#### Supplemental figure legends

Supplemental Figure 1. Functional TRPV1 expression in IEC and intestinal organoids. (A) PCR products generated with primers specific for Trpv1-6 and Gapdh (housekeeping gene) after 30 cycles of amplification, corresponding to Q-PCR results presented Figure 1A. The mRNA was isolated from primary IEC samples and used for cDNA synthesis. (B) PCR products after 30 cycles of amplification of mRNA obtained from small intestinal organoids, corresponding to Figure 1B. (C) Expression of *Trpv1* transcripts in intestinal organoids and freshly isolated IEC, respectively, derived from WT and  $Trpv1^{-/-}$  mice. Cldn6 (claudin-6), an epithelial cell marker. Gapdh, housekeeping gene. (D) Expression of TRPV1 mRNA in human colorectal cancer cell lines (HCT116 and HCA-7) and rodent cell lines (rat small intestinal cell line, IEC-6; murine colorectal cancer cell line, CMT-93). Positive control: homogenates from mouse hippocampus (CNS). (E) Validation of anti-TRPV1 Abs for Western blotting with cell lysates from CHO<sup>K1</sup> and TRPV1 overexpressing CHO<sup>TRPV1</sup> cells.  $\beta$ -Actin: loading control. (F) Single cell suspensions of CHO<sup>K1</sup> and CHO<sup>TRPV1</sup> cells were stained with anti-TRPV1 Abs (red lines) and detected with rabbit anti-goat AF488 conjugate Abs (flow cytometry). IgG control: goat IgG (black lines). (G) Validation of functional TRPV1 activity recordings. CHO<sup>TRPV1</sup> cells were loaded with 5  $\mu$ M Fura-2 AM and  $[Ca^{2+}]_{cvt}$  were measured in single cells during perfusion with PSS at room temperature (20 °C). Left panel: Stimulation with 1 nM capsaicin induced rapid [Ca<sup>2+</sup>]<sub>cvt</sub> elevations. **Right panel:** Pretreatment of CHO<sup>TRPV1</sup> cells with TRPV1 antagonist BCTC (1  $\mu$ M) prevented [Ca<sup>2+</sup>]<sub>cvt</sub> elevations induced by capsaicin (1 nM), but not ATP (10  $\mu$ M). Mean  $\pm$  SEM (n=52 for each condition). (H) Functional TRPV1 channel activity in the HCT116 cell line. Fura-2 loaded HCT116 cells were stimulated with increasing concentrations of capsaicin,

resulting in dose-dependent  $[Ca^{2+}]_{cyt}$  increases (left panel). These capsaicin-induced  $Ca^{2+}$  responses were attenuated by pretreatment with TRPV1 antagonist BCTC (10  $\mu$ M) (right panel).

Supplemental Figure 2. Functional effects of TRPV1 in IEC biology. (A) Gross crypt-villus architecture, Goblet cell and Paneth cell differentiation were unaffected in small intestines of Trpv1<sup>-/-</sup> mice. PAS staining (top panels) and immunostaining for matrix metalloprotease 7 MMP7 (bottom panels), respectively. (B) Numbers of Goblet cells and Paneth cells were similar in Trpv1--- and WT intestinal organoids. Immunostaining of fixed organoids in Matrigel with AF488 conjugated wheat germ agglutinin (WGA) (top panels) or anti-MMP7 Abs (bottom panels), Phalloidin AF546 and DAPI and imaged by confocal fluorescent microscopy. (C) Capsaicin treatment inhibited ligand-induced EGFR<sup>Tyr1068</sup> phosphorylation. WT mice were pretreated with regular chow or capsaicin 0.01% (w/w) mixed in chow for 2 days and then treated with EGF (2  $\mu$ g, i.p.) for 0 or 15 min. IEC lysates were analyzed by Western blotting. (D) WT mice received dietary capsaicin 0.01% (w/w) in chow or control chow for 4 weeks. Tissues were harvested and Ki67 staining with DAB detection of colon tissues was performed. Shown are representative pictures of colons used for Ki67 quantification. (E) Quantification of Ki67 positive cells of mice treated with dietary capsaicin or control diet (n=3 mice/group). (F) WT mice were treated with vehicle (EtOH), low dose capsaicin (0.3 mg/kg/day) or high dose capsaicin (3.0 mg/kg/day) for 5 consecutive days by gavage. Small intestinal tissues were harvested on day 5. Intestinal sections were stained with Alcian blue (counterstaining with nuclear fast red; top panels), or with anti-MMP7 Abs (counterstaining with hematoxylin; bottom panels). (G) Quantification of the area stained for mucin (Alcian blue) with ImageJ by using the Colour Deconvolution plugin (n=3 mice/group). (H) Quantification of the area

immunostained for MMP7 by immunohistochemistry with DAB (n=3 mice/group). Data are mean  $\pm$  SEM. \**P*<0.05 (t-test). Scale bars = 100  $\mu$ m.

Supplemental Figure 3. Analysis of transgenic TRPV1<sup>IEC</sup> mice. (A) Validation of conditional TRPV1 overexpression by PCR. Small intestinal and colonic IEC were harvested from TRPV1<sup>fl/fl</sup> and TRPV1<sup>IEC</sup> (Cre positive) mice, together with spinal cord homogenates. PCR was performed for Cre, Trpv1, a neuronal cell marker (Nefl; Neurofilament), an epithelial cell marker (Slc26a3), and housekeeping gene Gapdh. The results showed endogenous expression of Trpv1 in all tissues analyzed (TRPV1<sup>fl/fl</sup>), whereas overexpression was only found in IEC lysates from Cre positive (TRPV1<sup>IEC</sup>) mice. (B) Validation of conditional TRPV1 overexpression by Western blotting. Protein samples (IEC and spinal cord homogenates) were prepared for IB. Membranes were probed for TRPV1, a neuronal marker (Synaptophysin), an IEC marker (Occludin) and loading control β-Actin. TRPV1 overexpression was found only in IEC in Cre positive (TRPV1<sup>IEC</sup>) mice. (C) Reduced IEC proliferation in colons from TRPV1<sup>IEC</sup> mice. Representative pictures of Ki67+ cells in colon sections of TRPV1<sup>fl/fl</sup> and TRPV1<sup>IEC</sup> mice are shown. (D) Quantification of Ki67+ cells in colons of TRPV1<sup>fl/fl</sup> and TRPV1<sup>IEC</sup> mice, respectively. Data are shown as mean  $\pm$  SEM (n=3). (E) RNA from colonic IEC isolated from TRPV1<sup>fl/fl</sup> and TRPV1<sup>IEC</sup> mice were used for Q-PCR analysis for the expression of a proliferation marker, *Mki67*, normalized to *Gapdh*. Data are shown as mean  $\pm$  SEM (n=3). (F) Reduced expression of a proliferation marker, Mki67, and EGFR target genes, c-Fos, c-Jun and Egr1 in intestinal organoids generated from TRPV1<sup>IEC</sup> mice. RNA from organoids was used for Q-PCR analysis, expression data was normalized to Gapdh. Mean  $\pm$  SD (n=2). (G) Reduced staining intensity for

p-EGFR<sup>Y1068</sup> in TRPV1<sup>IEC</sup> compared to TRPV1<sup>fl/fl</sup> organoids. Scale bar = 100  $\mu$ m. \**P*<0.05 (t-test) in D, E and F.

Supplemental Figure 4. Functional association between TRPV1 and EGFR signaling in vitro. (A) TRPV1 inhibits ligand-induced EGFR<sup>Y1068</sup> phosphorylation. CHO<sup>K1</sup> cells were transfected with empty vector, wt-EGFR, or wt-EGFR + TRPV1. Cells were then stimulated with 10 ng/mL EGF for indicated times, followed by Western blot analysis of total cell lysates. (B) Detailed time course of EGF-stimulated HCT116 cells transfected with empty vector or TRPV1. Total cell lysates were analyzed for multiple EGFR phospho-sites including Y992, Y1045, S1046/S1047, and Y1068 by Western blotting. TRPV1 overexpression suppressed Y992, Y1045, and Y1068 phosphorylation, whereas it potentiated phosphorylation of S1046/S1047. The S1046/S1047 residues are known to be phosphorylated by  $Ca^{2+}/calmodulin-dependent$ kinase II (CaMKII), suggesting activation of CaMKII by TRPV1 under these experimental conditions. Representative results of three independent experiments are shown. (C) Transfection of HCT116 cells with empty vector, TRPV3, or TRPV4-FLAG plasmids. Three days after transfection, cells were stimulated with EGF (1 ng/mL) and total cell lysates were analyzed by Western blotting for indicated proteins. (D) HCT116 cells were transfected with empty vector or TRPV5-GFP construct and analyzed by widefield fluorescent microscopy on day 3. (E) Mock or TRPV5-GFP transfected HCT116 cells were stimulated with EGF (1 ng/mL) followed by Western blot analysis of total cell lysates. Results are representative of three independent experiments.

Supplemental Figure 5. Physiological effects of TRPV1 signaling in intestinal organoids. (A) Enhanced TA zone in *Trpv1<sup>-/-</sup>* crypts in intestinal organoids. WT and *Trpv1<sup>-/-</sup>* organoids, as shown in Figure 4E (cropped, dotted lines), were generated from small intestines and cultured under similar conditions. Ki67 staining was performed 3 days after passing, together with Phalloidin AF546 and DAPI. Scale bars = 100  $\mu$ m. (B) Ki67 staining of WT and *Trpv1<sup>-/-</sup>* organoids at the same growth stage by immunohistochemistry with DAB (brown) and hematoxylin counterstaining. Representative examples are shown. (C) Quantification of Ki67+ cells (%) relative to hematoxylin staining nuclei with ImageJ (ImmunoRatio plugin). Mean  $\pm$  SEM (n=7). \**P*<0.05 (t-test). (D) Proliferation in EGF-starved WT and *Trpv1<sup>-/-</sup>* organoids similar to those shown in Figure 4, G and H at lower magnification. Scale bars = 100  $\mu$ m.

Supplemental Figure 6. Identification of PTP1B activity as a downstream effector mechanism of TRPV1. (A) Validation of the CaM inhibitor, W-7, as a blocker of the downstream CaMKII pathway. HCT116 cells were pretreated with 0 or 10  $\mu$ M W-7 for 1 hr. Cells were then stimulated with 5  $\mu$ M ionomycin for 0, 10, or 30 min. Total cell lysates were analyzed by Western blotting for p-CaMKII, total CaMKII, and  $\beta$ -actin. W-7 pretreatment efficaciously inhibited constitutive and ionomycin-induced CaMKII phosphorylation. (B) Calmodulin (CaM) does not play a role in TRPV1-mediated effects on EGFR signaling. HCT116 cells were transfected with control or TRPV1 plasmid, followed by stimulation with EGF (1 ng/mL) for indicated times. CHO<sup>K1</sup> and CHO<sup>TRPV1</sup> cells were transfected with control or wt-EGFR plasmid, followed by EGF stimulation as indicated. Pre-treatment with 10  $\mu$ M W-7 (1 hr) before stimulation failed to reverse TRPV1-mediated inhibition of EGFR signaling in both HCT116 (left panel) and CHO cells (right panel). (C) Validation of FK506 as a potent inhibitor

of the Ser phosphatase activity of calcineurin in HCT116 cells. Cells were pretreated with 0, 1, or 10 µM FK506 (1 hr), followed by stimulation with or without 5 µM ionomycin (30 min). Total cell lysates were analyzed for calcineurin target, NFAT1, by using antibodies that recognize total and phosphorylated NFAT1. (D) Calcineurin inhibitor, FK506 (10 µM), did not reverse the inhibition of EGFR phosphorylation by TRPV1. The same experimental approach was applied with HCT116 and CHO cells, respectively, as in **B**. (E) Quantification of expression levels of PTP candidates in WT versus  $Trpv1^{-/-}$  colon crypts by Q-PCR, normalized for *Gapdh*. Results are shown as mean  $\pm$  SEM. (F) Dephosphorylation of PTP1B target Hrs in TRPV1 overexpressing cells. HCT116 cells were transfected with empty vector or TRPV1 plasmid. Two days later, cells were stimulated with EGF (1 ng/mL) for indicated time periods, followed by Western blot analysis with antibodies directed to phospho-EGFR<sup>Y1068</sup>, total EGFR, phospho-Hrs<sup>Y334</sup>, or β-actin. (G) Transfection of HCT116 cells with empty vector, WT PTP1B (PTP435), or PTP1B truncation mutants (PTP370, PTP377). Total cell lysates were analyzed by IB with anti-HA tag mAbs. (H) Validation of IEC fractionation. Q-PCR analysis showed enrichment for Lgr5, Mmp7, and Egr1 in the crypt fraction (normalized for Gapdh). Conversely, mature enterocyte marker *Slc26a3* (a chloride anion exchanger) expression was relatively enriched in villus fractions. Data are mean  $\pm$  SEM (n=3/group).

**Supplemental Figure 7. Functional association between TRPV1 and EGFR signaling pathways**. (A) Validation of PLCγ1 knockdown in control and *PLCG1* siRNA transfected cells by Western blotting. (B) Model for the triggering of TRPV1 by EGFR, as explained in text. DAG, diacylglycerol; IP<sub>3</sub>, inositol trisphosphate; PIP<sub>2</sub>, phosphatidylinositol-4,5-bisphosphate; PLC, phospholipase C.

Supplemental Figure 8. TRPV1 deficiency on the Apc<sup>min/+</sup> background. (A) Enhanced expression of *c-Fos* mRNA in mice with hetero- or homozygous deletion of *Trpv1* on the Min background as determined by PCR. (B) Increased expression of *c-Fos* in  $Apc^{min/+}/Trpv1^{+/-}$ (wt/ko) and Apc<sup>min/+</sup>/Trpv1<sup>-/-</sup> (ko/ko), compared to Apc<sup>min/+</sup>/Trpv1<sup>+/+</sup> (wt/wt) mice. Q-PCR was performed with IEC lysates from age-matched mice and data were normalized for *Gapdh*. (C) Systemic ablation of TRPV1+ sensory neurons in  $Apc^{min/+}$  mice did not recapitulate the phenotype of  $Apc^{min/+}/TrpvI^{-/-}$  mice.  $Apc^{min/+}$  mice were treated with 50 µg/kg resiniferatoxin (RTX) s.c. on day 1, 2, and 7 after birth. Ablation of TRPV1+ neurons was validated by a negative eye wipe test by ocular challenge with 0.01% (w/v) capsaicin in saline between 6-8 weeks. The cohort of  $Apc^{min/+}$  mice (n=16) is the same as shown in Figure 7D for direct comparison with  $Apc^{min/+}$  RTX mice (n=11). Median survival of  $Apc^{min/+}$  (control) and  $Apc^{min/+}$ RTX mice were 23.2 and 24 weeks, respectively. \*P=0.0397 (Log-rank test). (D) Left panel: Polyp counts (>1 mm) of  $Apc^{min/+}$  (control) and  $Apc^{min/+}$  RTX mice (n=12 and n=11, respectively). Control cohort is the same as shown in Figure 7B. Right panel: Hemoglobin levels of control and RTX-treated  $Apc^{min/+}$  mice (n=10 and n=18, respectively) at 20 weeks. Control cohort is the same as shown in **Figure 7C**.

Supplemental Figure 9. TRPV1 vs. TRPV5 in the regulation of the EGFR. The proposed mechanism of TRPV1-mediated feedback on EGFR signaling, which is initiated by PLC $\gamma$  activation and PIP<sub>2</sub> depletion. This mechanism is likely distinct from that mediated by TRPV5.

#### Supplemental Tables

#### Supplemental Table 1: Mouse primers.

Gene	Official full name	Accession No.	Sequence 5'-3'
Axin2	axin2	NM_015732.4	CAGTGAGCTGGTTGTCACCT
			CTGAGCTGCTCCTTGAAGTG
c-Fos	FBJ osteosarcoma	NM_010234.2	ACCATGATGTTCTCGGGTTTC
	oncogene		GCTGGTGGAGATGGCTGTCAC
c-Jun	Jun oncogene	NM_010591	GCAGAAAGTCATGAACCACG
			GCAACCAGTCAAGTTCTCAAG
с-Мус	myelocytomatosis	NM_001177353.1	TGGTGTCTGTGGAGAAGAGGCAAA
	oncogene	NM_010849.4	TTGGCAGCTGGATAGTCCTTCCTT
Cldn6	claudin 6	NM_018777.4	GGATGTCCTGTGTGGTTCAG
			AGGACAATGAGGAGGGTGAC
Egrl	early growth response 1	NM_007913	GTGTGCCCTCAGTAGCTTC
			GACATCAATTGCATCTCGGC
Fosl2	fos-like antigen 2	NM_008037.4	CGGGAACTTTGACACCTCG
(Fra-2)			TGATGGCGTTGATTGTGGG
Gapdh	glyceraldehyde-3-	NM_008084	TCAACAGCAACTCCCACTCTT
	phosphate dehydrogenase		ACCCTGTTGCTGTAGCCGTAT
Lgr5	leucine rich repeat	NM_010195.2	TGAGCGGGACCTTGAAGATTTCCT
	containing G protein		AGCCAGCTACCAAATAGGTGCTCA
	coupled receptor 5		
Mki67	antigen identified by	NM_001081117.2	TGCCCGACCCTACAAAATG
	monoclonal antibody Ki 67		GAGCCTGTATCACTCATCTGC
Mmp7	matrix metallopeptidase 7	NM_010810	AACACTCTAGGTCATGCCTTCGCA
			AGACCCAGAGAGTGGCCAAATTCA
Nefl	neurofilament, light	NM_010910	GGCCTTGGACATCGAGATTG

	polypeptide		TCTGCAAGCCACTGTAAGC
Ocln	occludin	NM_008756.2	TCCACACTCAAGGTCAGAGG
			CAATGGCCTACTCCTCCAAT
Slc26a3	solute carrier family 26,	NM_021353.3	AGGGAATGCTGATGCAGTTTGCTG
	member 3		AGTTGAAATGCTACACTTGCCGCC
Trpv1	transient receptor potential	NM_001001445	AGCTGCAGCGAGCCATCACCA
	cation channel, subfamily		ATCCTTGCCGTCCGGCGTGA
	V, member 1		
Trpv2	transient receptor potential	NM_011706	CCAGCCATTCCCTCATCAAAA
	cation channel, subfamily		AAGTACCACAGCTGGCCCAGTA
	V, member 2		
Trpv3	transient receptor potential	NM_145099	TGAAAGAAGGCATTGCCATTT
	cation channel, subfamily		GAAACCAGGCATCTGACAGGAT
	V, member 3		
Trpv4	transient receptor potential	NM_022017	TCACCTTCGTGCTCCTGTTG
	cation channel, subfamily		AGATGTGCTTGCTCTCCTTG
	V, member 4		
Trpv5	transient receptor potential	NM_001007572	CGTTGGTTCTTACGGGTTGAAC
	cation channel, subfamily		GTTTGGAGAACCACAGAGCCTCTA
	V, member 5		
Тгрvб	transient receptor potential	NM_022413	ATCCGCCGCTATGCACA
	cation channel, subfamily		AGTTTTTCTCCTGAATCTTTTTCCA
	V, member 6		

#### Supplemental Table 2: Other primers.

Gene	Accession number	Sequence 5'-3'
hGAPDH	NM_002046.3	ACCAAATCCGTTGACTCCGAC
		TTCGACAGTCAGCCGCATCT
hTRPV1	NM_080704.3	TGGATGGCTTGCCTCCCTTTA
	NM_018727.5	ACTGTAGCTGTCCACAAACAG
	NM_080706.3	
	NM_080705.3	
CRE	B449974.1	TCCATATTGGCAGAAGGAA
		CAGCTACACCAGAGACGGAA

















