Fig. S1



Fig. S1. Intestinal *Gata6* deletion alters gene networks controlling cell proliferation in the mature ileum. Network analyses on microarray data of *Ctl* and *Gata6* ΔIE ileum (n=3 in each group) revealed (A) an increase in targets of the tumor surpressor gene p53, and (B) a decrease in targets of the proto-oncogene c-MYC. (C) Legend defining symbols used in the networks. Arrows indicate the direction of the interaction. Red circles = up-regulated transcripts; Blue circles = down-regulated transcripts. Differentially expressed transcripts were determined at the 5% FDR level using Significance Analysis of Microarrays (SAM) and interaction networks were developed from the differentially expressed transcripts using Metacore.



Fig. S2. *Gata6* and *Spdef* mRNA abundance in ileum in each group of mice. *Gata6* Δ *IE* and *DKO* mice had significantly lower *Gata6* mRNA abundances than *Ctl* and *SpdefKO* mice. *SpdefKO* and *DKO* mice had significantly lower *Spdef* mRNA abundances than *Ctl* and *Gata6* Δ *IE* mice.

A qRT-PCR primers

hGata6:	Sense:	5' -GACTTGCTCTGGTAATAG-3'
	Antisense:	5' -CTGTAGGTTGTGTTGTGG-3'
hSpdef:	Sense:	5' -AACATCACCGCAGATCCC-3'
	Antisense:	5' -CGGTATTGGTGCTCTGTC-3'
hGapdh:	Sense:	5' -AAGTATGACAACAGCCTC-3'
	Antisense:	5' -ATGAGTCCTTCCACGATA-3'
mGata6:	Sense:	5' -CAGCAAGCTGTTGTGGTC-3'
	Antisense:	5' -GTCTGGTACATTTCCTCCG-3'
mSpdef:	Sense:	5' -CAACAAGGAGAAAGGCATCTT-3'
	Antisense:	5' -GCGGCTTAGTTTATCATAGTTCA-3'
mChga:	Sense:	5' -CACAGCCACCAATACC-3'
	Antisense:	5' -TCTTCCTCCTCCTCTTC-3'
mNeurog3:	Sense:	5' -CTCAGCAAACAGCGAAGAAG-3'
	Antisense:	5' -GGGAAGGTGGGCAGGAC-3'
mLyz:	Sense:	5' -ACTCCTCCTTGCTTTCTGTC-3'
-	Antisense:	5' -GTCGGTGCTTCGGTCTC-3'
mMuc2:	Sense:	5' -AACTACCACTGTGATGCCAATG-3'
	Antisense:	5' -ACAATGTTGATGCCAGACTCG-3'
mApoa1:	Sense:	5' -GGACTTCTGGGATAACCT-3'
	Antisense:	5' -GCACCTTCTGTTTCACTT-3'
mCar1:	Sense:	5' -AACAGAATTATGTCAGTGCTAA-3'
	Antisense:	5' -AGAGAATGAATCACTTAGTTGTAA-3'
mGapdh:	Sense:	5' -GCCTTCCGTGTTCCTACCC-3'
	Antisense:	5' -TGCCTGCTTCACCACCTTC-3'

B ChIP primers

<i>hSpdef</i> (-40 k	b), region 1: Sense: Antisense:	5' -AAAGGTCTTCCTCCAAAATG-3' 5' -CAAAGCAGTAAGACTTGGAG-3'
<i>hSpdef</i> (-40 k	b), region 2: Sense: Antisense:	5' -CAGTTTGAATCCATCCTCC-3' 5' -CCTATCTTCAAAGCATTTACTATCT-3'
<i>hSpdef</i> (-40 k	b), region 3: Sense: Antisense:	5' -CTATGGATTGGTCACATCTAC-3' 5' -GTTTGAGGGTCTTGAACTT-3'
Amy1 TSS:	Sense: Antisense:	5' -GGAAAATAAAGGGTTGGAGC-3' 5' -AGTGAATCATGTCAGTATAACAA-3'

C Genotype primers

Gata6:	Sense: Antisense:	5' -GTGGTTGTAAGGCGGTTTGT-3 ' 5' -ACGCGAGCTCCAGAAAAAGT-3'
Cre:	Sense: Antisense:	5' -CGTATAGCCGAAATTGCCAG -3' 5' -CAAAACAGGTAGTTATTCGG-3'
Spdef:	WTSense: Antisense: KO Sense:	5' -CCCACCTCCTATGTCAGCCATGGC-3' 5' -CAATCCTGTACCATATCTGGCATGG-3' 5' -GCATCGCATTGTCTGAGTAGGTGTCA-3'

Fig. S3. Primers used for: (A) qRT-PCR, (B) ChIP assays, and (C) genotyping.



Fig. S4. Timeline for study. Mice 4 wks of age were given Tamoxifen as indicated (*black circle*) beginning on Day 0, and an injection of BrdU 2 hr before tissue collection at Day 28.