wnt8b
MAPK/ERK1/2	Ren1, Grin2B, Tlr13, Gdf10, Pik3ap1, Bmp5, Bmp8b, Bmp6, Fgf18, Ccl2, Ccl21a, Ptpn22, Ccl8, Ccl5, Cd74, Ccl11, Ccl12, Ccl22, Gm10591, C1qtnf3, Pdgfra, Gpnmb, Pla2g5, Csf1r, F2r
Canonical Wnt signaling	Fz9, Fzd6, Wnt4, Wnt3, Wnt8b, Daam2, Sfrp1, Sfrp2, Lef1, Ptpu, Snai2, Gli1, Cam2b
Cytokine-cytokine receptor interaction	Relt, Ccl11, Ccl12, Ccl2, Ccl22, Ccl3, Ccl4, Ccl5, Ccl8, Ccr10, Ccr2, Ccr5, Ccr6, Cxcl12, Cxcl14, Cxcr5, Cx3cr1, Csf1r, Csf2rb2, Csf3r, Il10ra, Il18rap, Il2ra, Il6ra, Il7r, Il9r, Lep, Lifr, Ltb, Gm10591, Gm1987, Tnfrsf13b

## Supplemental Table 4

Differentially-expressed ZNF277 target genes identified by RNA-Seq in human HT29 cells. Increased expression in red and decreased expression in blue.

Top ZNF277 Gene Pathways and functional annotation clustering identified by DAVID	Gene names
Pathway in cancer	GNAI1, GNG4, KITLG, RASGPR2, WNT3A, WNT7A, WNT9A, WNT10B, WNT16, BMP4, F2R, COL4A3, COL4A4, COL4A5, COL4A6, CSF2RA, CCND1, CDKN1A (p21 <sup>WAF1</sup> ), CYCS, EDNRA, FGF2, FGFR1, LAMA4, LAMC3, LPAR1, LPAR3, PTGS2, RET, STAT1, SMO
Signaling pathways regulating pluripotency of stem cells	LHX5, SMAD9, WNT3A, WNT7A, WNT9A, WNT10B, WNT16, BMP4, BMPR2, DUSP9, ESRRB, FGF2, FGFR1, LIFR, ONECUT1, PAX6
Proteoglycans in cancer	TIAM1, WNT3A, WNT7A, WNT9A, WNT10B, WNT16, CAV1, CAV2, CCND1, CDKN1A, EZR, FGF2, FGFR1, FLNC, ITGA5, PLAU, SMO
PI3K-AKT signaling pathway	GNG4, KITLG, CREB3L1, F2R, COL4A3, COL4A4, COL4A5, COL4A6, COL6A1, COL6A2, CCND1, CDKN1A, FGF2, FGFR1, ITGA5, LAMA4, LPAR1, LPAR3, OSMR, PPKAA2, PPP2R2C, SGK1
HOX (Homeobox)	LHX2, LHX5, LHX6, LMX1B, MEIS2, NKX1-2, NKX2-8, NKX6-1, TLX3, GBX2, HOXD8, IRX3, IRX4, MKX, MSX1, ONECUT1, ONECUT3, PAX6, PITX3, PRRX2, SHOX2, ZFHX4