ORIGINAL ARTICLE



Deficiency of IL-17A, but not the prototypical Th17 transcription factor RORyt, decreases murine spontaneous intestinal tumorigenesis

Mia Shapiro^{1,2} · Bisweswar Nandi^{1,2} · Christine Pai¹ · Mehmet K. Samur^{2,3} · Dheeraj Pelluru^{1,3} · Mariateresa Fulciniti^{2,3} · Rao H. Prabhala^{1,2,3} · Nikhil C. Munshi^{2,3,4} · Jason S. Gold^{2,5,6}

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Abstract While inflammation has been associated with the development and progression of colorectal cancer, the exact role of the inflammatory Th17 pathway remains unclear. In this study, we aimed to determine the relative importance of IL-17A and the master regulator of the Th17 pathway, the transcription factor RORyt, in the sporadic intestinal neoplasia of APC^{MIN/+} mice and in human colorectal cancer. We show that levels of IL-17A are increased in human colon cancer as compared to adjacent uninvolved colon. Similarly, naïve helper T cells from colorectal cancer patients are more inducible into the Th17 pathway. Furthermore, IL-17A, IL-21, IL-22, and IL-23 are all demonstrated to be directly mitogenic to human colorectal cancer cell lines. Nevertheless, deficiency of IL-17A but not RORyt is associated with decreased spontaneous intestinal tumorigenesis in the APC^{MIN/+} mouse model, despite the fact that helper T cells from RORyt-deficient APCMIN/+

Mia Shapiro and Bisweswar Nandi: Co-first authors.

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☑ Jason S. Gold jgold4@partners.org

- ¹ Research Service, VA Boston Healthcare System, West Roxbury, MA, USA
- ² Harvard Medical School, Boston, MA, USA
- ³ Dana-Farber Cancer Center, Boston, MA, USA
- ⁴ Medicine Service, VA Boston Healthcare System, West Roxbury, MA, USA
- ⁵ Surgery Service, VA Boston Healthcare System (112), 1400 VFW Parkway, West Roxbury, MA 02132, USA
- ⁶ Brigham and Women's Hospital, Boston, MA, USA

mice do not secrete IL-17A when subjected to Th17-polarizing conditions and that II17a expression is decreased in the intestine of RORyt-deficient APC^{MIN/+} mice. Differential expression of Th17-associated cytokines between IL-17A-deficient and RORyt-deficient APC^{MIN/+} mice may explain the difference in adenoma development.

Keywords ROR γ t · IL-17A · Th17 · Carcinogenesis · Adenoma · Colorectal neoplasms

Abbreviations

ANCOVA	Analysis of covariance for independent
	samples
APC	Adenomatous polyposis coli
Apc	Adenomatous polyposis coli gene
ATCC	American Type Culture Collection
FAP	Familial adenomatous polyposis
HET	Heterozygote
IACUC	Institutional Animal Care and Use Committee
IBD	Inflammatory bowel disease
ILC	Innate lymphoid cell
ILC3	Group 3 innate lymphoid cell
IRB	Institutional review board
KO	Knockout
MIN	Multiple intestinal neoplasia
NKT cell	Natural killer T cell
qRT-PCR	Quantitative real-time polymerase chain
	reaction
siRNA	Small interfering ribonucleic acid
SPF	Specific pathogen-free
Treg	Regulatory T cell
VABHS	VA Boston Healthcare System
WT	Wild type

It is widely accepted that inflammation contributes to the development of cancer [1]. Pathologists have long noted the presence of inflammatory cells in tumors [2]. A relationship between inflammation and intestinal neoplasia is supported by the facts that inflammatory bowel disease predisposes patients to intestinal carcinomas and that the anti-inflammatory drugs aspirin [3–5], celecoxib [6], and rofecoxib [7], all have proven efficacy in preventing human colorectal adenoma development.

The APC^{MIN/+} mouse strain is the most widely studied murine model of intestinal neoplasia. This model mimics the human condition familial adenoma polyposis (FAP) characterized by the sporadic development of numerous intestinal adenomatous polyps. In the mouse, the small intestine is most affected, whereas the large intestine is most affected in humans. In APC^{MIN/+} mice, as in humans, a germline mutation in the adenomatous polyposis coli (*Apc*) gene is responsible for the phenotype.

Despite the fact that adenoma development is controlled by a mutation in a tumor suppressor gene, inflammation is critically important for carcinogenesis in murine models with germline *Apc* mutations. Adenoma formation is increased after adoptive transfer of pro-inflammatory CD4⁺CD45RB^{hi} lymphocytes [8] and decreased after the adoptive transfer of anti-inflammatory regulatory T cells (Tregs) [9]. Similarly, adenoma formation is decreased in the absence of the adapter molecule MyD88, which is essential for mediating inflammation from bacterial and viral products [10], as well as the inflammatory cytokine IL-6 [11] and the inflammatory chemokine receptor CCR6 [12].

CD4⁺ T cells polarized to the Th17 pathway as characterized by the secretion of interleukin 17A (IL-17A) as well as IL-17F, IL-21, and IL-22 are essential for the development of several chronic inflammatory diseases such as multiple sclerosis [13], rheumatoid arthritis [14], psoriasis [15], asthma [16], and inflammatory bowel disease (IBD) [17]. The transcription factor ROR γ t has been identified as the master regulator of polarization of helper T cells toward the Th17 pathway [18].

Several lines of evidence implicate the Th17 pathway in intestinal carcinogenesis [19–28]. The role of the Th17 in cancer, however, is not clear cut as conflicting evidence supports an anti-tumor role for IL-17A [29, 30]. In this study, we aimed to determine the relative importance of IL-17A and the transcription factor ROR γ t in sporadic intestinal carcinogenesis.

Methods

Human samples

Tissue samples were acquired immediately after resection from patients undergoing surgery for colorectal cancer at VA Boston Healthcare System (VABHS). Blood was also obtained prior to surgery. Written consent was obtained for collection of all specimens. Oral antibiotic bowel preparation and intravenous antibiotic prophylaxis were routinely used. The protocol was approved by the VABHS Institutional Review Board (IRB) (VABHS IRB protocol #2199).

Animals

APC^{MIN/+} mice (C57BL/6J-Apc^{MIN}/J) and RORyt^{GFP/GFP} (RORytKO) mice with the green fluorescent protein (GFP) transcript inserted in place of the first exon of the RORyt gene in both alleles (B6.129P2(Cg)-Rorctm2Litt/J), and wildtype C57BL/6J mice were purchased from the Jackson Laboratory (Bar Harbor, ME, USA). IL-17A^{-/-} (IL-17AKO) mice were generously donated from Dale T. Umetsu (Boston Children's Hospital, Boston, MA, USA) with the permission of Yoichiro Iwakura (Tokyo, Japan) [31]. APC^{MIN/+} and RORyt^{GFP/GFP} mice were intercrossed to generate ROR $\gamma t^{GFP/GFP}$ -APC^{MIN/+} (ROR γKO -APC^{MIN/+}) and $ROR\nu t^{GFP/+} - APC^{MIN/+}$ (ROR $\nu HET - APC^{MIN/+}$) mice. $APC^{MIN/+}$ mice were also crossed with IL17A^{-/-} mice to generate IL17A^{-/-}-APC^{MIN/+} (IL17KO-APC^{MIN/+}) mice. Mouse genotypes were confirmed by performing PCR on DNA from tail snips using primers and conditions provided by the Jackson Laboratory and Dr. Iwakura. Experimental mice were not routinely littermates or co-housed. The mice were housed and maintained in the specific pathogen-free (SPF) animal research facility of VABHS. The experimental protocols were approved by VABHS Institutional Animal Care and Use Committee (IACUC) (VABHS IACUC protocols #247, 289, and 291).

Cell lines

The murine colon cancer cell line MC38 was provided by Michael T. Lotze of the University of Pittsburgh [32]. Cell lines were grown in RPMI 1640 (Life Technologies, Grand Island, NY, USA) with 10 % fetal bovine serum (FBS) (Life Technologies) and antibiotic–antimycotic (Life Technologies). The human colorectal adenocarcinoma cell lines HCT 116, HT-29, and SW480 were purchased from American Type Culture Collection (ATCC, Manassas, VA, USA).

Proliferation assay

Cells were plated at 10,000 cells per well in RPMI 1640 with 10 % FBS in 96-well plates either in the presence or absence of IL-17A, IL-21, IL-22, or IL-23 (R&D Systems, Minneapolis, MN, USA) at 50 ng/mL for 48 h. Cells were then pulsed with ³H-thymidine (0.5 micro Ci/well) for 6 h and harvested, and radioactivity was counted using the LKB Betaplate scintillation counter.

Quantification of polyps

Following euthanasia, the small intestine was dissected from the pylorus to the ileocecal valve. The intestines were cannulated and flushed with phosphate-buffered saline (PBS) to remove fecal material. The cleaned intestine was divided into three equal length segments: duodenum (first segment), jejunum (second), and ileum (third, most distal). Each segment was cut longitudinally and examined under a dissecting microscope at $4 \times$ magnification. Polyps were counted and measured at their greatest diameter [12]. Tumor load was determined by multiplying the number of polyps by their measured diameters [25].

Tumor challenge

MC38 cells were cultured until they reached 80 % confluence. 5×10^5 cells in 100 µL of sterile PBS were injected subcutaneously into the right hindquarter of 6-week-old sex-matched C57BL/6J (WT) and RORγKO mice. Tumors were measured by their greatest dimension 3 times a week until they reached an endpoint of 2 cm in largest diameter or ulceration.

Human IL-17A concentration in colon cancer specimens

Snap-frozen sections of human tissue were stored at -80 °C until use. Upon thawing, tissues were homogenized. Homogenates were made in a volume of 200 µL of PBS using an Eppendorf micropestle in a 1 mL Eppendorf tube (Eppendorf, Hauppauge, NY, USA). The homogenized samples were then centrifuged at 13,000 rpm for 10 min to remove debris, and the supernatant was collected. Protein concentration was measured using the Quick Start Bradford protein assay (Bio-Rad Life Sciences, Hercules, CA, USA). IL-17A levels were assessed by multiplexed ELISA using a Q-plex assay chemiluminescent kit (Quansys Bioscience, UT). All buffers/solutions were provided as part of the Quansys 16-plex kit. Briefly, supernatants were diluted as per protocol in sample diluent buffer. Recombinant cytokine was serially diluted to create a standard curve. Fifty microliters of each dilution and 50 µL of diluted test sample supernatants were added to a 96-well Q-Plex Array plate. The plate was incubated for 1 h at room temperature on an orbital shaker (500 rpm). Following incubation, plates were washed 3 times with wash buffer and incubated with 50 mL of detection buffer for 1 h on a shaker (500 rpm). The wells were again washed 3 times with buffer, and 50 μ L of 1× Streptavidin–HRP solution was added to each well for 15 min at room temperature on a shaker (500 rpm). Next, the wells were washed 3 times with wash buffer and incubated for 15 min with 50 μ L per well of mixed substrates A and B. The plate was then imaged using a Q-View imager (Quansys Bioscience). Results were analyzed using Q-view software (Quansys Bioscience).

Expression data from The Cancer Genome Atlas

Data from The Cancer Genome Atlas (TCGA) for colon adenocarcinoma (COAD) [33] were downloaded from the Broad Institute's Firehose pipeline using RTCGAToolbox [34]. Agilent microarray-based data were used to compare expression of *IL17A* in colon cancers and normal colon. The empirical Bayes method for assessing differential expression was used for statistical analysis. Microarrayand RNAseq-based gene-level expression data were used to test the correlation between *IL17A* and *RORC* expression using Pearson correlation coefficients. R and the Bioconductor limma package were used for statistical analysis.

RNA extraction and qRT-PCR

Murine small intestinal tissue was dissected from 15-weekold mice and cleaned as was done for quantification of polyps above. Once clean, the third portion of the intestine (ileum) was snap frozen and stored at -80 °C until use. Upon thawing, cell lysates were prepared from ~100 mg of intestinal tissue by addition of 900 µl of QIAzol Lysis Reagent (Qiagen) in an Eppendorf tube followed by tissue homogenization using the Tissue Lyser II (Qiagen). RNA extraction was performed using the RNeasy Plus Universal Kit (Qiagen, Valencia, CA, USA). For cDNA synthesis, 5 µg of total RNA was reverse-transcribed using the Super-Script III First-Strand Synthesis System (Life Technologies) where oligo-dT was used as the first-strand primer. qPCR for 1123, 1122, 1121, 1117a, 116, Ifng, Tnfa, 1110, Ki67, Bad, Bax, Bbc3 (PUMA), Bcl2, and Bcl2l1 (Bcl-xL) was performed from cDNA. The primers are listed in Supplementary Table 1. Fold change was calculated using the $2-\Delta\Delta$ Ct method with normalization to GAPDH.

Mouse Th17 differentiation

Spleens were harvested from 6- to 8-week-old mice, mechanically dissociated using a $100 \,\mu$ M filter, and washed

with RPMI. DNase I (Roche Life Science, Indianapolis, IN, USA) was added (1:200 dilution 50 µg/mL). Red blood cells were lysed using BD Pharm Lyse (BD Biosciences, San Jose, CA, USA). Naïve CD4⁺ cells were extracted using the CD4⁺CD62L⁺ T Cell Isolation Kit II (Miltenyi Biotec, Cologne, Germany). Two rounds of negative selection were performed followed by 1 round of positive selection per the manufacturer's protocol. A 12-well polystyrene plate was coated with anti-CD3 (clone 145-2C11, eBioscience, San Diego, CA, USA) (4 µg/mL) and anti-CD28 (2 µg/mL) (clone 37.51.1, Invitrogen, Grand Island, NY, USA) diluted in PBS and incubated for 3 h at 37 °C. PBS was removed, and 1×10^6 cells were added to each well along with anti-IL-4 antibody (1:250, clone 11B11, eBioscience), IL-6 (50 ng/mL) (R&D Systems), TGF-B1 (5 ng/ mL) (R&D Systems), IL-23 (10 ng/mL) (R&D Systems), anti-IFN-y antibody (5 µg/mL, clone XMG1.2, eBioscience), and IL-1B (20 ng/mL) (R&D Systems). After 6 days, cells were stimulated with phorbol 12-myristate13-acetate (PMA) (50 ng/mL) and ionomycin (750 ng/mL) for a total of 5 h. Brefeldin A (25 µg/mL) was added (Sigma-Aldrich, St. Louis, MO, USA) after the first hour. Following stimulation, rat anti-mouse CD16/32 (clone 2.4G2, BD Biosciences) was added at a 1:100 dilution for 10 min. Anti-mouse CD3 APC (clone 145-2C11, BD Biosciences) and anti-mouse CD4 AlexaFluor700 (clone GK1.5, eBioscience) were added, and cells were incubated for 20 min. Cells were fixed with 2 % paraformaldehyde for 20 min and then permeabilized with 0.5 % saponin (Sigma-Aldrich) and stained with anti-mouse IL-17A-PE (clone TC11-18H10, BD Biosciences) and anti-mouse IFNy-PE-Cy7 (clone XMG1.2, BD Biosciences).

Human Th17 differentiation

PBMCs were isolated from whole blood using a Ficoll-Paque gradient (GE Healthcare, Piscataway, NJ, USA). PBMCs were polarized with a cocktail consisting of IL-1β (10 ng/mL), IL-6 (20 ng/mL), and IL-23 (10 ng/mL) in addition to anti-IFN-y (10 µg/mL), anti-IL-4 (10 µg/mL), TGF- β (1 ng/mL) (all from R&D Systems) and anti-CD3 and anti-CD28 antibodies (BD Biosciences) for 6 days in RPMI 1640 supplemented with 10 % FBS. Cells were further expanded with IL-2 (20 U/mL; R&D Systems) for an additional 6 days prior to restimulation with PMA and ionomycin (Sigma) for 6 h in the presence of GolgiStop (eBioscience). The cells were stained for CD4 (BD Biosciences), and after washing, cells were fixed with Cytofix/Cytoperm buffer (BD Biosciences). Intracellular staining for IFN-y and IL-17A was performed using conjugated anti-IFN-y (BD Biosciences) and anti-IL-17A (eBioscience) antibodies and analyzed using flow cytometer (BD Biosciences) as described previously [35].

Results

As several reports suggest that the Th17 pathway may promote intestinal carcinogenesis, we first sought to assess the expression of IL-17A in human colorectal cancer. IL-17A protein levels were measured in matched samples of human colorectal cancer tumors and adjacent uninvolved colon. The mean level of IL-17A expression was found to be significantly greater in the tumors. Of the 19 matched samples analyzed, there was more IL-17A detected in the tumor in 13 (68 %) with less noted in only 3 (16 %). In 10 matched samples (53 %), the tumor had at least double the level of IL-17A. The magnitude of increase in IL-17A level ranged up to over 47,000 % (Fig. 1a). An increase in IL17A gene expression was confirmed using data from The Cancer Genome Atlas (TCGA) (Fig. 1b). A more modest increase in IL17A expression in colon cancers as compared to nondiseased colon was seen (fold change 1.14, p = 0.01). In contrast, expression of RORC, the gene encoding RORyt, the prototypical transcription factor that drives Th17 differentiation, was decreased in tumors as compared to nondiseased colon (fold change 1.7, p < 0.001). (Supplementary Fig. 1). Interestingly, a correlation between IL17A and RORC expression was not found (Supplementary Fig. 2).

We next assessed whether helper T cells in colorectal cancer patients were more prone to be induced into the Th17 pathway. PBMCs from colorectal cancer patients and healthy controls were purified and subjected to Th17-polarizing conditions and then non-specifically stimulated and analyzed by flow cytometry with intracellular staining. The mean number of CD4⁺ cells expressing IL-17A after polarization was approximately 7.5-fold higher in the cancer patients as compared to healthy controls. In contrast, the level of IFN- γ -expressing CD4⁺ cells was approximately 2.5-fold lower in the cancer patients. We did not note a significant variation with stage in the percent of CD4⁺ cells expressing either IL-17A or IFN- γ (Fig. 1c).

We then sought to explore a causal relationship between the Th17 pathway and tumor growth. We first examined whether cytokines associated with the Th17 pathway were directly mitogenic to colorectal cancer cells. We subjected 3 different human colorectal cancer cell lines to exposure to IL-17A, IL-21, IL-22, and IL-23 and measured proliferation. Two of the cell lines (Hct 116 and HT-29) appeared responsive to the Th17 cytokines as exposure to each of the 4 cytokines significantly increased proliferation from 203 to 879 %. Proliferation of the third cell line (SW480) was not affected by IL-17A, IL-21, IL-22, or IL-23 (Fig. 2).

To further assess the role of the Th17 pathway in intestinal tumorigenesis, we used a well-validated model of spontaneous intestinal tumorigenesis that is driven by loss of a tumor suppressor gene. APC^{MIN/+} or multiple intestinal



Fig. 1 IL-17A levels are overexpressed in human colon cancers, and helper T cells from human colorectal cancer patients are more inducible into the Th17 pathway. **a** IL-17A expression in samples of human colon cancer and matched adjacent normal tissue (n = 19) was assayed by ELISA and normalized to total protein measured by the Bradford assay. The values for cancer and normal tissue were compared for each patient, and the Wilcoxon signed-rank test was used to assess significance. **b** *IL17A* gene expression in 153 colon cancers and 19 normal colon samples was determined from microarray data. The *boxes* represent the interquartile range (IQR); the *lines* represents the median; the *error bars* represent the lowest and highest data still within 1.5 IQR of the upper and lower quartiles, respectively; and the *circles* represent the data points outside of 1.5 IQR of the upper and

neoplasia (MIN) mice are heterozygous for a germline mutation in the adenomatous polyposis coli (*Apc*) gene, resulting in a non-functional gene product. Mice spontaneously develop multiple intestinal adenomas mimicking the human condition FAP. We bred these animals to mice deficient in ROR γ t (ROR γ tKO mice), to create APC^{MIN/+} mice that were either heterozygous or homozygous for loss of ROR γ t (ROR γ tHET–APC^{MIN/+} mice and ROR γ tKO–APC-^{MIN/+} mice, respectively). We first confirmed that helper T cells from ROR γ tKO–APC^{MIN/+} do not secrete IL-17A when subjected to Th17-polarizing conditions. While the number of IFN- γ -expressing cells after activation with a

lower quartiles. The empirical Bayes method for assessing differential expression was used for statistical analysis. **c** PBMCs were isolated from normal donors and colon cancer patients and then activated with a Th17-polarizing cocktail for 12 days. Intracellular IL-17A and IFN- γ were measured by flow cytometry following PMA/ionomycin restimulation. The *right panels* are representative dot plot analyses from one healthy patient and one patient with stage IV colon cancer showing the percentage of cells that are positive for intracellular IL-17A or IFN- γ within the gated CD4⁺ population. The cells were single-stained for each cytokine. Composite results are presented in the *bar graph* on the *left* as mean values with standard error of the mean (*p < 0.05, compared to healthy controls). The Student's *t* test was used for statistical comparison

Th17-polarizing cocktail was similar between the different genotypes, there were minimal IL-17A-expressing cells from the IL-17A knockout (IL-17AKO), ROR γ tKO, and ROR γ tKO–APC^{MIN/+} mice while IL-17A production was maintained in the WT and APC^{MIN/+} mice (Fig. 3).

We then compared polyp development in APC^{MIN/+}, IL-17AKO–APC^{MIN/+}, ROR γ tHET–APC^{MIN/+}, and ROR γ tKO–APC^{MIN/+} mice. At 15 weeks of age, the total polyp number and total polyp load in the small intestine were significantly decreased for IL-17AKO–APC^{MIN/+} mice as compared to APC^{MIN/+} mice (Fig. 4a). The polyp number and load were also found to be significantly decreased in each segment of the small bowel (duodenum, jejunum, and ileum; Fig. 4b). In contrast, polyp number and load were preserved for both the RORytHET-APC^{MIN/+} and $RORvtKO-APC^{MIN/+}$ mice in the entire small intestine and in each segment (Fig. 4). Polyps were also quantified for RORytHET-APC^{MIN/+} and RORytKO-APC^{MIN/+} mice at 22 weeks, and no difference was found in the polyp number overall or in individual segments. A very small but significant decrease in polyp load was seen in RORytKO-APC-MIN/+ mice that were restricted to the ileum (Supplementary Fig. 3). We assessed relative expression of genes associated with proliferation and apoptosis in the adenoma-bearing intestines of APCMIN/+, RORytKO-APCMIN/+, and IL-17AKO–APC^{MIN/+} mice. The proliferation gene Ki67 was modestly downregulated in the intestines of RORytKO-APC^{MIN} as compared to APC^{MIN} mice, whereas expression



Fig. 2 Human colorectal cancer cell lines proliferate in response to Th17 cytokines. Proliferation of cell lines cultured with or without 50 ng/mL of IL-17A, IL-21, IL-22, or IL-23 for 48 h was assessed by thymidine incorporation. Thymidine incorporation of cells cultured with cytokines is shown as a percentage of that for the same cell line cultured without cytokine. *Error bars* show the standard error of the mean (*p < 0.05, compared to thymidine incorporation in the absence of cytokine). Data are representative of 3 independent experiments. The Student's *t* test was used for statistical comparisons

did not appear to be altered in the intestines of IL-17AKO– APC^{MIN/+} mice (Supplementary Fig. 4a). In contrast, adenoma-bearing intestine from IL-17AKO–APC^{MIN/+} mice had decreased expression of the anti-apoptotic genes *Bad*, *Bax*, and *Bbc3* (PUMA) with increased expression of the pro-apoptotic genes *Bcl2* and *Bcl2l1* (Bcl-xL) in comparison with APC^{MIN} mice, whereas the expression of these genes was only altered modestly if at all in RORγtKO– APC^{MIN/+} mice (Supplementary Fig. 4b).

We also explored whether deficiency of ROR γ t affected tumor growth after subcutaneous injection of the colorectal cancer cell line MC38, which is syngeneic for C57/ Bl6 mice. In this model, we did not detect any difference in tumor growth or in survival between WT and ROR γ tKO mice after tumor challenge (Fig. 5).

Lastly, in order to attempt to explain the difference in intestinal tumorigenesis observed between RORytKO-APC^{MIN/+} and IL-17AKO-APC^{MIN/+} mice, we assessed the relative expression of cytokines in the adenoma-bearing intestines of APCMIN/+, RORytKO-APCMIN/+, and IL-17AKO-APC^{MIN/+} mice. The inflammatory cytokines 1123, 1117a, and 116 were decreased in the intestines of $ROR\gamma tKO-APC^{MIN/+}$ and $IL-17AKO-APC^{MIN/+}$ mice as compared to those of APC^{MIN/+} mice. The magnitude of the decrease in Il17a expression was greater in the IL-17AKO-APC^{MIN/+} mice. There appeared to be alterations in the expression of Il22 and Il21 between the intestines of RORytKO-APC^{MIN/+} and IL-17AKO-APC^{MIN/+} mice as compared to those of APCMIN/+ mice. While these cytokines were both decreased in RORytKO-APC^{MIN/+} mice intestines, Il22 was only marginally decreased and *Il21* was increased in the IL-17AKO-APC^{MIN/+} intestines (Fig. 6). As IL-21 can be associated with increased tumor immunity through granzyme B- and perforin-dependent mechanisms [36], we also assessed expression of *Gzmb1* and Prf1. Consistent with the hypothesis that the increased *Il21* expression in the intestine of IL-17AKO-APC^{MIN/+}



Fig. 3 Helper T cells from ROR γ tKO–APC^{MIN/+} do not produce IL-17A when subjected to Th17-polarizing conditions. Naïve CD4⁺ T cells were purified from the spleens of 6- to 8-week-old mice from WT, IL-17AKO, ROR γ tKO, APC^{MIN/+}, and ROR γ tKO–APC^{MIN/+} mice using magnetic column purification. Cells were cultured in anti-CD3/anti-CD28 coated wells along with anti-IL-4, IL-6, TGF- β , IL-23, anti-IFN- γ , and IL-1 β for 6 days and then stimulated with

PMA and ionomycin followed by Golgi blockade with brefeldin A. Cells were then analyzed by flow cytometry after intracellular staining for IL-17A and IFN- γ . **a** Representative flow cytometry data. **b** The percent of cells expressing IL-17A were compared. p = 0.05 by the Student's *t* test. Data are derived from 5 independent experiments (n = 5 of each APC^{MIN/+} and ROR γ tKO–APC^{MIN/+} mice)



Fig. 4 Lack of IL-17A but not ROR γ t is associated with decreased development of intestinal adenomas. Polyp number and polyp load of 15-week-old mice are shown for **a** the entire small intestine and **b** the duodenum, jejunum and ileum individually (n = 7-12 mice per

group) (*p < 0.05 compared to APC^{MIN/+} mice, $^{\dagger}p < 0.05$ compared to ROR γ tHET–APC^{MIN/+} mice, $^{\ddagger}p < 0.05$ compared to ROR γ tKO–APC^{MIN/+} mice). *Error bars* show the standard error of the mean. The Student's *t* test was used for statistical comparisons

Fig. 5 Lack of ROR γ t does not affect tumor growth or survival after subcutaneous implantation of a syngeneic colon cancer. **a** Mean maximal tumor diameter and **b** survival versus time are shown after injection of 5 × 10⁵ MC38 cells subcutaneously into the flank of WT and ROR γ tKO mice (n = 5-6 per group). Mice were euthanized if they

reached endpoints of a maximal tumor diameter of 2 cm or ulceration. Data are representative of 2 independent experiments. One-way analysis of covariance for independent samples (ANCOVA) and the log-rank test were used to assess significance of tumor growth and survival, respectively

Fig. 6 Relative cytokine expression in adenoma-bearing intestine from APC^{MIN/+}, ROR γ tKO–APC^{MIN/+}, and IL-17AKO–APC^{MIN/+} mice. Ileum of 15-week-old mice was lysed and RNA extracted. cDNA was then synthesized. qRT-PCR was performed. The relative quantification of gene expression was calculated. The fold difference in gene expression for ROR γ tKO–APC^{MIN/+} and IL-17AKO–APC-^{MIN/+} mice relative to APC^{MIN/+} mice is shown on a log₂ scale. *Bars above* the *horizontal axis* represent increased expression relative to APC^{MIN/+} mice, while those *below* the *horizontal axis* represent decreased expression relative to APC^{MIN/+} mice

mice may contribute to the decreased development of intestinal adenomas, we also noted that the expression of *Gzmb1* and *Prf1* appeared to mirror the expression of *Il21* in the intestines of ROR γ tKO–APC^{MIN/+} and IL-17AKO–APC-^{MIN/+} mice (Supplementary Fig. 5). We did not see marked differences in the expression of *Ifng*, *Tnfa*, and *Il10* in the intestine of ROR γ tKO–APC^{MIN/+} versus IL-17AKO–APC-^{MIN/+} mice (Fig. 6).

Discussion

The aim of this paper was to determine the relative importance of IL-17A and RORyt in sporadic intestinal carcinogenesis. We have demonstrated that IL-17A is overexpressed in human colorectal cancer and that peripheral helper T cells from colorectal cancer patients are more inducible into the Th17 pathway. We have also shown that IL-17A, IL-21, IL-22, and IL-23, cytokines associated with the Th17 pathway, are directly mitogenic to human colorectal cancer cell lines. Interestingly, however, deficiency of IL-17A but not RORyt, the master regulator of the Th17 pathway, is associated with decreased murine sporadic intestinal carcinogenesis. Similarly, deficiency of RORyt does not impair tumor growth in a syngeneic mouse adoptive transplant model. Lastly, we explored the cytokine milieu of the adenoma-bearing intestines to better understand the different phenotype between the IL-17A- and RORyt-deficient APC^{MIN/+} mouse models. The inflammatory cytokines *Il17a* and *Il23* as well as *Tnfa* and *Il6* were all suppressed in both models. Unexpectedly, though, expression of *Il21* was increased in the intestine of IL-17A-deficient APC^{MIN/+} mice, while it was decreased in ROR γ t-deficient APC^{MIN/+} mice.

Our finding of the immune system being skewed toward the Th17 pathway in colorectal cancer is supported by other studies. For example, our finding that IL-17A is upregulated at the protein level in human colorectal carcinomas is consistent with the finding that IL17A [19, 20, 25] and IL23 [25] are upregulated at the mRNA level. While an increased infiltration of T cells expressing IL-17A infiltrating colorectal tumors as detected by immunohistochemistry has been previously reported [20], to our knowledge it has not been previously demonstrated that peripheral blood helper T cells from colorectal cancer patients are more apt to be induced to the Th17 pathway. Interestingly, however, we noted that RORC, the gene encoding RORyt, did not appear to be upregulated in human colon cancer and that tumor expression of RORC did not correlate with that of *IL17A*, possibly suggesting that $ROR\gamma t$ may not be as significant in colon cancer and may not account for the increased IL-17A.

While a link of inflammation with the development and progression of colorectal cancer has been established, the mechanisms behind this relationship are poorly understood. Attention has been focused on the importance of the Th17 pathway. In mouse models of sporadic intestinal carcinogenesis, the Th17-associated cytokines IL-17A [22, 25], IL-17F [23], IL-22 [27], and IL-23 [25, 28] have all been shown to promote tumor development even in the absence of other frank inflammatory pathologies. It has been suggested that effect of IL-23 in promoting intestinal neoplasia may be at least partially mediated through IL-17A [25, 28]. We are not aware of other data demonstrating a direct mitogenic effect of the Th17-associated cytokines IL-17A, IL-21, IL-22, and IL-23 on neoplastic epithelial cells in colorectal cancers. This has been shown, however, for related cytokines such as IL-6 and TNF- α [37].

In this study, we confirm that deficiency of IL-17A in APC^{MIN/+} mice results in decreased spontaneous intestinal tumorigenesis [22]. Surprisingly, however, we found that deficiency of ROR γ t, the master regulator of the Th17 pathway, does not reduce adenoma development in the same model. We also did not find reduced growth of transplanted MC38 syngeneic mouse colon cancer tumors. Of note, growth of MC38 in the same adoptive transplant model was not impacted by host deficiency of IL-17A in one study [38], despite the fact that another study using this model showed that abrogation of IL-17A with an adenovirus siRNA vector decreased growth of MC38 [39]. Perhaps deficiency of host IL-17A without ablating any

contribution of IL-17A from tumor cells is not sufficient to reduce tumor growth.

Our finding of a lack of effect of RORyt deficiency on sporadic intestinal polyposis appears to conflict with another study that reported decreased adenomas in the setting of RORyt deficiency in a similar mouse model of sporadic intestinal tumorigenesis induced by an Apc mutation [26]. The model used in the conflicting study, APC^{Δ 468} mice, has 1 allele of the Apc gene with exons 11 and 12 deleted, resulting in a prematurely terminated 468 amino acid protein [40]. In contrast, APC^{MIN/+} mice have 1 Apc allele with a T to A transversion at nucleotide 2549, resulting in a premature stop codon at codon 850 [41]. While it is not clear how the shorter Apc truncation in the other study could result in such a different result, there is a previous instance of a dramatic difference between these 2 models relating to the influence of the immune system on tumorigenesis. A study using APC^{$\Delta 468$} mice reported that mast cells promote adenoma development [42], whereas a separate study using APCMIN/+ mice demonstrated that mast cells exert a protective effect against polyposis [43] despite the fact that both studies used mast cell-deficient Kit^{W-sh/} Wsh (SASH) mice on backgrounds of the respective Apc mutant animals. Further study will be needed to determine which of these mouse systems better models human colorectal carcinogenesis.

It is not obvious why APC^{MIN/+} mice deficient in IL-17A have dramatically decreased intestinal tumorigenesis while those deficient in RORyt do not. Despite the fact that RORyt is the most important transcriptional regulator of IL-17A expression in helper T cells [44] and that helper T cells from RORyt-deficient mice do not polarize into Th17 cells [18], it is conceivable that deficiency of RORyt might not abrogate Th17 polarization in APCMIN/+ mice. Nevertheless, we have shown that helper T cells from RORyt-deficient APC^{MIN/+} mice, as expected, do not secrete IL-17A after exposure to Th17-polarizing conditions. It is also possible that despite the absence of $ROR\gamma t$, IL-17A might still be expressed in the adenoma-bearing intestine of APC^{MIN/+} mice as IL-17A can be secreted by cells other than the helper T cells that were assayed in the polarization assays and as RORyt-independent secretion of IL-17A has been described [45]. While *Il17a* expression is dramatically reduced in the adenoma-bearing intestine of RORyt-deficient APC^{MIN/+} mice, it may not be decreased to the same extent as IL-17A-deficient APCMIN/+ mice. It is possible that a threshold reduction of IL-17A is required for a reduction in adenoma growth. The concept of RORytindependent secretion of IL-17A in colorectal cancer may be supported by the fact that RORC is actually decreased in human colon tumors despite the increased expression of IL-17A and that expression of RORC and IL17A in tumors is not correlated.

Another theoretic possibility is that an effect of RORyt deficiency other than loss of IL-17A expression is responsible for the different phenotypes. RORyt is involved in several immune functions beyond the development of classical Th17 cells. For example, RORyt-deficient mice fail to develop lymphoid organs such as lymph nodes and Peyer's patches [46]. RORyt is also critical for the development and function of innate lymphoid cells (ILCs) [47, 48]. In addition, RORyt is involved in the development of other immune cell subsets including subsets of $\gamma\delta$ T cells [49] and NKT cells [50]. There is considerable plasticity between Tregs and Th17 cells, and in fact, RORyt may be involved in the development of Tregs, including Tregs that subsequently turn off RORyt and do not generate IL-17A [51]. Despite the multifaceted role of ROR γ t in the immune system, loss of RORyt eliminates other IL-17A-mediated pathology in the gut [52].

An intriguing possibility is that another alteration in cytokine production may counterbalance the loss of IL-17A. In the panel of cytokines that we assessed, we did not find differences in expression of the prototypical Th1 cytokine Ifng or the regulatory cytokine Il10 in the adenoma-bearing intestine of IL-17A-deficient or RORytdeficient APC^{MIN/+} mice. We also did not see differences in the expression of the inflammatory cytokines Il23, Tnfa, or Il6. We did, however, detect upregulation of Il21 in the IL-17A-deficient animals as compared to downregulation of *Il21* in the RORyt-deficient animals. As IL-21 can elicit anti-tumor immune responses [36], it is conceivable that the decreased *Il21* in RORyt-deficient APC^{MIN/+} mice may contribute to the difference in intestinal tumorigenesis. The differences in the levels of *Prf1* (perforin) and *Gzmb1* (granzyme B) in the 2 models also support this hypothesis. Further experimentation is needed, however, to definitively determine whether the alterations in *Il21* expression explain the difference in adenoma growth and to delineate the mechanism of this effect.

Interestingly, the cytokine expression in the adenoma environment of the different mouse models corresponds to the expression of the proliferation gene Ki67 as would be expected based on the effect of the cytokines on proliferation seen in vitro. APC^{MIN/+} mice deficient in RORyt have modestly decreased intestinal expression of Ki67 consistent with the decreased expression of the cytokines Il17a, 1121, 1122, and 1123, all of which were shown to promote proliferation of colorectal cancer cell lines. In contrast, IL-17A-deficient APC^{MIN/+} mice, which have decreased intestinal expression of *Il17a* and *Il23* without a change in *Il22* expression and with increased Il21, did not appear to have a change in Ki67 expression compared to APC^{MIN/+} mice. It seems that changes in apoptosis may be more important than changes in proliferation in regulating adenoma growth in these mouse models. Decreased expression of the anti-apoptotic genes *Bad*, *Bax*, and *Bbc3* (PUMA) and increased expression of the pro-apoptotic genes *Bcl2* and *Bcl2l1* (Bcl-xL) mirrored the decreased polyp number for IL-17A-deficient APC^{MIN/+} mice, whereas the expression of these genes was only altered modestly if at all in ROR γ tKO–APC^{MIN/+} mice, consistent to what was observed for adenoma growth. These alterations in apoptotic gene expression are also similar to alterations in *Gzmb1* and *Prf1*, which code for proteins secreted by lymphocytes and natural killer cells to induce the apoptosis-mediated cell death of target cells.

In summary, our results suggest that loss of IL-17A but not ROR γ t diminishes adenoma development in a mouse model of sporadic intestinal carcinogenesis. The lack of measurable change in adenoma development despite the loss of ROR γ t occurs despite the fact that production of IL-17A is impaired in this model. These data imply that strategies directed against the Th17 pathway in colorectal cancer should focus on inhibition of IL-17A itself rather than on ROR γ t.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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