Histology and Immunohistochemistry

Mice were individually euthanized, dissected, and their resected intestinal tissue was immediately fixed in 10% neutral-buffered formalin (Millipore Sigma) 40h after treatment with tamoxifen. Tissue specimens were fixed for 16-24 hours, washed in PBS three times, then placed in 70% ethanol and shipped to Histowiz (Brooklyn, NY) for paraffin embedding, sectioning, staining, and high-resolution (40X) scanning according to standard protocols. H&E histologic severity quantitation was performed using a scoring system previously described for genetic IBD models affecting the small intestine (score range 0-15) or colon (score range 0-12) (114). For CC3-stained sections (Cell Signaling Technology 9661), 10 consecutive crypts from each of 5 sections for small intestine and colon (50 crypts per mouse per segment) were counted. Chromogenic RNA *in situ hybridization* (RNA-ISH) was performed using RNAscope[®] Probe Mm-*Lta* (Advanced Cell Diagnostics 317231) and RNAscope 2.5 HD Assay-Brown Kit (Advanced Cell Diagnostics 322370) according to the manufacturer's specifications. Slides were immediately covered with a coverslip using Cytoseal. Slides were then scanned at 40X magnification with a Zeiss Axio ScanZ.1 (Zeiss). DAB-positive areas were identified against a hematoxylin background in all 5 sections of small intestine and colon on each slide using QuPath v0.2.3. At least two and up to three blinded readers assessed histologic disease severity, CC3 quantitation, and *Lta* quantitation and the results were averaged.

Antibodies and reagents

Antibodies directed against A20 (Cell Signaling Technology, 5630), ABIN-1 (Millipore Sigma HPA037893), Caspase 8 (Cell Signaling Technology mouse specific 4927 and 4790), CC8 (Cell Signaling Technology 8592), RIPK1 (Cell Signaling Technology 3493), Caspase 3 (Cell Signaling Technology 9662), CC3 (Cell Signaling Technology 9661), PARP (Cell Signaling Technology 9542), mouse RIPK3 (Cell Signaling Technology 95702), phospho-RIPK3 (T231/S232) (Cell Signaling Technology 91702), phospho-RIPK1 (Ser166) rodent specific (Cell Signaling Technology 31122), and GAPDH (Millipore MAB374), were used for western blot. Anti-CC3 (Cell Signaling Technology 9661) was also used for IHC studies. Mouse IFN- β ELISA (PBL assay bioscience) was used according to the manufacturer's instructions. Recombinant mouse TNF, mouse IL-18, mouse TL1A, mouse LT $\alpha_1\beta_2$, human LT α_3 , mouse TNFR1-Fc, and mouse LTBR-Fc were purchased from R&D systems. Recombinant mouse RANKL was purchased from BioLegend. Recombinant mouse IL-1 β and mouse LIGHT were purchased from Peprotech. Recombinant human TNF was purchased from Enzo Life Sciences. Pam3CSK4, poly(I:C) HMW, and LPS from E. coli O111:B4 were purchased from Invivogen. Necrostatin-1s (BioVision 2263) and Emricasan (MedChem Express HY-10396) were used at 50µM final concentration. S5H3.2.2 hybridoma cell line (ATCC) expressing monoclonal hamster anti-mouse LTα (65) was provided to BioXcell (Lebanon, New Hampshire) for production of purified monoclonal antibody for in vivo use. InVivoMAb Armenian hamster IgG isotype control anti-glutathione S-transferase (BioXcell #BE0260) was used as a negative control.

LTα₃-Fc conditioned media

Codon-optimized gblocks (IDT) encoding mouse $LT\alpha$ (amino acids 1-202; UniProtKB/Swiss-Prot P09225.1) followed by a GSG linker and mouse IgG2a Fc fusion (amino acid 99-330; UniProtKB/Swiss-Prot P01863) were synthesized and then cloned into pENTR (Addgene Plasmid #17398) using InFusion cloning according to manufacturer's instructions (Clontech). The insert sequence was confirmed and then cloned into pLEX 307 (Addgene Plasmid #41392) using the Gateway Cloning LR reaction according to manufacturer's instructions (ThermoFisher Scientific). Lentivirus particles were prepared by co-transfection with the packaging plasmid psPAX2 (Addgene Plasmid #12260) and the envelope plasmid pMD2.G (Addgene Plasmid #12259) into HEK293T (ATCC) cells using Lipofectamine 2000 (ThermoFisher Scientific). The lentivirus-containing medium was collected at 72 h post-transfection. After centrifugation, the lentivirus medium was filtered using a 0.45 µm syringe filter. HEK293T cells were transduced with diluted lentivirus and selected with puromycin 5 ug/ml for several passages. The puromycin was removed and mLTα-Fc conditioned media (mLTα-Fc CM) was collected and filtered over $0.22 \mu m$ filter. The mLT α -Fc CM was guantitated by ELISA with goat polyclonal LT α capture antibody (R&D systems AF749), rat monoclonal LT α detection antibody (R&D systems MAB749) biotinylated with ChromaLink biotin according to manufacturer's instructions (Trilink Biotechnologies), a recombinant mouse lymphotoxin alpha ELISA reference (R&D systems), peroxidase streptavidin (Jackson ImmunoResearch), and 1-Step Ultra TMB-ELISA (ThermoFisher Scientific).

References:

114. Erben U et al. A guide to histomorphological evaluation of intestinal inflammation in mouse models. *International journal of clinical and experimental pathology* 2014; 7(8):4557-4576.



Supplemental Figure 1. Tamoxifen-induced deletion of floxed *A20* and *Abin-1* on a *Vil-cre-ER*^{T2+} background in vivo and in vitro. (A) Immunoblotting analysis of IECs with the indicated genotypes 40h after treatment with 2mg tamoxifen per os (p.o.). (B) Immunoblotting analysis of *A20/Abin-1*^{T-Δ/EC}*Tn*^{F/-} enteroid cultures treated with the indicated concentration of 4-OHT for 24h. For panels (**A**,**B**), lysates were immunoblotted with the antibodies indicated on the right. (**C**) Representative confocal microscopy images of PI-stained enteroids (pseudocolor red) from indicated genotypes of mice treated with vehicle or 4-OHT for 24h followed by 24 h of 2.5 ng/ml TNF as indicated. Scale bars, 100µm. (**D**) Quantitative luminescent cell viability assay of enteroid cultures treated as described in (**C**) (mean ± SEM). For panel (**D**), significance was assessed by two-way ANOVA with Dunnett's multiple comparison test comparing *A20/Abin-1*^{T-Δ/EC} and *A2*



Supplemental Figure 2. Editing of MyD88 in A20/Abin-1^{T-ΔIEC}Tnf-/- mice. Nucleotide chromatogram and sequence of the MyD88 PCR amplicon surrounding the CRISPR-Cas9-targeted site for (A) MyD88 C1 and (B) MyD88 C2 in A20/Abin-1^{T-ΔIEC}Tnf-/- mice.

Supplemental Figure 3



Supplemental Figure 3. IL18 stimulation of A20/Abin-1^{T-ΔIEC}Tnf^{-/-} enteroids and editing of Trif in A20/Abin-1^{T-ΔIEC}Tnf^{-/-} mice. (A) Quantitative luminescent cell viability assay of A20/Abin-1^{T-ΔIEC}Tnf^{-/-} enteroid cultures treated with the indicated stimuli (mean \pm SEM; IL-18 25ng/ml). (B) Agarose gel electrophoresis and (C) nucleotide chromatogram and sequence of the Trif PCR amplicon surrounding the CRISPR-Cas9-targeted site in Trif for A20/Abin-1^{T-ΔIEC}Tnf^{-/-} mice. (D) IFN-β ELISA of mouse splenocytes with the indicated genotype stimulated with 100 µg/ml poly(I:C). For panel (A) significance was assessed using one-way ANOVA with Dunnett's multiple comparison test relative to vehicle alone. For panel (D), significance was assessed using unpaired t-test. Only significant differences are shown; ****=p< 0.0001. Data represent at least two independent experiments.



Supplemental Figure 4. Deletion of *Casp8* in *A20/Abin-1^{T-ΔIEC}Ripk3^{-/-}* mice and poly(I:C) stimulation of *A20/Abin-1^{T-ΔIEC}Ripk3^{-/-}* and *A20/Abin-1^{T-ΔIEC}Ripk3^{-/-}* Casp8^{-/-} enteroids. (A) Nucleotide chromatogram and sequence of the *Casp8* PCR amplicon surrounding the CRISPR-Cas9-targeted site in *Casp8* for *A20/Abin-1^{T-ΔIEC}Ripk3^{-/-}* mice. (B) Immunoblot of intestinal epithelial cells from *A20/Abin-1^{T-ΔIEC}Ripk3^{-/-}* mice with the indicated *Casp8* genotypes. (C) Quantitative luminescent cell viability assay of enteroids with the indicated genotype treated with indicated stimuli, (mean ± SEM). For panel (C) statistical significance was assessed by two-way ANOVA with Bonferroni's multiple comparison test between genotypes for each stimulation condition. Only significant differences are shown; ***=p < 0.001; ****=p< 0.0001. Data represent at least two independent experiments.



Supplemental Figure 5. Anti-LT α alone, in the absence of TNF neutralization or depletion, does not improve survival in A20/Abin-1^{*T*}-AVEC mice. (A) Weight curve and (B) Kaplan-Meier survival curves of the indicated genotypes of tamoxifen-treated mice treated with anti-LT α or isotype control. For panel (A), significance was assessed by two-way ANOVA with Bonferroni's multiple comparison test. For panel (B), significance was assessed by Log-rank Mantel-Cox test. Only significant differences are shown.





Supplemental Figure 6. A20 and ABIN-1 protein levels are relatively decreased in inflamed intestinal mucosa in patients with IBD. (A) qPCR for the indicated mRNA transcripts in colonic mucosal biopsies (median and interquartile ranges) and (B) immunoblot of lysates from a subset of biopsies from inflamed (red) and/or non-inflamed (blue) areas from patients with IBD or without IBD (non-IBD, black). Patient and biopsy number are shown, along with location (R-right colon; L-left colon). Biopsy samples were obtained at one or two locations and processed separately, Each data point represents one patient. RKO cell line mRNA and lysate were used as a reference. (Median and interquartile ranges). (C) Relative A20 (TNFAIP3) protein (left panel), mRNA (middle panel), and protein/mRNA (right panel) quantitation from a subset of patients in (A) (mean ± SD). (D) Correlation between A20 (TNFAIP3) protein and mRNA in mucosal biopsies. (E) Relative ABIN-1 (TNIP1) protein (left panel), mRNA (right panel) quantitation from a subset of patients in (A) (mean ± SD). (D) Correlation between A20 (TNFAIP3) protein and mRNA in mucosal biopsies. (E) Relative ABIN-1 (TNIP1) protein and mRNA (middle panel), and protein/mRNA (right panel) quantitation from a subset of patients in (A) (mean ± SD). (P) Correlation between A8IN-1 (TNIP1) protein and mRNA in mucosal biopsies. For panels (A, C, E), significance was assessed using one-way ANOVA with Tukey's multiple comparison test. For panels (D,F), pearson correlation coefficient and corresponding p-value were calculated. ns=not significant, *=p<0.05, **=p<0.01. Data represent two independent measurements.

Characteristic		Non-IBD (n=19)	IBD (n=33)	p-value
Gender (F/M)		11/8	13/20	0.25
Age, years median (IQR1–IQR3)		53 (36-61)	37 (30-48)	0.005
Disease type n (%)				<0.0001
	UC	0	20 (61%)	
	CD	0	13 (39%)	
	non-IBD	19 (100%)	0	
Racial and Ethnic Category n (%)				0.61
	White	15 (79%)	28 (85%)	
	Hispanic	1 (5%)	0	
	Black	2 (11%)	2 (6%)	
	Asian	1 (5%)	2 (6%)	
	Other/unknown	0	1 (3%)	
Medication				<0.0001
	Steroids n (%)	0	1 (3%)	
	Mesalamine n (%)	0	6 (18%)	
	Immunomodulator n (%)	1 (6%)	13 (39%)	
	Biologic n (%)	0	25 (76%)	
	anti-TNF	U	23 (70%)	
	other biologic		2 (6%)	
Endoscopic inflammation n (%)		0	13 (39%)	0.0018

Supplemental Table 1. Baseline demographic and clinical data for study participants. Categorical variables were analyzed by Fisher's exact or Chi-square tests where appropriate, and non-normally distributed continuous variables were compared using Mann-Whitney test.