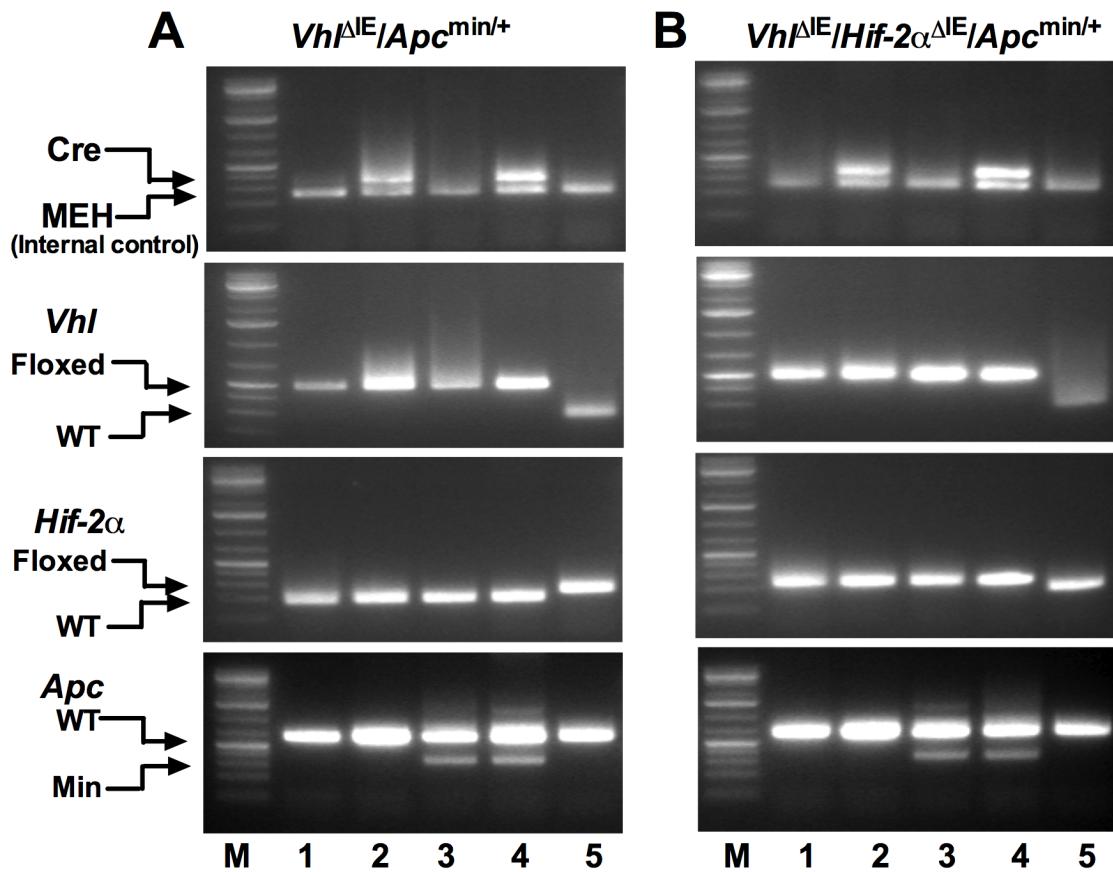
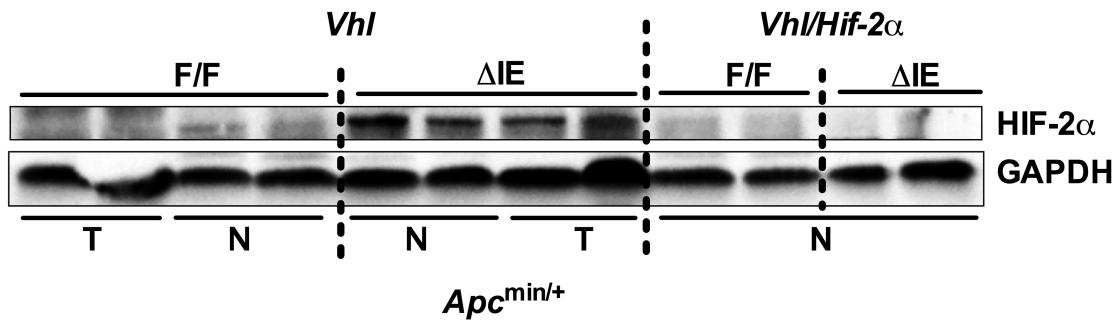


Supplemental Data: Figures 1-5 and Tables 1-3



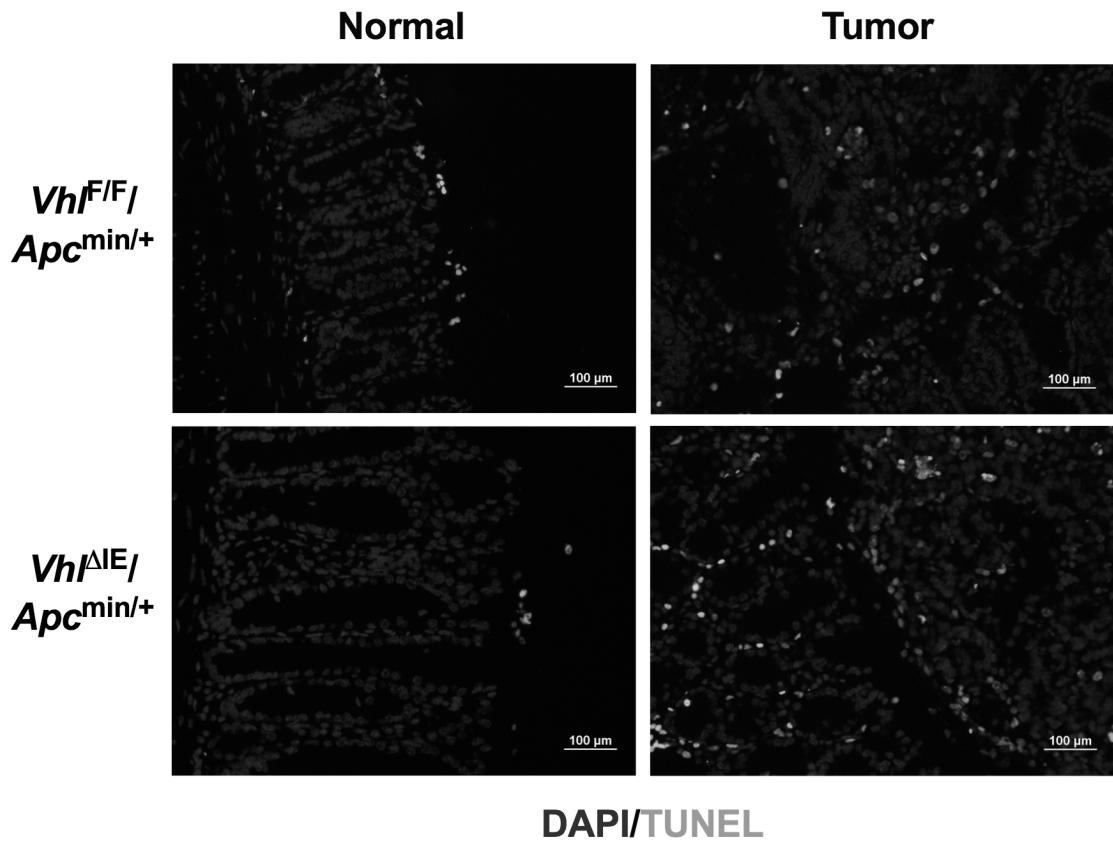
Supplemental Figure 1 Representative genotyping for *Vhl*^{ΔIE}/*Apc*^{min/+} and *Vhl*^{ΔIE}/*Hif-2α*^{ΔIE}/*Apc*^{min/+} mouse lines.

(A) Tail DNA was extracted and genotyping for *Cre*, *Vhl*, *Hif-2α* and *Apc* were performed in *Vhl*^{ΔIE}/*Apc*^{min/+} mouse line. M, marker; 1, *Vhl*^{F/F}; 2, *Vhl*^{ΔIE}; 3, *Vhl*^{F/F}/*Apc*^{min/+}; 4, *Vhl*^{ΔIE}/*Apc*^{min/+}; 5, *Hif-2α*^{ΔIE}. (B) Tail DNA was extracted and genotyping for *Cre*, *Vhl*, *Hif-2α* and *Apc* were performed in *Vhl*^{ΔIE}/*Hif-2α*^{ΔIE}/*Apc*^{min/+} mouse line. M, marker; 1, *Vhl*^{F/F}/*Hif-2α*^{F/F}; 2, *Vhl*^{ΔIE}/*Hif-2α*^{ΔIE}; 3, *Vhl*^{F/F}/*Hif-2α*^{F/F}/*Apc*^{min/+}; 4, *Vhl*^{ΔIE}/*Hif-2α*^{ΔIE}/*Apc*^{min/+}; 5, *Vhl*^{F/F}.



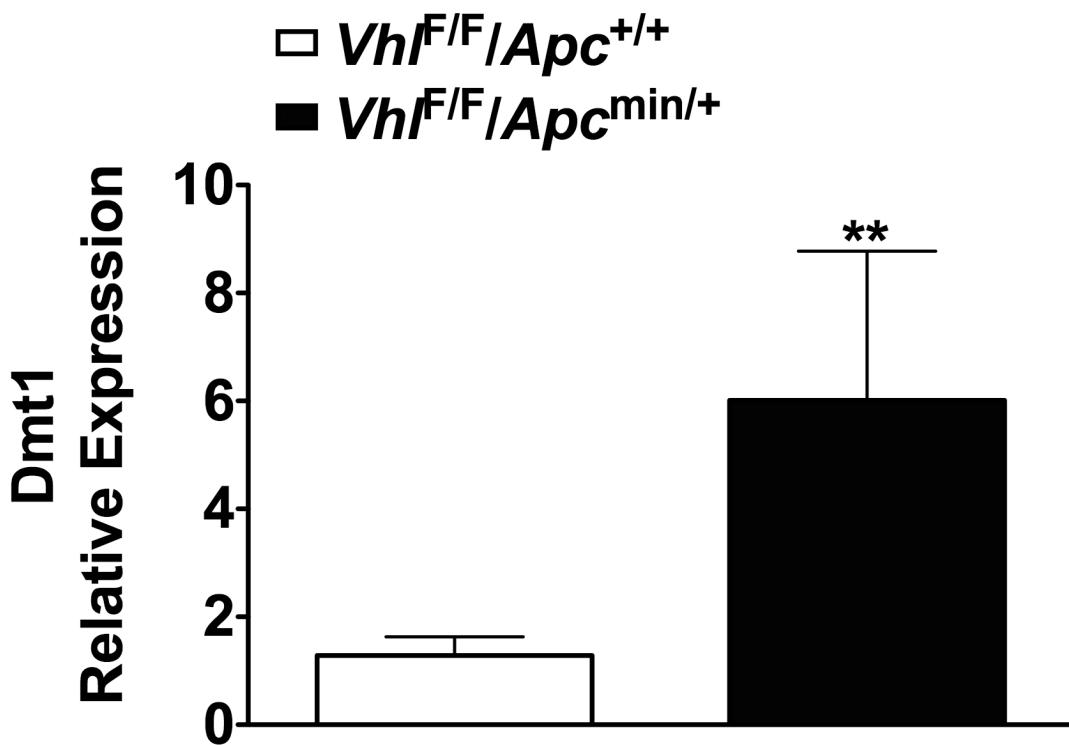
Supplemental Figure 2 HIF-2 α Expression in the colons of *Vhl*^{ΔIE}/*Apc*^{min/+}, *Vhl*^{ΔIE}/*Hif-2 α* ^{ΔIE}/*Apc*^{min/+} and their wild-type littermate control mice.

Western Blot analysis for HIF-2 α and GAPDH from normal and tumor colon tissues of 6-month-old *Vhl*^{F/F}/*Apc*^{min/+}, *Vhl*^{ΔIE}/*Apc*^{min/+}, *Vhl*^{F/F}/*Hif-2 α* ^{F/F}/*Apc*^{min/+} and *Vhl*^{ΔIE}/*Hif-2 α* ^{ΔIE}/*Apc*^{min/+} mice. N, normal tissue; T, tumor tissue.



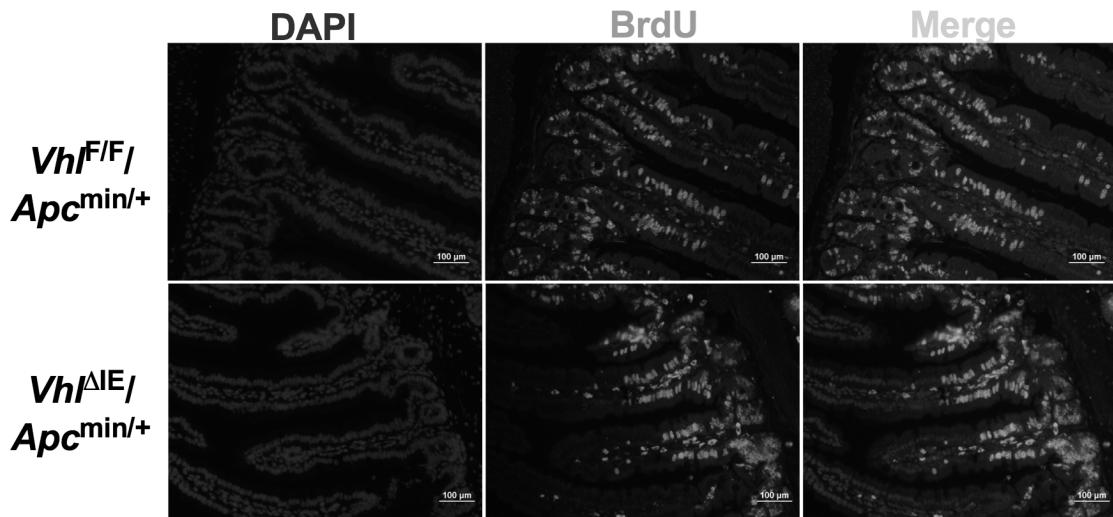
Supplemental Figure 3 TUNEL assay in the normal and tumor colons of *Vhl*^{F/F}/*Apc*^{min/+} and *Vhl*^{ΔIE}/*Apc*^{min/+} mice.

TUNEL assay for the normal and tumor colons of 6-month-old *Vhl*^{F/F}/*Apc*^{min/+} and *Vhl*^{ΔIE}/*Apc*^{min/+} mice.



Supplemental Figure 4 DMT-1 Expression in the small intestine of $Vhl^{F/F}/Apc^{min/+}$ mice.

qPCR analysis in the small intestine of $Vhl^{F/F}/Apc^{min/+}$ and $Vhl^{F/F}/Apc^{+/+}$ mice. Each bar represents the mean value \pm S.D. ** $p < 0.01$ compared to $Vhl^{F/F}/Apc^{+/+}$ mice.



Supplemental Figure 5 Disruption of *Vhl* in the small intestine does not increase proliferation.

BrdU staining of small intestine epithelial cells from 3-month-old *Vhl*^{F/F}/*Apc*^{min/+} and *Vhl*^{ΔIE}/*Apc*^{min/+} mice.

Supplemental Table 1 Primers List

QPCR Primer	Sequence (5'-3')
β-actin F	TATTGGCAACGAGCGGTTCC
β-actin R	GGCATAGAGGTCTTACGGATGT
Cndl F	GGGTGGGTTGGAAATGAAC
Cndl R	TCCTCTCCAAAATGCCAGAG
Dmt1 F	TGTTTGATTGCATTGGGTCTG
Dmt1 R	CGCTCAGCAGGACTTCGAG

Supplemental Table 2 List of top 10 changed genes from the colons of *Vhl*^{F/F} vs *Vhl*^{ΔIE} and *Vhl*^{F/F}/*Apc*^{min/+} vs *Vhl*^{ΔIE}/*Apc*^{min/+}.

Symbol	Description	Microarray*	Oncomine#
Xist	inactive X specific transcripts	Decrease	Decrease (5)
Myl7	myosin, light polypeptide 7, regulatory	Increase	No Change
Mpp4	membrane protein, palmitoylated 4 (MAGUK p55 subfamily member 4)	Increase	No Change
Nrn1	neuritin 1	Increase	Decrease (1)
Ndufa4l2	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 4-like 2	Increase	Increase (3)
Slc11a2	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2	Increase	Increase (19)
Als2cr4	amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 4	Increase	Increase (14), Decrease (1)
Dnase1l2	deoxyribonuclease 1-like 2	Increase	Increase (4)
Ankrd37	ankyrin repeat domain 37	Increase	Increase (1) Decrease (2)
Steap4	STEAP family member 4	Increase	Decrease (4)

* Gene changes in *Vhl*^{ΔIE} compared to *Vhl*^{F/F} mice.

Numbers in parenthesis indicate independent microarray reports for gene changes in colon cancer compared to normal tissue.