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## Inhibition of a novel fibrogenic factor TL1a reverses established colonic fibrosis

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### Abstract

Intestinal fibrostenosis is among the hallmarks of severe Crohn's disease. Patients with certain *TNFSF15* (gene name for TL1A) variants over-express TL1A and have a higher risk of developing strictures in the small intestine. Additionally, sustained TL1a expression in mice leads to small and large intestinal fibrostenosis under colitogenic conditions. The aim of this study was to determine whether established murine colonic fibrosis could be reversed with TL1a antibody. Treatment with neutralizing TL1a antibody reversed colonic fibrosis back to the original pre-inflamed levels, potentially as result of lowered expression of connective tissue growth factor (Ctgf), Il31Ra, transforming growth factor (Tgf)  $\beta$ 1 and insulin-like growth factor-1 (Igf1). Additionally, blocking TL1a function by either neutralizing TL1a antibody or deletion of death domain receptor 3 (Dr3) reduced the number of fibroblasts and myofibroblasts, the primary cell types that mediate tissue fibrosis. Primary intestinal myofibroblasts expressed Dr3 and functionally responded to direct TL1a signaling by increasing collagen and Il31Ra expression. These data demonstrated a direct role for TL1A-DR3 signaling in tissue fibrosis and that modulation of TL1A-DR3 signaling could inhibit gut fibrosis.

### Keywords

Fibrosis; TL1A; DR3

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## Introduction

Crohn's disease (CD) is a chronic inflammatory condition with pathological features such as patchy transmural inflammation and fibrostenosis. It is accepted that chronic intestinal inflammation leads to the development of fibrosis. However, even when treated with potent anti-inflammatory therapies, approximately 20% of CD patients still develop stricturing complications that require surgical intervention.<sup>1,2</sup> A study using the *S. typhimurium* colitis model showed that despite the attenuation of intestinal inflammation with antibiotic treatment, fibrosis not only persisted, but actually progressed and that myofibroblast activation and fibrogenesis were not completely resolved by early removal of the inflammatory trigger.<sup>3</sup> Several other studies have shown that pathways independent of inflammation also drive fibrosis,<sup>4-6</sup> and that removal of the inciting inflammatory stimulus does not reverse established fibrosis.

TL1A (a protein encoded by *TNFSF15*) is a member of the TNF superfamily that binds to death domain receptor 3 (DR3) and modulates the adaptive immune response.<sup>7</sup> A particular *TNFSF15* haplotype is associated with higher TL1A expression, increased risk of CD, intestinal fibrostenosis, and greater need for surgery.<sup>8-11</sup> In addition to human reports, studies in mice also implicate the T11a/Dr3 signaling pathway in mucosal inflammation and fibrosis. As shown by our group and others previously, constitutive T11a expression in mice leads to mild spontaneous ileitis and increased collagen deposition.<sup>12-15</sup> Under colitogenic conditions, *T11a* transgenic mice develop worsened small and large intestinal inflammation and fibrostenosis.<sup>10</sup>

T11a antibody (Ab) has been shown to prevent and treat murine dextran sodium sulfate (DSS) colitis;<sup>16</sup> however, whether targeting T11a independently reduces gut fibrosis has not been established. In the present study, we used two distinct chronic colitis models, DSS and adoptive T cell transfer, to determine whether the reversal of colonic fibrosis subsequent to treatment with T11a Ab was independent of its previously reported effect in amelioration of inflammation. We found that the anti-fibrotic effect of was associated with reversal of the fibrogenic program, leading to reduced numbers of fibroblasts and myofibroblasts. Further, to determine whether the fibrogenic effect of T11a was through direct signaling of intestinal fibroblasts, we generated mice that were deficient of Dr3 (*Dr3*<sup>-/-</sup>), the only known receptor for T11a. *Dr3*<sup>-/-</sup> mice had significantly fewer intestinal fibroblasts, and showed that Dr3 was preferentially expressed on intestinal myofibroblasts. Furthermore, T11a directly increased collagen and Il31Ra expression on wildtype (WT) but not *Dr3*<sup>-/-</sup> intestinal fibroblasts. These findings indicate that TL1A/DR3 signaling pathway blockade may constitute a novel therapeutic approach for the treatment of CD, not only by inhibiting inflammation, but also directly reducing fibrogenesis and reversal of established fibrosis.

## RESULTS

### T11a Ab reversed established colonic fibrosis

Mice with constitutive T11a expression were previously shown to develop increased gut fibrosis.<sup>10, 14, 15</sup> Therefore, to determine the effect of blocking T11a signaling on colonic fibrosis, we used two mouse models of chronic colitis; adoptive T cell transfer and chronic

DSS. In the adoptive T-cell transfer model, immune-deficient *RagI*<sup>-/-</sup> mice were adoptively transferred with naïve CD4<sup>+</sup>CD45RB<sup>hi</sup> T-cells. T11a Ab (80-mg/kg) or isotype control Ab (Iso Ab, 80-mg/kg) was administered two times per week beginning on day 29 (week 4) post-transfer when colitis was established (Figure 1a, left panel). Sirius red stain was used to measure the degree of collagen deposition. By the 4<sup>th</sup> week after naïve T cell transfer, increased collagen deposition was found in the colons of mice in the Pre-Tx group compared to the baseline *Rag Co* group (Figure 1b, left and middle panels). The degree of collagen deposition in the colon was greater by the 8<sup>th</sup> week in mice receiving control Iso Ab. Treatment with T11a Ab led to significant reduction in collagen deposition compared to mice that received the Iso Ab or the Pre-Tx groups (Figure 1b, left and middle panels). Notably, collagen deposition was not significantly different when the T11a treated mice were compared to normal *Rag Co* mice (Figure 1b, left and middle panels). The Sircol assay, a dye-binding method designed to quantitatively measure acid and pepsin-soluble collagen, was used to measure colonic collagen and which showed increased soluble collagen in the Pre-Tx group compared to the *Rag Co* group (Figure 1b, right panel). Addition of control Iso Ab led to further increase in soluble collagen, whereas T11a Ab administration reduced soluble collagen to levels similar to the baseline *Rag Co* group (Figure 1b, right panel).

In the chronic DSS model, T11a (20-mg/kg) or isotype Ab (20-mg/kg) was administered twice a week beginning at day 15 when colitis was established (Figure 1a, right panel). Reduction in collagen deposition and soluble collagen in the colon with T11a Ab treatment was observed when compared to the Iso Ab and the Pre-Tx groups (Figure 1c). Together, these data indicated that blocking T11a signaling not only prevented further accumulation of collagen, but also reversed collagen to similar levels measured prior to the onset of inflammation.

### **T11a Ab administration reduced, but did not completely reverse established colitis**

We next sought to determine whether the reversal of collagen levels seen with blocking T11a signaling by T11a Ab would be paralleled by a like reduction of clinical and histologic colonic inflammation. In both chronic colitis models, administration of T11a Ab led to a significant lowering of disease activity index (DAI) as compared to the Iso Ab group (Figure 2a). However, while T11a Ab reduced collagen deposition back to the baseline, the decrease in disease activity, as measured by DAI, did not reach baseline levels and that clinical parameters of inflammation persisted (weight loss, loose stool, and stool blood).

Histologic examination of the colon revealed reduced inflammation characterized by reduced cellular infiltrate, mucin depletion, crypt abscesses, muscularis propria thickness, and architectural changes with T11a Ab therapy compared to Iso Ab group in both the adoptive transfer and chronic DSS colitis models (Figure 2b, c). The reduction in colonic histological inflammation was also significant compared to the Pre-Tx group in both models of chronic colitis (Figure 2b, c). Although histologic inflammation was reduced with T11a Ab treatment, colonic inflammation was still significantly higher compared to baseline in the WT control (Co) group in the chronic DSS colitis model (Figure 2c).

Given that there was still significantly elevated clinical disease activity in both models and that in the DSS model there was worsened histologic colitis as compared to the WT Co

despite treatment with T11a Ab, the reversal of collagen deposition is consistent with at least in part, a direct fibrosis-ameliorating effect of T11a Ab.

### **Blocking T11a-Dr3 signaling reduced numbers of intestinal fibroblasts and myofibroblasts**

Colonic myofibroblasts are a cell population involved in gut fibrogenesis.<sup>17</sup> To study the cellular mechanisms of collagen deposition reduction with T11a Ab, fibroblast expression of vimentin and myofibroblast coexpression of vimentin and alpha smooth muscle actin ( $\alpha$ SMA) were measured to assess the numbers of these cell types. After naïve T-cell transfer in both the Pre-Tx and Iso Ab groups, the numbers of colonic fibroblasts and myofibroblasts were increased (Figure 3a). However, treatment with T11a Ab led to a reduction in the number of fibroblasts and myofibroblasts to levels similar to normal *Rag Co* (Figure 3a).

In the chronic DSS model, mice treated with T11a Ab exhibited a similar reduction in the number of colonic fibroblasts and myofibroblasts compared to the Iso or the Pre-Tx groups (Figure 3b). Consistent with what was observed in the adoptive transfer model, the number of gut fibroblasts and myofibroblasts with T11a Ab treatment reduced to a level that was not statistically different from WT baseline control (Figure 3b). Because there was still significantly worsened colitis with T11a Ab treatment as compared to WT Co group in the chronic DSS colitis model, the reduced numbers of myofibroblasts and fibroblasts is consistent with at least in part, a direct consequence of neutralizing T11a, rather than solely a secondary effect through reduced inflammation.

We next assessed whether there were Dr3 expression changes in association with fibrotic changes in these murine models of chronic colitis. Immunofluorescent staining revealed increased Dr3 expression in the Pre-Tx and Iso Ab groups as compared to both baseline control groups (*Rag Co* and WT Co) and the T11a Ab treated groups in both the adoptive transfer and chronic DSS colitis models (Figure 3c, d). Notably, there was expression of Dr3 in a percentage of fibroblasts in the Pre-Tx and Isotype Ab groups (Figure 3c, d). Real-time quantitative reverse transcriptase-PCR analysis showed that the expression of Dr3 was significantly higher in the Iso Ab group as compared to mice in the both baseline control (*Rag Co* and WT Co) and T11a Ab treatment groups in both models (Figure 3e). Additionally, T11a mRNA expression was significantly increased in the Iso Ab group as compared to un-inflamed controls (*Rag Co* and WT Co) and the T11a Ab treatment groups in both the adoptive transfer and chronic DSS colitis models (Figure 3f). These results are consistent with a direct relationship between Dr3-T11a expression and increase in intestinal fibrosis.

To determine whether the reduction in the number of intestinal fibroblasts and myofibroblasts could be due to direct T11a-Dr3 signaling, we generated Dr3 deficient (*Dr3*<sup>-/-</sup>) mice (Supplementary Figure S1A, B). Although there was no spontaneous colitis in either WT or *Dr3*<sup>-/-</sup> mice up to 8 weeks of age (Figure 4a, top panel), there were significantly fewer intestinal fibroblasts in *Dr3*<sup>-/-</sup> as compared to WT littermate mice as shown by immunofluorescent staining of vimentin (Figure 4a, middle panel) and quantitation of the total recovered fibroblasts per colon (Figure 4a, bottom panel). There were no morphological differences between WT and *Dr3*<sup>-/-</sup> fibroblasts by immunofluorescent staining with vimentin and  $\alpha$ SMA (Figure 4a, middle panel) or with

light microscopy (Figure 4a, bottom panel). *Ex vivo* CellTrace Violet assay and Annexin V stain were used to determine whether the difference in the numbers of intestinal fibroblasts between WT and *Dr3*<sup>-/-</sup> mice was due to proliferation and/or apoptosis, respectively. Flow cytometric analysis showed similar rates of proliferation as evidenced by the overlapping CellTrace Violet intensity between WT and *Dr3*<sup>-/-</sup> intestinal fibroblasts (Figure 4b). No differences were observed in the rate of apoptosis between WT and *Dr3*<sup>-/-</sup> intestinal fibroblasts (Figure 4c).

### Reversal of fibrogenesis with T11a Ab administration

To study the molecular mechanisms of reversal of established intestinal fibrosis with T11a Ab treatment, we measured the expression of collagen, II31 receptor (II31Ra), fibrogenic program mediators (Tgfb1, Ctgf, Igf1, and Pten), and factors (Mmp and Timp) involved in extracellular matrix (ECM) remodeling.<sup>17, 18</sup> Lower levels of collagen expression were found in both the adoptive transfer (Table 1) and chronic DSS (Table 2) models. T11a Ab treatment resulted in normalization of the fibrogenic program mediators as evidenced by lower expression of Tgfb1 and Igf1 in the adoptive transfer (Table 1) and Tgfb1 in the chronic DSS (Table 2) models. *Il31Ra*, a gene known to be expressed on fibroblasts, also appeared to be regulated by T11a (Table 1, 2). The expression of connective tissue growth factor (Ctgf), a down-stream mediator of Tgfb signaling, was reduced with T11a Ab administration as compared to Pre-Tx and Iso Ab groups in the adoptive transfer model (Table 1).

ECM remodeling was assessed by measuring the expression of metalloproteases (Mmp) and tissue inhibitors of metalloproteases (Timp). Compared to the isotype Ab group, the expression of genes involved in ECM degradation was reduced in mice treated with T11a Ab in the adoptive transfer model (Mmp2, Mmp3; Table 1) and in the chronic DSS model (Mmp2, Mmp3, Mmp13; Table 2). Notably, the expression of Timp was lower with T11a Ab treatment in the adoptive transfer model (Timp2, Table 1) and in the chronic DSS model (Timp1, Timp2; Table 2). Although Mmp expression was decreased, it is possible that reduction in Timp expression resulted in a net increase in MMP activity that led to enhanced removal of established ECM components rather than induction of tissue damage. These results demonstrate that treatment with T11a Ab reduced the fibrogenic program, leading to decreased collagen synthesis.

### Intestinal fibroblasts express Dr3 and respond to T11a stimulation

To determine whether intestinal fibroblasts functionally respond to direct T11a signaling, mRNA levels of Dr3 were measured and found to be expressed at low levels in WT (0.0018±0.001 %β-actin) but undetectable in Dr3 deficient primary intestinal fibroblasts. We next performed flow cytometric analysis to determine whether Dr3 was expressed on vimentin<sup>+</sup>αSMA<sup>-</sup> fibroblasts or vimentin<sup>+</sup>αSMA<sup>+</sup> myofibroblasts. Our results showed that Dr3 was expressed preferentially on vimentin<sup>+</sup>αSMA<sup>+</sup> myofibroblasts as compared to vimentin<sup>+</sup>αSMA<sup>-</sup> fibroblasts (Supplementary Figure S1C). Additionally, there was a direct correlation of Dr3 expression with αSMA levels on myofibroblasts; with a higher proportion of Dr3 expression on myofibroblasts with the highest αSMA expression (Figure 5a). Additionally, sorted αSMA positive primary intestinal fibroblasts that were immunostained

with  $\alpha$ SMA and Dr3 showed co-staining of Dr3 in WT but not in Dr3 deficient myofibroblasts, indicating that Dr3 was expressed on  $\alpha$ SMA positive primary intestinal fibroblasts (Figure 5b).

To determine whether intestinal fibroblasts could respond to direct T11a stimulation, changes in the expression of collagen (Col1a2, marker for fibroblast function) and Il31Ra (*Il31Ra* is expressed on fibroblasts) were measured with the addition of exogenous T11a protein. Results showed a T11a dose-dependent increase in the expression of Col1a2 and Il31Ra in murine primary intestinal fibroblasts *ex vivo* (Figure 5c). The specificity of T11a stimulation was demonstrated by the blunted T11a induction of Col1a2 and Il31Ra in *Dr3*<sup>-/-</sup> murine intestinal fibroblasts *ex vivo* (Figure 5d). In contrast, a differential induction of Col1a2 or Il31Ra was not seen using known fibroblast growth factors (Tgf $\beta$  and Igf1) or proinflammatory stimuli (Tnf $\alpha$ ) (Figure 5d). These data indicated that intestinal fibroblasts expressed Dr3 and could functionally respond to direct T11a signaling.

## DISCUSSION

The results of this study showed that blocking T11a function reversed established colonic fibrosis in two murine chronic colitis models. Although the decrease in inflammation resulting from blocking T11a is likely responsible for a component in the reduction of fibrosis, several lines of evidence indicated that *TNFSF15* could be a profibrogenic gene that perpetuates the process of fibrosis independent of inflammation. First, constitutive T11a expression in mice resulted in increased colonic collagen deposition in the absence of detectable histologic inflammation in the colon.<sup>14, 15</sup> Second, primary ureteral strictures independent of visible inflammation or nephrolithiasis were found in transgenic mice that constitutively express T11a and the same was also found in humans with elevated TL1A levels.<sup>10</sup> Third, there is a growing body of evidence showing *TNFSF15* risk variants are associated with severe gut inflammation, intestinal fibrostenosis, and more rapid progression to surgery.<sup>8-11</sup> In line with the role of TL1A in fibrosis, in this study Dr3, the only known receptor for T11a, was shown to be expressed on primary murine intestinal fibroblasts (Supplementary Figure 1C, Figures 3c, 3d, 5a, 5b) and that stimulation with T11a can induce the expression of Col1a2 and Il31Ra on intestinal fibroblasts (Figure 5c, d). Despite the persistence of histologic inflammation with T11a Ab in the chronic DSS model (Figure 2c), collagen deposition was reversed back to the original un-inflamed baseline by inhibition of T11a (Figure 1c). The reduced collagen in the chronic DSS model may partly be due to the direct anti-fibrosis effect of T11a Ab rather than solely a secondary effect of reduced inflammation. Additionally, blocking T11a signaling can normalize the fibrogenic program, resulting in reduced expansion of fibroblasts and myofibroblasts and reversal of histologic fibrosis. To our knowledge, this is the only *in vivo* model in which blocking a single IBD associated gene product resulted not only in mitigation of inflammation, but also in the reversal of fibrosis back to the original pre-inflamed level.

Fibrogenesis depends on the balance between production of ECM proteins and its enzymatic degradation. Fibroblasts are involved in fibrogenesis and the synthesis of ECM components.<sup>17</sup> Proliferation and activation of fibroblasts to myofibroblasts lead to increased ECM production. Under normal wound healing, fibroblasts and myofibroblasts either stop



proliferating, acquire an inactive phenotype or undergo apoptosis leading to a higher rate of ECM removal.<sup>19, 20</sup> The crucial step that distinguishes normal wound healing from fibrosis may be the perpetuation of fibroblasts and myofibroblasts that results in excessive accumulation of ECM and formation of fibrotic scar.<sup>18</sup> Our data suggested that one of the fibrogenic mechanisms of T11a is through its direct role in increasing the number of intestinal fibroblasts and myofibroblasts. Interestingly, the reduced number of fibroblasts in Dr3 deficient mice was not due to differences in proliferation or apoptosis. We hypothesize that TL1A-DR3 may have a role in fibroblast formation, either through epithelial-mesenchymal transition, endothelial-mesenchymal transition, circulating fibrocytes, or possibly other mechanisms yet to be characterized.

Molecularly, the process of fibrosis results from the complex interaction between mediators involved in the fibrogenic program (Tgf $\beta$ , Ctgf, Igf1, Pten) and ECM remodeling factors (Mmp and Timp).<sup>17</sup> The profibrotic effects of Tgf $\beta$  and Igf1 are numerous and include expansion and activation of fibroblasts, collagen synthesis, and ECM remodeling.<sup>18</sup> The reduced Tgf $\beta$  and Igf1 expression with T11a Ab treatment suggested decreased fibrogenesis. Ctgf, a down-stream mediator of Tgf $\beta$  signaling, is involved in fibroblast and myofibroblast proliferation and ECM synthesis. Reduced numbers of fibroblasts, myofibroblasts and collagen synthesis with T11a Ab administration may have been in part due to reduced Ctgf expression.

IL31 signals through the heterodimeric receptor IL31RA and oncostatin M receptor (OSMR), and has been linked with the development of atopic dermatitis and allergic asthma.<sup>21–23</sup> The role of IL31RA in fibrosis has not been directly demonstrated, but is implicated by its expression in fibroblasts and myofibroblasts<sup>24–26</sup> and from a recent report showing that IL31RA is upregulated in aldosterone induced cardiac fibrosis.<sup>27</sup> IL31 has been shown to stimulate the secretion of inflammatory cytokines, chemokines and MMPs in human colonic subepithelial myofibroblasts.<sup>26</sup> The data in this report suggest that IL31Ra, a T11a regulated protein in primary intestinal fibroblasts, might play a role in fibrosis via production of inflammatory mediators or ECM remodeling.

Although T11a Ab treatment led to reduced Mmp, a major mediator for ECM degradation, the expression of Timp (Inhibitor of Mmp) was also reduced. Decreased expression of both Mmp and Timp with T11a Ab may result in the removal of the fibrotic ECM component while minimizing the unwanted effect of excessive Mmp activity, including tissue injury, which may ultimately lead to fibrosis.<sup>28–31</sup>

Although Dr3, the only known receptor for T11a, is expressed on approximately 25% of vimentin<sup>+</sup> $\alpha$ SMA<sup>+</sup> myofibroblasts (Supplementary Figure S1C), this represents a small proportion of the total intestinal fibroblasts in mice with colitis (approximately 2–4%) and without colitis (approximately 1–2%); thus, raising the issue of biological relevance of direct T11a-Dr3 signaling in intestinal fibrosis. Additionally, the relative contribution of direct (effect on fibrosis independent of inflammation) and indirect (effect on fibrosis through inflammation) fibrogenic effects of T11a was not determined in this study. Constitutive *Dr3* knockout in *Rag* sufficient or deficient genetic background would still not separate the relative contribution of T11a-Dr3 signaling on fibroblast from other inflammatory cells that

also express Dr3. One approach to assess the biological relevance of direct T11a-Dr3 signaling on fibroblasts in intestinal fibrosis would be to create mice with tissue specific Dr3 deficiency in fibroblasts, but not in other cell types. The fibroblast specific Dr3 knockout mouse would enable assessment of the *in vivo* fibrogenic role of T11a-Dr3 signaling on fibroblasts independent from its role inflammatory cells.

Existing medical therapies that target intestinal inflammation have a relatively poor record for treating CD complications such as fibrostenosis and altering the natural history of CD.<sup>32–34</sup> Novel therapies are needed that target both inflammation and fibrosis. T11a Ab is successful in murine experimental models in mitigation of intestinal inflammation and reversal of fibrosis back to baseline levels. Targeting the TL1A signaling pathway may lead to a novel class of therapeutics that could not only treat inflammation, but also reverse the fibrogenic process and alter the natural history of CD.

## METHODS

### Induction of chronic colitis and treatment

C57BL/6J mice were purchased from the Jackson Laboratory. Chronic DSS colitis using four cycles of 2.5% DSS in drinking water was performed as described.<sup>10</sup> In the adoptive-transfer model, colitis was induced by intraperitoneal injection of 500,000 CD4<sup>+</sup>CD45RB<sup>hi</sup> naïve T-cells isolated from WT mice to *Rag1*<sup>-/-</sup> mice. Hamster anti-mouse T11a Ab (12F6A, TEVA, North Wales, PA) that specifically blocked the function of T11a but did not mediate depletion of activated T cells (data not shown) was used. For the adoptive transfer model, T11a Ab or control immunoglobulin (Ig) G (Leinco Technologies, St. Louis, MO) were injected into mice intraperitoneally twice per week beginning on day day 29 (Figure 1a, left panel). For the chronic DSS model, T11a Ab or control IgG Ab were injected intraperitoneally twice per week beginning on day 15 (Figure 1a, right panel). Baseline controls (Rag Co or WT Co) were mice analyzed prior to DSS treatment or adoptive transfer of naïve T-cells. Pre-treatment (Pre-Tx) controls were mice analyzed at day 14 for the chronic DSS model and day 28 for the adoptive-transfer model. Treatment groups were mice analyzed at day 28 for the chronic DSS model and day 56 for the adoptive transfer model (Figure 1a). All mice were maintained under specific pathogen-free conditions in the Animal Facility at Cedars-Sinai Medical Center (CSMC). This study was carried out in strict accordance with the Guide for the Care and Use of Laboratory Animals of the National Institutes of Health. Animal studies were approved by the CSMC Animal Care and Use Committee (protocol 3813).

DAI and histopathological analyses. DAI was calculated by scoring from 0 to 4 abnormalities regarding changes in body weight (0, no weight loss; 1, 1–5% weight loss; 2, 5–10% weight loss; 3, 10–15% weight loss; 4, more than 15% weight loss), stool consistency (0, firm dry stool; 1, moist stool; 2, soft adherent stool; 3, large soft pliable stool; 4, liquid stool), stool blood performed on Hemocult Sensa (Beckman Coulter, Brea, CA; 0, no color; 1, flecks of blue; 2, up to 50% blue; 3, more than 50% blue; 4, gross red blood) and summing the results.<sup>10</sup> DAI score was determined every other day for the DSS model and twice a week for the adoptive-transfer model. Tissue samples were processed and stained with hematoxylin and eosin (H&E) by the CSMC Histology-Core. Sirius red staining



was performed using the NovaUltra Sirius Red Stain Kit according to manufacturer's protocol (IHC World, Woodstock, MD). Immunofluorescent stain was performed on 4  $\mu\text{m}$  frozen sections fixed with 10% formalin, blocked in 10% BSA, 0.1% Triton X-100 TBST, and stained overnight at 4 degree C with primary antibodies: rabbit polyclonal  $\alpha\text{SMA}$  Ab (Abcam, Cambridge, MA) at 1:100 dilution, anti-mouse Dr3 Ab conjugated with PE (Biolegend, San Diego, CA) at 1:50 dilution, and  $\alpha\text{-Vimentin}$  Ab (Covance, San Diego, CA) at 1:2000 dilution. The next day, secondary antibody was added for 2 hours at room temperature with donkey  $\alpha\text{-rabbit IgG}$  and goat  $\alpha\text{-chicken IgY}$  (Abcam, Cambridge, MA) and goat  $\alpha\text{-chicken IgY\&L}$  (Abcam, Cambridge, MA) secondary antibodies at 1:500 dilution. Histopathological scores were assigned in a blinded manner by two trained animal pathologists (DQS and JC) as described.<sup>10</sup> Area of collagen deposition was determined using the ImageJ 1.47 software package.<sup>35</sup> The numbers of fibroblasts and myofibroblasts were quantitated from colonic mucosa and submucosa using a Leica TCS SP spectral confocal microscope.

### Sircol Assay

Acid and pepsin-soluble collagen from mid-colon (3 mm piece) was measured using the Sircol Soluble Collagen Assay according to the manufacturer's protocol (Biocolor, Northern Ireland, UK). Acid and pepsin-soluble collagen were measured using a colorimetric absorbance using the 550 nm filter and concentration determined using the collagen standard provided in the Sircol Soluble Collagen Assay Kit. Recovered mid-colon collagen is normalized to total acid and pepsin-soluble protein measured using Bradford Protein Assay Kit according to manufacturer's protocol (Thermo Scientific, Rockford, IL).

### Generation of *Dr3*<sup>-/-</sup> mice

Cloning of *Dr3* targeting vector and generation of *Dr3*<sup>+/-</sup> founder mice were performed in collaboration with genOway (genOway, Lyon, France). Briefly, *Dr3* endogenous locus containing 1.5 kb upstream of exon 1 and 3 kb downstream of exon 8 were generated by PCR amplification using genomic DNA from C57BL/6J mice and cloned into the pCR4-TOPO vector (Invitrogen, Carlsbad, CA). Subsequently, two *loxP* sites were inserted flanking *Dr3* exons 2 to 5 (Supplementary Figure S1A). Positive selection neomycin gene flanked by FRT sites was inserted to the intron between exon 1 and 2 to generate the targeting vector (Supplementary Figure S1A). Every step of the cloning process was validated through restriction enzyme analysis and sequencing. The *Dr3* gene targeting construct was linearized and electroporated into genOway proprietary embryonic stem (ES) cells with C57BL/6J background. Homologous recombinants were selected by G418 and confirmed by Southern blot analysis. ES clones with correct 5' and 3' recombination were microinjected into C57BL/6J blastocysts and introduced into pseudopregnant C57BL/6J mice. Male chimeric offspring were bred to obtain germ line mutant mice which were then bred to *Flpe* delete mouse strain to remove the neomycin cassette, then bred to Cre delete mice to excise the *loxP* flanked sequences (Supplementary Figure S1A), confirmed by Southern blot, and maintained on the C57BL/6J genetic background.

## Expression analysis

Total RNA was isolated using RNeasy Microarray Tissue Mini Kit (Qiagen, Valencia, CA), and reverse-transcription polymerase chain reaction (RT-PCR) was performed using RT2 HT First Strand and gene expression was measured using the RT2 Custom Fibrosis Array CAPM11248 (Qiagen, Valencia, CA) kits per manufacturer's protocols. Validated Dr3 qPCR assay Mm.PT.51.17321439, Il31Ra qPCR assay Mm.PT.56a.32787326, Tl1a qPCR assay Mm.PT.51.17321439, Col1a2 qPCR assay Mm.PT.51.5206680, and  $\beta$ -actin qPCR assay Mm.PT.39a.22214843 were purchased from IDT Technologies (Skokie, IL).

## Cell isolation, culture, flow cytometry, and cell sorting

Isolation and culture of lamina propria mononuclear cells (LPMC) and mesenteric lymph node (MLN) cells were carried out as previously reported.<sup>14</sup> The whole colon and the distal 10 cm of the ileum were used for LPMC isolation. CD4<sup>+</sup> T-cells were isolated from spleen and MLN by negative selection (StemCell Technologies Inc, Vancouver, Canada). MLN, LPMC and CD4<sup>+</sup> T-cells were cultured in 96-well round-bottom plates at  $1.5 \times 10^6$  cells/mL of T-cell medium (RPMI 1640 containing 2mmol/L glutamine, 25 mmol/L HEPES, 10% fetal bovine serum, 100  $\mu$ M NEAA, 1 mM sodium pyruvate, 50  $\mu$ M  $\beta$ -mercaptoethanol, 50  $\mu$ g/mL gentamicin) under conditions indicated in Supplementary Figure 1 with Il12 (5 ng/mL), Tnf $\alpha$  (5 ng/mL), Il6 (100 ng/mL), anti-CD3 $\epsilon$  (0.5  $\mu$ g/mL, 145-2C11), anti-CD28 (1  $\mu$ g/mL, 37.51), Tl1a Ab (50  $\mu$ g/mL, 12F6A, TEVA, North Wales, PA) and control isotype immunoglobulin (Ig) G (50  $\mu$ g/mL, Leinco Technologies, St. Louis, MO). Cells and media were harvested 48 hours after culture and assessed for Ifn $\gamma$  production via ELISA kit (eBioscience, San Diego, CA), proliferation and apoptosis assay as described below.

Mouse primary colonic fibroblasts were isolated from colon that were incubated in 1 mM DTT (Fisher Scientific, Tustin, CA), 37 °C, 15 min, and then 1 mM DTT with 5 mM EDTA (Promega, Madison, WI), 37 °C, 30 min. The remaining colonic tissues were rinsed by 1 $\times$  HBSS (Corning Cellgro, Swedesboro, NJ), minced and then digested for 30 min at 37 °C with 1.5 mg/mL Collagenase II (Worthington, Lakewood, NJ), 0.3 mg/mL DNase I and 3 mg/mL Hyaluronidase (Sigma, St. Louis, MO) in DMEM (Corning Cellgro, Swedesboro, NJ). The isolated cells were cultured in DMEM supplemented with 10% FCS, Penicillin/Streptomycin (100 IU/mL), Fungizone (0.5  $\mu$ g/mL). The number of fibroblasts was counted using a hemocytometer and represented as total fibroblast recovered per colon. The criteria to score cells as fibroblast are based upon morphology and positive vimentin staining. Primary intestinal fibroblasts were used at passage 2. Intracellular staining and antibodies used were described previously.<sup>14, 36</sup> Growth factors Tgf $\beta$  100 ng/mL and Igf1 10 ng/mL (Cell Signaling, Danvers, MA) and cytokines Tl1a 10–200 or 100 ng/mL (R&D Systems, Minneapolis, MN) and Tnf $\alpha$  5 ng/mL (Biolegend, San Diego, CA) were added to passage 2 fibroblasts for 4 hours before harvesting for expression analysis.

For flow cytometry, cells were acquired on a LSR II flow-cytometer (BD Biosciences, San Jose, CA) and analyzed using FlowJo analysis software. For cell sorting, cells were sorted using the MoFlo Cell Sorter with the assistance from Flow Cytometry Core Facility at CSMC.

### **Ex vivo intestinal fibroblast proliferation and apoptosis assay**

Primary intestinal fibroblasts were isolated and stained with CellTrace Violet (Invitrogen, Carlsbad, CA) per manufacturer's instructions. Stained cells were then incubated with 100 ng/mL of T11a in DMEM supplemented with 10% FCS, Penicillin/Streptomycin (100 IU/mL), and Fungizone (0.5 µg/mL). After 48 hours, cultured intestinal fibroblasts were stained using Annexin V Apoptosis Detection Kit (eBioscience, San Diego, CA) per manufacturer's instructions. After Annexin V stain, fibroblasts were harvested, washed and fixed with 2% paraformaldehyde and subjected to flow cytometric analysis with BD LSR II flowcytometer (BD Biosciences, San Jose, CA) and analyzed by FlowJo software.

### **Statistical analysis**

Data are presented as the mean ± standard deviation (SD). Comparison between two groups was performed by a two-tailed Fisher's Exact Test for categorical variables and Student's *t*-test for continuous variables. Parametric and non-parametric tests were used depending on the fulfillment of the test assumptions. Comparison between three groups was done using ANOVA, followed by pair wise posthoc analysis with Turkey's HSD and Behrens-fisher-Test correction for the multiple comparisons. *P* < 0.05 was considered significant.

### **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

### **ACKNOWLEDGEMENTS**

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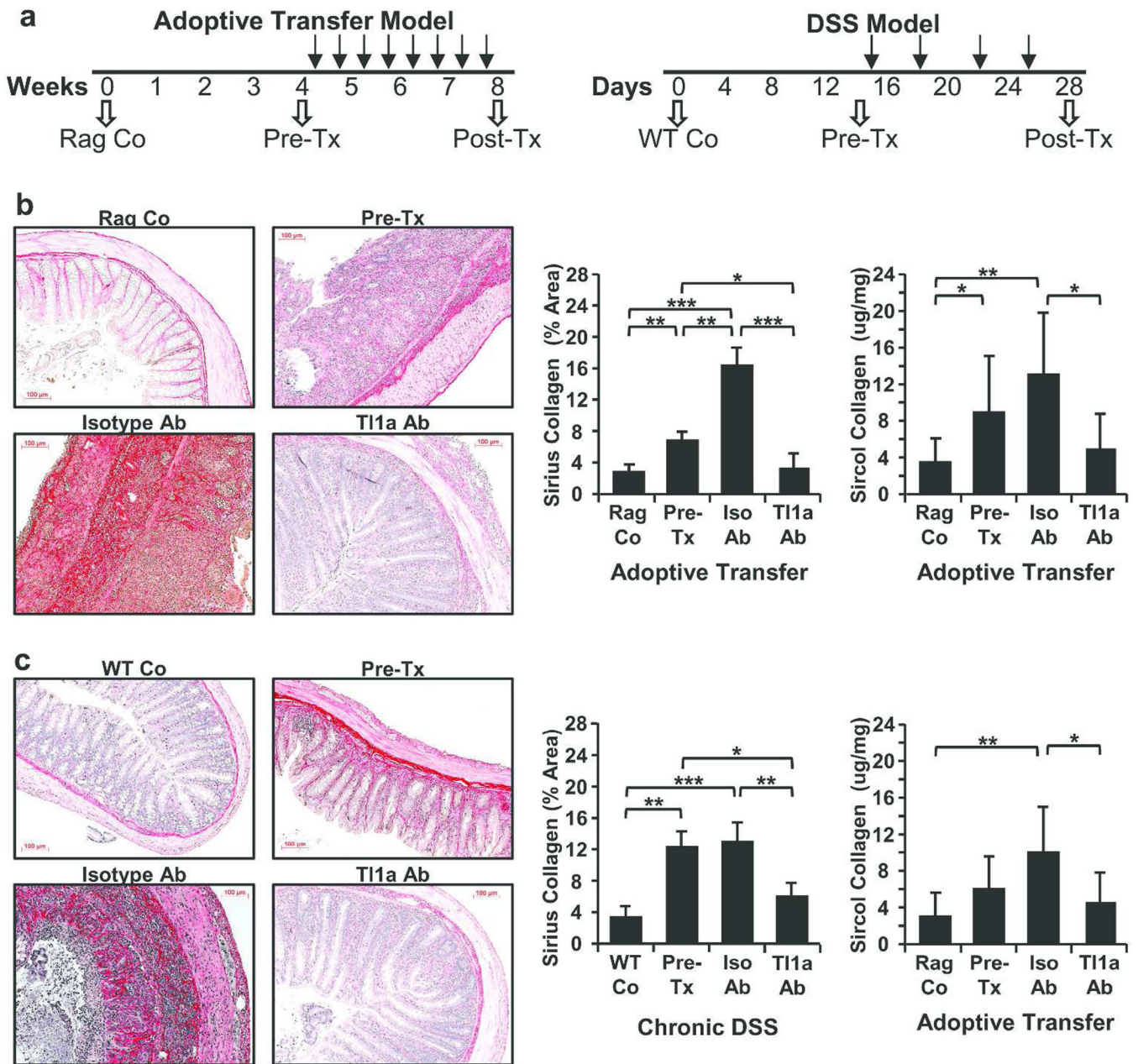
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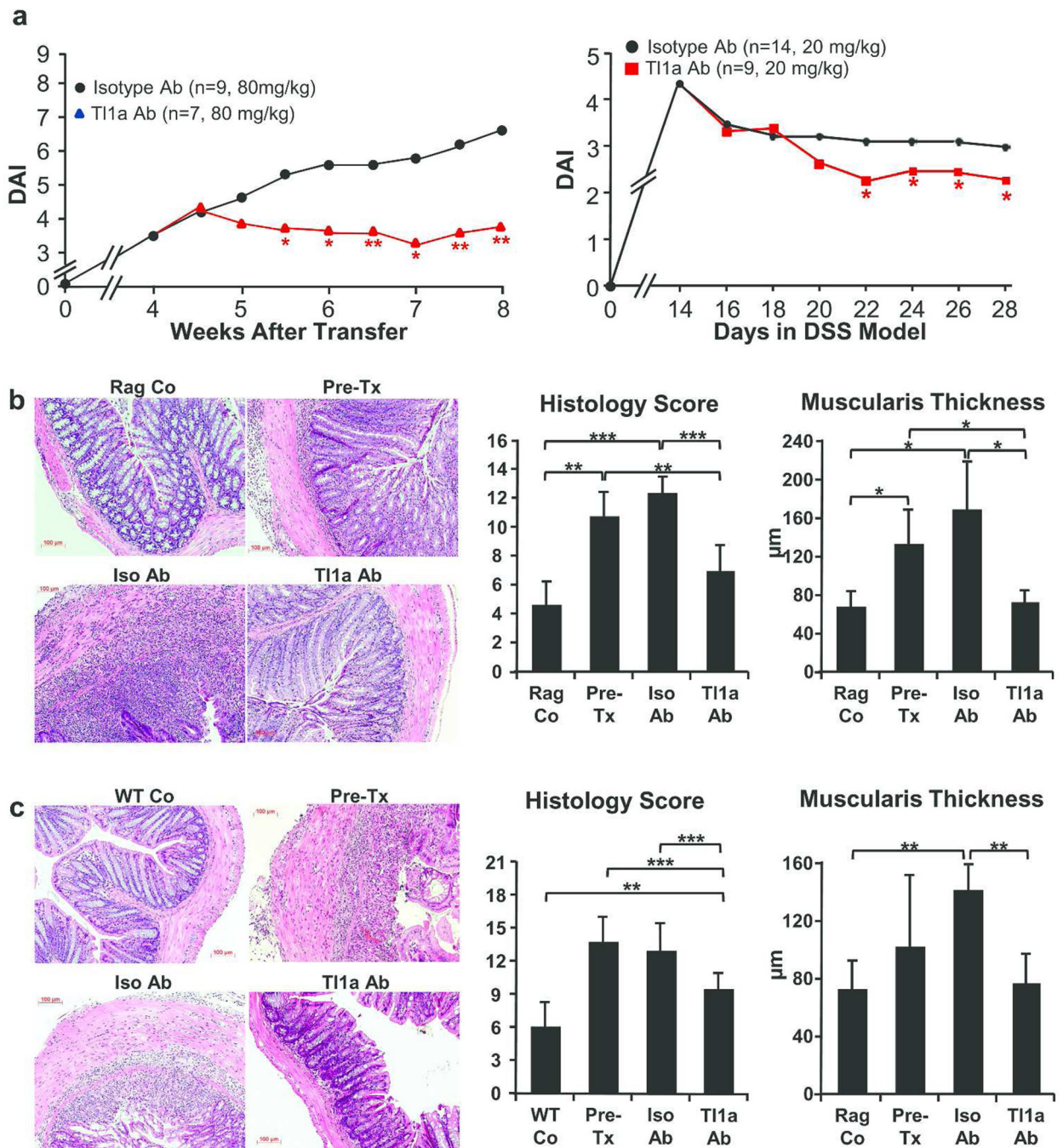




**Figure 1.** Reversal of established fibrosis with T11a Ab therapy. (a) T11a Ab treatment schematics for the adoptive transfer model (left panel) and the chronic DSS colitis model (right panel); baseline control mice (*Rag* n=5 or WT Co n=5), pre-treatment group (Pre-Tx, n=5 for transfer, n=6 for DSS), post treatment group (Post-Tx, n=7–14). Representative Sirius red staining of collagen deposition in mid-colon tissue sections at 100× magnification is shown for adoptive transfer model in (b, left panels) and chronic DSS model in (c, left panels). Percent of colon with collagen staining were quantitated and expressed as mean ± SD for the adoptive transfer model in (b, middle panel) and for the chronic DSS model in (c, middle panel). Quantitation of soluble collagen from the colon were determined and expressed as

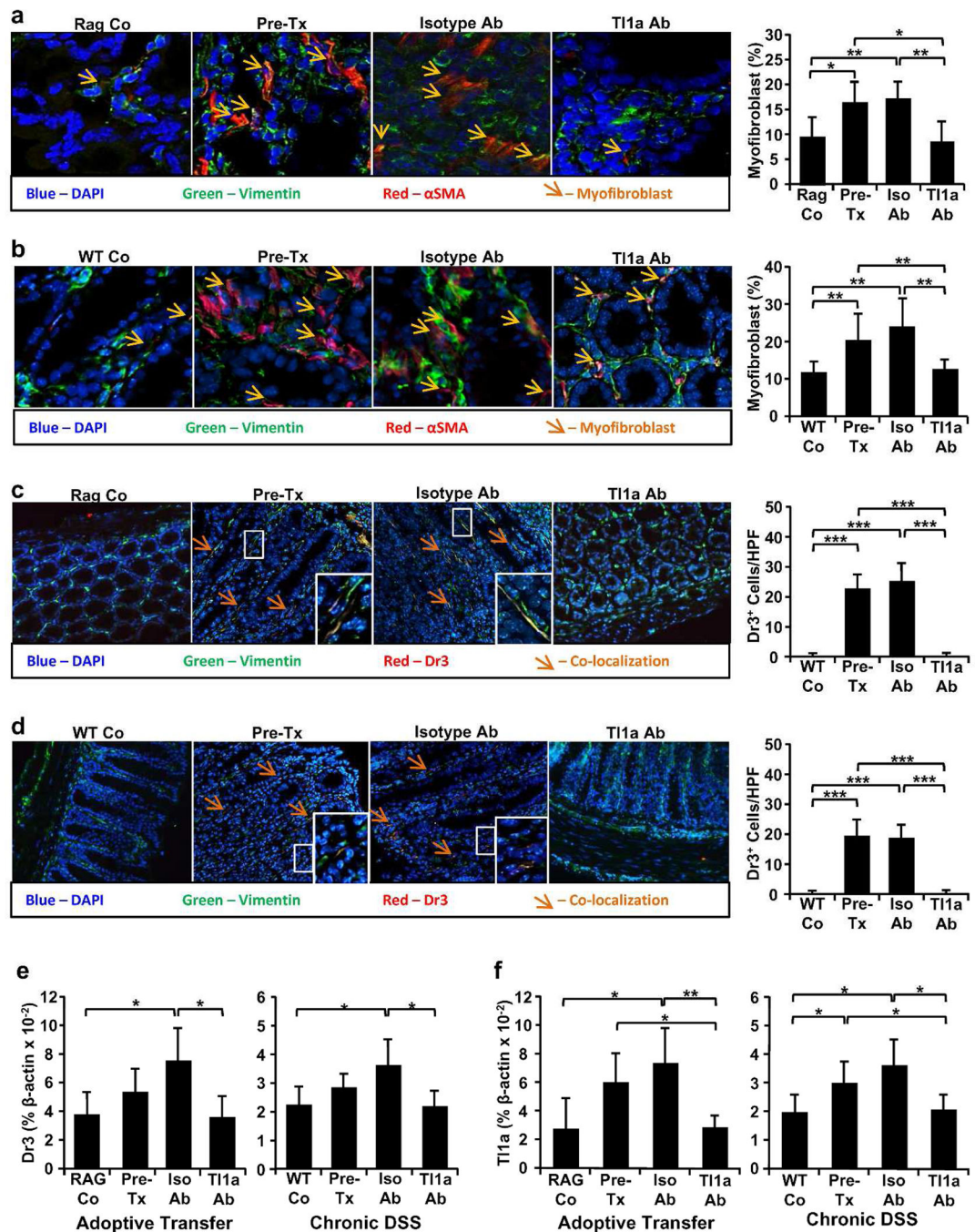


mean  $\pm$  SD for the adoptive transfer model in (**b**, right panel) and for the chronic DSS model in (**c**, right panel). At least 20 independent fields per group are scored and data are expressed as mean  $\pm$  SD. \*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001.



**Figure 2.**

T11a Ab reduced established chronic colitis. (a) DAI of the adoptive transfer (left panel) and chronic DSS colitis model (right panel) is compared between isotype Ab and T11a Ab treated groups. Representative H&E stained mid-colon sections from the adoptive transfer model (b) and chronic DSS colitis model (c) at 100× magnification is shown and quantitative histology scores (b and c, middle panels) and thickness of muscularis propria (b and c, right panels) are shown. At least 20 independent fields per group are scored and data are expressed as mean ± SD. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ .

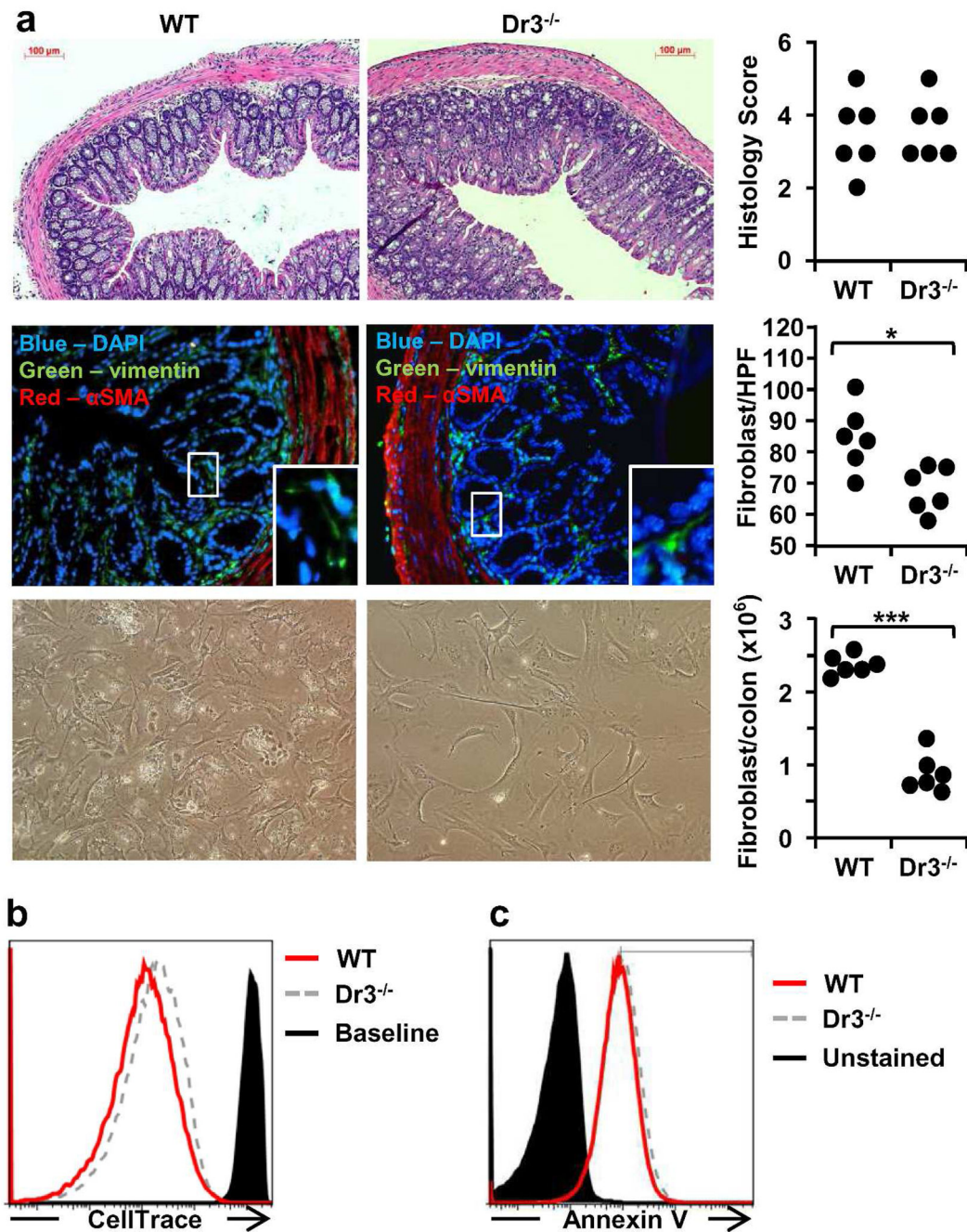
**Figure 3.**

T11a Ab reduced myofibroblast number and expression of Dr3 and T11a. Representative immunofluorescent staining of vimentin (green) and  $\alpha$ SMA (red) from mid-colon sections from the adoptive transfer model (**a**) and chronic DSS model (**b**) at 630 $\times$  magnification are shown. Orange arrows denote myofibroblasts that co-express vimentin and  $\alpha$ SMA.

Percentages of myofibroblasts from the mid-colon sections were quantitated and expressed as mean  $\pm$  SD for the adoptive transfer model (**a**, right panel) and chronic DSS model (**b**, right panel). At least 10 independent fields were scored per group for (**a**) and (**b**).

Representative immunofluorescent staining of vimentin (green) and Dr3 (red) from mid-colon sections are shown from the adoptive transfer model (c) and chronic DSS model (d). Figure insets for (c) and (d) are larger view of the images that were acquired at 200× magnification. At least 8 independent fields were quantitated per group and plotted as Dr3<sup>+</sup> cells per high power fields (HPF). Colonic Dr3 (e) and T11a (f) mRNA was quantitated and shown as mean ± SD (n=5–14). T11a Ab treated groups are compared to baseline *Rag* Co, Wt Co, Pre-Tx, and Iso Ab experimental groups. \**P* < 0.05, \*\**P* < 0.01, \*\*\**P* < 0.001.





**Figure 4.** Reduced intestinal fibroblasts with Dr3 deficiency. (a) Representative H&E stained colon at 100 $\times$  magnification with quantitation of inflammation is shown on the upper panels. Representative Vimentin/ $\alpha$ SMA stained colon at 200 $\times$  magnification (insets are larger view at 200 $\times$  magnification) with quantitation of fibroblasts per HPF is shown in the middle panels. Representative photographs of intestinal fibroblasts recovered from littermate WT and Dr3<sup>-/-</sup> colon and individual total fibroblasts per colon are shown (a, bottom panels). Representative flow cytometric histograms of proliferating fibroblasts (b) and fibroblasts

undergoing apoptosis (c) from WT and *Dr3*<sup>-/-</sup> mice are shown. Decreased CellTrace violet fluorescence intensity indicates proliferation. Increased Annexin V staining indicates apoptosis. Representative flow cytometric histograms of at least 6 independent experiments with similar results are shown. \**P* < 0.05, \*\**P* < 0.01, \*\*\**P* < 0.001.

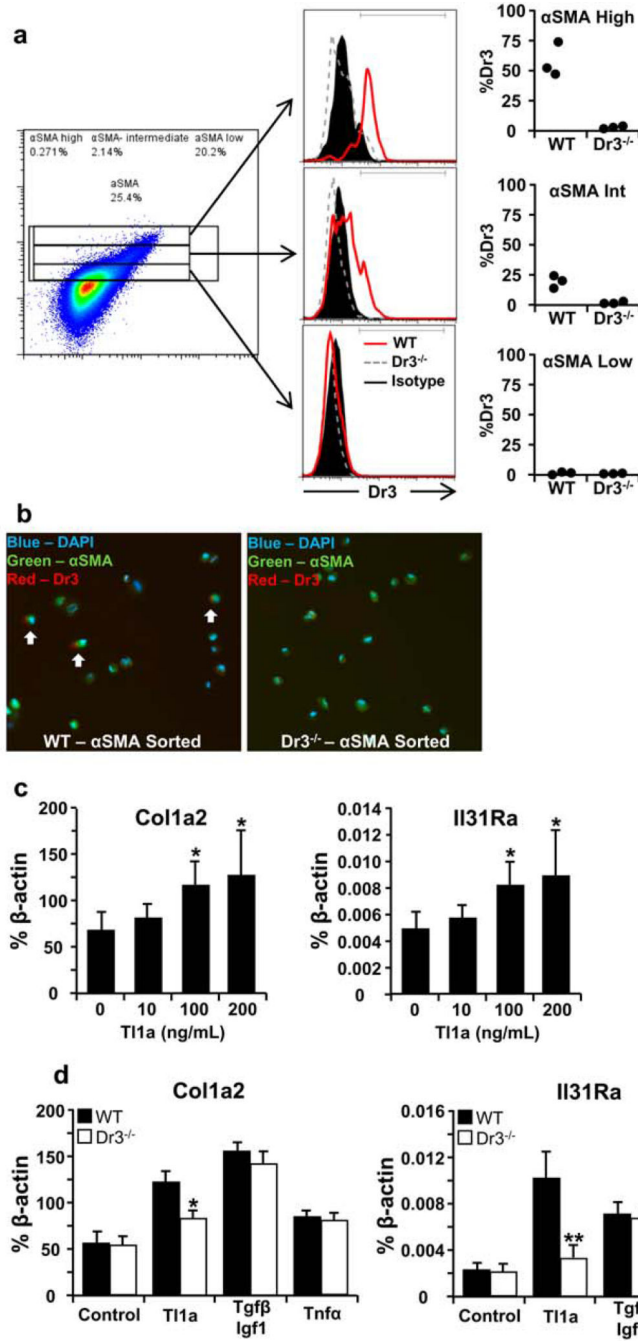
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**Figure 5.** Intestinal fibroblasts express Dr3 and respond to T11a stimulation. **(a)** Primary intestinal fibroblasts were stained with Dr3, αSMA and vimentin and analyzed by flow cytometry. Fibroblasts expressing high, intermediate, and low αSMA were gated as shown and Dr3 staining is preferentially found in αSMA high > intermediate > low. Three independent experiments were performed. **(b)** Data are representative of 3 independent sorted αSMA positive myofibroblasts at 200× magnification. There was co-staining of Dr3 in WT, but not in Dr3 deficient αSMA positive myofibroblasts. **(c)** Expression of Col1a2 and Il31Ra

mRNA in WT primary intestinal fibroblasts with increasing T11a stimulation (0–200 ng/mL) and represented as mean  $\pm$  SD are shown (n=3). (d) Induction of Col1a2 and Il31Ra mRNA by T11a, Tgf $\beta$ /Igf1, and Tnf $\alpha$  in WT and *Dr3*<sup>-/-</sup> intestinal are shown and represented as mean  $\pm$  SD (n=3). \**P* < 0.05, \*\**P* < 0.01.

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**Table 1**

Expression analysis of fibrosis mediators in the adoptive transfer colitis model.

	baseline		Pre-Tx		Iso Ab		TIIa Ab - 80 mg/kg		<i>p</i> vs.
	% $\beta$ -actin n=6	% $\beta$ -actin n=6	% $\beta$ -actin n=6	% $\beta$ -actin n=7	% $\beta$ -actin n=6	% $\beta$ -actin n=6	Reg	Pre-Tx Iso Ab	
<b>colla1</b>	0.19 $\pm$ 0.12	0.19 $\pm$ 0.12	0.19 $\pm$ 0.12	0.21 $\pm$ 0.10	0.11 $\pm$ 0.03	0.11 $\pm$ 0.03	ns	0.024	0.03
<b>colla2</b>	0.49 $\pm$ 0.29	0.76 $\pm$ 0.32	0.76 $\pm$ 0.32	1.23 $\pm$ 0.78	0.39 $\pm$ 0.12	0.39 $\pm$ 0.12	ns	0.024	0.026
<b>col3a1</b>	12.69 $\pm$ 3.61	16.45 $\pm$ 3.93	16.45 $\pm$ 3.93	16.08 $\pm$ 4.04	9.66 $\pm$ 3.44	9.66 $\pm$ 3.44	ns	0.0073	0.014
<b>col4a1</b>	1.54 $\pm$ 0.32	1.95 $\pm$ 0.32	1.95 $\pm$ 0.32	1.88 $\pm$ 0.84	1.19 $\pm$ 0.33	1.19 $\pm$ 0.33	ns	0.00055	ns
<b>Tgfb1</b>	0.16 $\pm$ 0.06	0.40 $\pm$ 0.16	0.40 $\pm$ 0.16	0.50 $\pm$ 0.17	0.25 $\pm$ 0.06	0.25 $\pm$ 0.06	0.018	0.046	0.003
<b>Ctgf</b>	0.66 $\pm$ 0.13	1.04 $\pm$ 0.40	1.04 $\pm$ 0.40	1.04 $\pm$ 0.32	0.54 $\pm$ 0.08	0.54 $\pm$ 0.08	ns	0.021	0.007
<b>Igfl</b>	0.32 $\pm$ 0.06	0.53 $\pm$ 0.18	0.53 $\pm$ 0.18	0.73 $\pm$ 0.36	0.41 $\pm$ 0.15	0.41 $\pm$ 0.15	ns	ns	0.047
<b>Pten</b>	3.80 $\pm$ 0.75	2.28 $\pm$ 0.53	2.28 $\pm$ 0.53	1.86 $\pm$ 0.25	2.03 $\pm$ 0.67	2.03 $\pm$ 0.67	0.0015	ns	ns
<b>IL13Ra</b>	0.003 $\pm$ 0.001	0.005 $\pm$ 0.002	0.005 $\pm$ 0.002	0.007 $\pm$ 0.003	0.004 $\pm$ 0.001	0.004 $\pm$ 0.001	ns	ns	0.034
<b>Mmp2</b>	0.32 $\pm$ 0.072	0.43 $\pm$ 0.12	0.43 $\pm$ 0.12	0.44 $\pm$ 0.13	0.28 $\pm$ 0.07	0.28 $\pm$ 0.07	ns	0.015	0.015
<b>Mmp3</b>	0.046 $\pm$ 0.022	1.15 $\pm$ 1.15	1.15 $\pm$ 1.15	1.18 $\pm$ 0.70	0.34 $\pm$ 0.37	0.34 $\pm$ 0.37	0.043	ns	0.036
<b>Mmp13</b>	0.047 $\pm$ 0.015	0.20 $\pm$ 0.17	0.20 $\pm$ 0.17	0.18 $\pm$ 0.08	0.10 $\pm$ 0.06	0.10 $\pm$ 0.06	ns	ns	ns
<b>Timp1</b>	0.038 $\pm$ 0.014	0.19 $\pm$ 0.24	0.19 $\pm$ 0.24	0.18 $\pm$ 0.12	0.14 $\pm$ 0.07	0.14 $\pm$ 0.07	0.016	ns	ns
<b>Timp2</b>	1.11 $\pm$ 0.27	0.86 $\pm$ 0.15	0.86 $\pm$ 0.15	0.79 $\pm$ 0.12	0.59 $\pm$ 0.18	0.59 $\pm$ 0.18	0.001	0.001	0.048

ns = not significant

Table 2

Expression analysis of fibrosis mediators in the DSS model

	WT		Pre-Tx Co		Isotype Co		TIIa Ab - 20 mg/kg		P vs.
	% $\beta$ -actin n=6	% $\beta$ -actin n=5	% $\beta$ -actin n=5	% $\beta$ -actin n=5	% $\beta$ -actin n=5	% $\beta$ -actin n=5	WT	Pre-Tx	
<b>colla1</b>	0.54 $\pm$ 0.45	0.55 $\pm$ 0.36	0.67 $\pm$ 0.45	0.30 $\pm$ 0.17	ns	ns	ns	ns	ns
<b>colla2</b>	0.67 $\pm$ 0.26	1.20 $\pm$ 0.94	1.19 $\pm$ 0.93	0.63 $\pm$ 0.31	ns	ns	ns	ns	ns
<b>col3a1</b>	35.79 $\pm$ 10.95	38.64 $\pm$ 18.02	35.18 $\pm$ 9.74	23.28 $\pm$ 3.47	0.044	ns	ns	ns	0.036
<b>col4a1</b>	2.60 $\pm$ 1.08	2.62 $\pm$ 1.37	2.70 $\pm$ 0.54	1.83 $\pm$ 0.10	ns	ns	ns	ns	0.010
<b>Tgfb1</b>	0.21 $\pm$ .06	0.38 $\pm$ 0.15	0.43 $\pm$ 0.03	0.22 $\pm$ 0.04	ns	0.041	ns	0.041	6.943E-05
<b>Ctgf</b>	0.97 $\pm$ .27	1.1 $\pm$ .32	1.14 $\pm$ .36	0.84 $\pm$ .21	ns	ns	ns	ns	ns
<b>Igf1</b>	0.48 $\pm$ 0.18	0.85 $\pm$ 0.63	1.09 $\pm$ 0.52	0.65 $\pm$ 0.28	ns	ns	ns	ns	ns
<b>Pten</b>	0.004 $\pm$ 0.003	0.008 $\pm$ 0.004	0.012 $\pm$ 0.003	0.008 $\pm$ 0.002	ns	ns	ns	ns	ns
<b>Il31Ra</b>	3.13 $\pm$ 0.65	2.56 $\pm$ 0.61	2.67 $\pm$ 0.97	2.69 $\pm$ 0.31	0.020	ns	ns	ns	0.046
<b>Mmp2</b>	0.52 $\pm$ 0.16	0.60 $\pm$ 0.29	1.21 $\pm$ 0.38	0.52 $\pm$ 0.25	ns	ns	ns	ns	0.007
<b>Mmp3</b>	0.03 $\pm$ 0.015	2.07 $\pm$ 3.70	1.91 $\pm$ 0.10	0.42 $\pm$ 0.27	0.003	ns	ns	ns	1.48E-05
<b>Mmp13</b>	0.05 $\pm$ 0.011	0.61 $\pm$ 1.07	1.02 $\pm$ 0.21	0.21 $\pm$ 0.15	0.014	ns	ns	ns	0.0003
<b>Timp1</b>	0.04 $\pm$ 0.024	0.32 $\pm$ 0.37	0.33 $\pm$ 0.06	0.15 $\pm$ 0.07	0.003	ns	ns	ns	0.004
<b>Timp2</b>	1.02 $\pm$ 0.32	0.88 $\pm$ 0.17	1.12 $\pm$ 0.11	0.74 $\pm$ 0.14	ns	ns	ns	ns	0.005

ns = not significant