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T regulatory cells and TGF- β 1: Predictors of the host response in mesh complications

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Abstract

Polypropylene mesh is frequently used in urogynecology procedures; however, pain and mesh exposure into the vagina occur in ~10% of cases. Mesh-induced pain, which occurs with or without exposure, persists after removal in 50% of cases. Chronic pain history predicts poor response to mesh removal but only a fraction have this diagnosis. We hypothesize that mesh induced pain is correlated with fibrosis and failure to improve with a heightened inflammatory and fibrotic host response. Women undergoing mesh removal were offered participation in a mesh biorepository. Standardized questionnaires including visual analog scale (VAS) pelvic pain scores were completed at enrollment and 6 months after removal. Responders were considered those with 13 mm VAS improvement. 30 mesh-tissue explants were randomly selected for analysis. Samples were labeled for CD8, CD4 (T_h) and FoxP3 (T_{regs}). Peri-fiber collagen deposition (fibrosis) was measured using a customized semi-quantitative assay. Concentrations of TGF- β 1, bFGF, MCP-1, PDGF-BB, and IGFBP-1 in tissue were determined by immunoassay and compared to vaginal control biopsies with pathway analysis. VAS pain scores were correlated with degree of histologic fibrosis. Responders had more T_{regs} (7.8 vs 0.3 per mm², $p = 0.036$) and patients were 1.6 times as likely to be a responder for every additional T_{reg}/mm² ($p = 0.05$). Pro-fibrotic TGF- β 1 was doubled in nonresponders ($p = 0.032$). On pathway analysis, decreased bFGF and increased PDGF-BB provide a possible mechanism for upregulation of TGF- β 1. In conclusion, fibrosis is a

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Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

Supplementary materials

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plausible mechanism of pain complications and the adaptive immune response likely contributes to mitigation/prevention of complications and recovery in affected patients.

Keywords

Polypropylene mesh; T regulatory cells; TGF- β 1; Fibrosis; Prolapse

1. Introduction

Pelvic organ prolapse (POP) and stress urinary incontinence (SUI) are associated with severe impairments in female health, including sexual dysfunction, reduced activity, and reduced socialization [1,2]. Up to 20% of women will undergo surgery for POP or SUI in their lifetime [3,4]. Polypropylene mesh is widely used in prolapse surgery to overcome the high failure rate of suture repairs and in SUI to minimize the invasiveness of the procedure; however, even when the implanted mesh has optimal porosity and stiffness, women are still affected by mesh-induced pain and mesh exposure into the vagina in ~10% of cases [5]. Mesh-induced pain is difficult to treat and persists in 50% of cases, despite medication, physical therapy and multiple surgeries for mesh removal [6–9]. Mesh-induced pain can occur with or without exposure of the mesh through the vaginal epithelium. Although mesh-induced pain and exposure are sometimes considered two distinct complication phenotypes, there is likely pathophysiologic and symptomatic overlap between them that is poorly characterized. Understanding the pathophysiology of mesh complications and characterizing who will respond to removal has the potential to reduce poor outcomes related to mesh implants and better treat those who have already experienced these complications.

Fibrosis, driven by increased TGF- β 1, appears to be an important element present in explants from women with pain and exposure mesh complications [10]. In mesh complications, pore collapse can result in crowding of mesh fibers with an overlapping foreign body response, bridging fibrosis, encapsulation and myofibroblast-induced contracture [11,12]. Indeed, both pain and exposure complications demonstrate increased collagen surrounding mesh fibers that suggests collagen over-production by fibroblasts. Fibrosis and contracture have also been implicated in adverse consequences for many other medical implants including artificial heart valves, breast implants, and joint replacements, supporting their likely importance in mesh complications [13,14].

Soluble factors such as growth factors, cytokines and chemokines play a critical role in normal wound healing and are often altered in states of chronic inflammation, chronic or heightened foreign body responses and dysregulated wound healing [15,16]. For example, platelet-derived growth factor BB (PDGF-BB) is a profibrotic growth factor that stimulates fibroblast proliferation and increases TGF- β 1 activity, which in turn stimulates fibroblast activation. Fibroblast growth factor (bFGF) antagonizes TGF- β 1 signaling and promotes angiogenesis during wound healing [17,18]. Monocyte chemoattractant protein (MCP-1) is a chemokine that recruits monocytic phagocytes and prolonged expression in chronic wounds causes an extended inflammatory response. Similarly, insulin-like growth factor binding protein-1 (IGFBP-1) binds anti-fibrotic insulin-like growth factor (IGF), preventing IGF

from activating its receptor. High levels of IGFBP-1 are associated with severe pulmonary and liver fibrosis [19,20].

In addition to fibrosis, a strong pro-inflammatory reaction is observed in mesh complications, including a powerful M1 response, which has been shown to influence fibrosis surrounding biomaterials; a localized but robust T cell response; and a predominance of pro-inflammatory cytokines [8,21–23]. This may contribute directly to mesh pain through macrophage-stimulated release of pro-inflammatory factors, such as tumor necrosis factor α (TNF- α) and interleukin 1- β (IL1- β), which stimulate nociceptor neurons, and through cytotoxic T-cells, result in neuropathic pain [24–26]. We observed that T_{reg} concentrations were inversely related to the amount of fibrosis in mesh-tissue complexes excised from women with complications, suggesting the immune system and normal wound repair mechanisms must synchronize to prevent complications after mesh implantation [10].

Despite these important laboratory findings, it is unknown whether histologic measures of fibrosis, the host immune response and/or levels of soluble biofactors observed in mesh-tissue explants removed for complications correlate with patient-reported measures such as pain. Pelvic pain attributed to mesh complications is subjective and may be influenced by factors other than the host response to the biomaterial such as pain syndromes or symptoms, expectations, fear or other motivation such as a plan to pursue litigation [27,28]. In addition, it is not known if persistent symptoms following removal is due to factors such as chronic neuroinflammation and scarring or if it is because the pain is multifactorial or not attributable to the mesh. The overall goal of this study was to define biologic factors involved in fibrogenesis following mesh implantation and correlate them with patient-reported pain outcomes and response to removal. We hypothesized that key regulators of wound healing, namely TGF- β 1, PDGF-BB, bFGF, MCP-1 and IGFBP-1, would be elevated in mesh complications compared to controls. In addition, we hypothesized that patients with greater concentrations of T_{regs} and lower levels of fibrosis would be more likely to have improvement in pain after mesh removal than those with a profibrotic profile, more CD8⁺ cells and fewer T regulatory cells.

2. Methods

2.1. Patient selection and study design

After approval by the Institutional Review Board at the University of Pittsburgh Medical Center (9/11/2017), patients who underwent vaginal mesh removal for the primary indications of pain or exposure were invited to participate in the Magee Mesh Biorepository. Exclusion criteria included active infection as determined by their surgeon, chronic immunosuppression or an autoimmune disorder. Patients were consented for collection of the excised mesh-tissue complexes and an optional adjacent full thickness vaginal biopsy at the time of mesh excision. Because the vaginal biopsy was considered optional for study enrollment, additional control biopsies were obtained from women undergoing surgery for pelvic organ prolapse or stress urinary incontinence without prior mesh placement. Demographic data including age, BMI, gravity, parity, smoking status, menopausal status, use and type of hormone replacement therapy, and comorbidities as well as variables

describing the initial mesh placement and removal, and complications such as tenderness and mesh deformation were collected at the time of enrollment using a standardized form developed *a priori*. Questionnaires on pain and symptoms including visual analog scale (VAS) pelvic pain scores and PFD-20 questionnaires were completed at enrollment and 6 months after removal (supplementary Fig. 1). Patients were considered responders (i.e. responded positively to their mesh removal) if they had at least 13 mm of improvement on the VAS scale, previously demonstrated to be the minimally important clinical difference [29]. They were considered nonresponders if they had < 13 mm of improvement or worsening of pelvic pain on the VAS scale.

30 total samples removed for pain in the absence of exposure ($n = 15$) or exposure with or without pain ($n = 15$) were randomly selected for analysis from samples in the Mesh Biorepository that had been removed for pain or exposure. When mesh-tissue complexes are obtained from women, the amount of tissue obtained is limited, especially when a portion of the specimen is needed for litigation. For analysis, specimens were divided for histology and biochemistry with the former prioritized. In order to maintain random sampling, specimens were of varied quantity which precluded our ability to perform all assays on all of the specimens. To overcome this limitation, the sample size was increased until the appropriate assays could be performed. Thus, of the 30 samples, 29 had tissue available for histology and 21 had sufficient tissue for biochemistry and one sample was added to the biochemistry group that did not have histology available (Fig. 1). The data presented in the current study represents a secondary analysis of collagen quantification, immune cell type characterization and TGF- β 1 quantification previously characterized in the 29 patients with tissue available for histology (the 30th patient was included only in the biochemical analysis) [10]. As part of the current analysis, these data were correlated with patient-centered outcomes. Additionally, concentrations of TGF- β 1, bFGF, MCP-1, PDGF-BB, and IGFBP-1 were determined in 21 specimens with tissue sufficient for biochemistry to further characterize the host response as shown in Fig. 1.

2.2. Histologic preparation

Mesh tissue complexes were collected in a sterile manner, placed on ice and then imbedded in O.C.T. compound (Tissue-Tek; Sakura Finetek USA Inc, Torrance, CA), flash frozen in liquid nitrogen, sectioned at 7 μ m and preserved at -80 °C.

2.3. Immunofluorescent labeling

As previously described [10], tissue sections were incubated with primary antibodies to CD4 (CD4 anti-goat 1:50, Santa Cruz) and FoxP3 (FoxP3 anti-mouse 1:150, Abcam) followed by secondary antibody incubation (Biotin anti-Goat and HRP anti-mouse at 1:500) and amplification with Alexa Flour 647 streptavidin and Alexa Flour 594 tyramide HRP conjugate. Aqueous mounting medium containing DAPI (Vectashield with DAPI, Vector Laboratories, Burlingame, CA) was applied. Separate slides were triple labeled for CD8⁺, FoxP3⁺, and DAPI to assess T_{reg}/cytotoxic T cell ratios. Whole tissue section images were taken and analyzed using a Nikon ECLIPSE 90i upright microscope and Nikon NIS Elements Imaging Software (Nikon USA, Melville, NY). Fiber areas were measured. Samples were compared to negative controls to ensure specific labeling and

were run in duplicate or triplicate. CD4⁺ and FoxP3⁺ cells per 0.25 mm² were obtained as concentrations and total CD4⁺ and CD8⁺ cell counts were obtained per tissue section and normalized by fiber area. T_{regs} per 100 CD8⁺ cells and T_{regs} per 100 CD8⁺ cells per cm² fiber area were calculated. To correlate the location of CD4⁺ and FoxP3⁺ cells to fibrotic response, we used a previously described customized quantification system [10], also described in Section 2.4 below. Representative IHC images can be found in Artsen et al, *Acta Biomater* 2019 [10].

2.4. Trichrome staining and fibrosis quantification

As previously described [10], a standardized Masson's trichrome staining protocol omitting hematoxylin cell nucleus staining was applied and samples imaged using the Nikon ECLIPSE 90i upright microscope. A digital grid composed of 0.25 mm² squares was laid over the trichrome images. In each grid, the intensity of collagen fibrosis was quantified using a digital intensity profile. Intensity in each grid was then averaged and the grids were classified as mild, moderate, and high or pathologic fibrosis, which was more than 2.5 standard deviations higher than controls (Fig. 2).

2.5. Cytokine quantification

21 of the 30 patients with mesh complications in this cohort had sufficient tissue available for biochemistry. The number of patients and autologous biopsies analyzed for each assay depended on the concentration of protein needed for each assay as detailed in Fig. 1. For TGF- β 1 analysis, 16 patients with mesh complications had sufficient tissue for analysis with optional *autologous control* biopsies available in 8 patients (Fig. 1). 14 *matched control* biopsies were matched based on age (< age 50, 50–75, > 75), BMI (by NIH obesity class) and presence or absence of prolapse (stage 2) or stress incontinence. Protein concentrations were determined using a DC protein assay (Bio-Rad, Hercules, CA). TGF- β 1 was quantified using a commercially available enzyme-linked immunosorbent assay (R&D Systems, Minneapolis, MN). Samples were run in duplicate with 50 μ g protein.

A commercially available custom Luminex (R&D, Minneapolis, MN) kit was then used to determine concentrations of MCP-1, PDGF-BB and IGFBP-1. These chemokines and growth factors had previously been shown to demonstrate a difference between mesh and controls in preliminary studies employing a Luminex platform of proinflammatory and wound healing markers. 17 patients with mesh complications had sufficient tissue for comparison to 5 available autologous control biopsies and 10 matched controls. Those without appropriate controls were excluded from the paired analysis. Samples were run in duplicate with 80 μ g protein.

The concentration of bFGF in the mesh-tissue complexes was found to be relatively high in preliminary studies. Therefore, bFGF analysis required a much smaller concentration of tissue, and bFGF was determined using a commercially available enzyme-linked immunosorbent assay kit (R&D) in 19 patients with biochemistry tissue, 6 autologous controls and 13 matched biopsies. Samples were run in duplicate with 15 μ g protein.

2.6. Pathway analysis

Pathway analysis was performed using Ingenuity Pathway Analysis Software (Qiagen, Hilden, Germany) to define interactions between bFGF, TGF- β 1 and PDGF-BB and references were reviewed by the primary author for relevance.

2.7. Statistical analysis

For cytokine analysis, sample size calculation suggested 11 mesh samples and 11 pooled autologous and matched controls were needed to detect an expected 45% difference at a significance level of < 0.05 and power of 80%. Sample size for histologic samples was determined by the sample size calculation for the primary analysis, namely that 14 samples were needed with a power of 80% and the standard deviation of 0.00014 seen in preliminary data to detect an expected difference in T cell populations in areas of high versus low fibrosis using a Mann–Whitney U test.

Mann–Whitney U tests and regression models were used to compare groups and determine correlation with pain score, fibrosis, or response to removal. Signed rank tests were used to compare cytokines near mesh complications with autologous and matched control biopsies. Hypothesis testing was considered significant at the $p < 0.05$ level. All statistical tests were completed in STATA software (version 15.1; StataCorp, College Station, Texas) or SPSS (IMB, Armonk, NY).

3. Results

3.1. Patient demographics

Women enrolled in the study were middle aged (median 56 years), overweight (29.0 kg/m²), and parous (median parity 3). Pain samples included 10 mid-urethral slings and 5 prolapse meshes. Exposure samples included 7 mid-urethral slings and 8 prolapse meshes. Median duration of implantation was 49.9 months. There were no differences in age, BMI, parity, duration of implantation, tobacco use, rate of diabetes, menopausal status, hormone use, or prior hysterectomy based on removal indication (Table 1, $p > 0.05$). As expected due to the study design, there were no differences in women with mesh complications and their matched controls.

3.2. Baseline pain scores correlated linearly with degree of fibrosis

Overall median baseline VAS score was 45 (IQR 24–76) and 78.6% of patients answered “yes” to “Do you have pelvic pain?” demonstrating that many women with exposure also have pain. Median bother was 4 (scale 1–4). Women with pain in the absence of exposure had higher baseline pelvic pain scores than women with mesh removed for exposure [73 (45–95) vs 30.5 (23–47), $p = 0.46$]. At baseline, VAS score was associated with presence of pain and bother on the PFD-20 questionnaire ($p < 0.002$) and VAS scores increased linearly with amount of fibrosis ($p = 0.013$). Baseline VAS scores were not associated with T_{reg} concentration or TGF- β 1, PDGF-BB, MCP-1, IGFBP-1 or bFGF concentrations.

3.3. T regulatory cell concentration predicted response to removal

Six month follow up data was available for 17 of 29 patients with histology data. Of these 17 patients, there were 9 responders with at least 13 mm improvement on the VAS pelvic pain score and 8 were nonresponders.

Responders were equally likely to have prolapse or sling mesh (5 patients vs 4 patients, $p > 0.05$) and equally likely to have pain or exposure complications (4 patients vs 5 patients, $p > 0.05$). The average age and BMI of responders and nonresponders were similar and reflected the overall demographics of the group. There was no difference in the amount of fibrosis between responders and nonresponders. However, there was a trend toward lower duration of implantation in responders (22.0 vs 55.5 months, $p = 0.17$).

Responders had more T_{regs} (7.8 vs 0.3 per mm^2 , $p = 0.036$) and a trend toward higher $T_{reg}/CD8$ ratios (2.7 vs 1.1 $T_{reg}/100$ CD8 cells, $p = 0.07$, Table 2). On logistic regression modeling, the cell density of T_{regs} predicted responder status. Patients were 1.6 times as likely to be a responder for every additional T_{reg}/mm^2 of tissue, ($p = 0.05$).

3.4. TGF- β 1 was doubled in nonresponders

Six month follow up data was available for 9 out of 16 patients with TGF- β 1 data. Nonresponders had nearly twice the level of TGF- β 1 [12.8 (12.3–13.2) vs 6.6 (6.1–9.0) pg/ μ g protein] compared to responders ($p = 0.032$). bFGF, PDGF-BB, MCP-1 and IGFBP-1 concentrations were similar between responders and nonresponders (Table 2).

3.5. bFGF was decreased in mesh tissue complexes compared to controls

The median concentration of bFGF was nearly 50% decreased in mesh tissue complexes vs all controls combined (Figs. 3A and 4A, $p = 0.0003$). This was statistically significant when mesh complication levels were compared to autologous controls 7.0 (6.5–7.8) vs 20.2 (15.4–26.4) pg/ μ g protein, $n = 6$, $p = 0.028$, and matched controls, [6.0 (4.4–7.6) vs 10.7 (9.7–12.0) pg/ μ g protein, $n = 13$, $p = 0.0046$]. bFGF was similar in pain and exposure groups and did not correlate with collagen content or baseline VAS.

3.6. PDGF-BB was higher in mesh tissue complexes than controls

PDGF-BB was increased more than 5 times in mesh tissue complexes at 0.5 (0.4–0.7) pg/ μ g protein compared to 0.08 (0.05–0.27) pg/ μ g protein in control biopsies ($n = 15$, Figs. 3B and 4B, $p = 0.002$). This was seen for both autologous control biopsies [0.4 (0.4–0.5) vs 0.05 (0.03–0.2), $n = 5$, $p = 0.0$] and matched biopsies [0.5 (0.3–0.7) vs 0.10 (0.7–0.3) $n = 10$, $p = 0.02$].

3.7. IGFBP-1 is increased in exposure complications but not pain complications

IGFBP-1 was statistically higher in tissue explanted for exposure at 24.6 (10.2–31.6) pg/ μ g protein compared to controls, 11.4 (8.9–14.8) pg/ μ g protein ($p = 0.018$) but not in mesh explanted for pain without exposure (Figs. 3D and 4D). MCP-1 and was similar between mesh-tissue complexes and controls ($p > 0.05$, Figs. 3C and 4C). There was no difference between pain and exposure groups and these analytes did not correlate with collagen content or baseline VAS.

3.8. TGF- β 1 is regulated by bFGF on Pathway Analysis

Pathway analysis revealed direct regulation of TGF- β 1 by bFGF and indirect regulation via PDGF-BB (Fig. 5). bFGF demonstrated inhibition of TGF- β 1 dependent differentiation of fibroblasts into myofibroblasts [17]. bFGF's dose-dependent inhibition of PDGF-BB was demonstrated in pharmacologic binding studies and PDGF-BB was seen to upregulate TGF- β 1 [30,31].

4. Discussion

Polypropylene mesh complications have limited the use of transvaginal mesh for prolapse and stress urinary incontinence despite improvement in long-term success of these repairs over suture alone. A strong pro-inflammatory response has been demonstrated in mesh complications, but data increasingly also suggests a pro-fibrotic mechanism of pain resulting from poor tissue integration into mesh [8,10,22]. The results of this study support this mechanism and our hypothesis with the following important findings: 1) patient-reported outcomes, specifically, pain scores, were positively correlated with degree of fibrosis; 2) increasing number of T_{regs}/mm² predicted likelihood of improvement in pain after mesh removal; and 3) TGF- β 1, a pro-fibrotic cytokine, was doubled in nonresponders. Additionally, decreased bFGF and increased PDGF-BB were shown in a pathway analysis to be a possible mechanism for the upregulation of TGF- β 1.

4.1. Patient reported pain scores were correlated with degree of fibrosis

The difficulty of studying pelvic pain is well documented, particularly because the differential is broad and the degree of over-lap between pelvic pain symptoms is high [27,32]. However, many clinicians have documented a tender, painful band of mesh on examination [33] resulting from pore collapse, mesh contraction and bridging fibrosis seen in biomechanical and histologic analyses [11,12,34,35]. Our findings are in line with these studies by providing one of the first empiric data linking patient-perceived pain to histologic fibrosis in mesh pain complications.

4.2. Increasing T_{reg} concentration predicted likelihood of improvement in pain after mesh removal

Improvement in pain and dyspareunia after mesh removal is reported to be as low as 57%, with only the presence of other chronic pain syndromes predicting a lower likelihood of improvement [6,28]. Here we saw a similar percentage responding to removal at 53% with similar improvements regardless of complication type and mesh type. Notably, increasing T_{reg} concentrations predicted a positive response to mesh removal, regardless of fibrosis severity. We have previously shown that T_{regs} are inversely related to fibrosis in mesh-tissue explants and T_{regs} are well known for decreasing inflammation through suppressing T helper cells [36]. Our findings suggest that even in the case of severe fibrosis, the adaptive immune response may be responsible for prevention of complication and recovery through reducing fibrosis and inflammation; and that the altered wound healing that occurs in mesh complications could be modified by altering the adaptive immune response. In addition, pelvic pain appeared more likely to improve when mesh removal was performed after a

shorter duration of implantation. This observation deserves further study, as it could be that early referral to a specialist to evaluate for removal may improve outcomes.

4.3. TGF- β 1 was doubled in nonresponders and a pro-fibrotic growth factor profile seen in mesh-tissue explants may explain increased TGF- β 1

We have previously shown that TGF- β 1, which has been shown to facilitate myofibroblast contraction of the collagen matrix, is increased in women with mesh complications and here we demonstrate high levels in women who don't respond to mesh removal [10,37]. The importance of TGF- β 1 has been documented in the foreign body response [38] and our results further support the role of TGF- β 1 and fibrosis in mesh complications and ongoing pain after mesh removal.

Decreased bFGF in mesh-tissue explants and increased PDGF-BB may account for the increased TGF- β 1 observed in mesh tissue complexes from women with complications (Fig. 5). Our pathway analysis revealed that bFGF interacts directly and indirectly with TGF- β 1 through PDGF-BB. bFGF is produced by fibroblasts and facilitates reepithelization and tissue remodeling and can inhibit TGF- β 1 dependent differentiation of fibroblasts into myofibroblasts [18,33]. Normally, myofibroblasts undergo apoptosis with resolution of wound healing, but their persistence is a key feature in fibrosis [39]. Thus, bFGF may become depleted in counteracting TGF- β 1 in this state of dysregulated wound healing.

PDGF-BB is released by platelets after injury and stimulates macrophages to produce TGF- β 1, increases the proliferation and activation of fibroblasts, thus increasing ECM production and contraction [18,30]. The increase in PDGF-BB is supportive of repetitive cycles of injury [18] and repair which have been suggested as the stiffer polypropylene mesh moves against the softer vagina in a so called "brillo pad effect". bFGF and PDGF-BB participate in reciprocal inhibitory feedback [40,41]. This negative feedback may result in an unchecked pro-fibrotic state, with elevated PDGF-BB directly contributing to decreased bFGF and resulting in increased TGF- β 1 levels. Alternatively, as bFGF and PDGF-BB are both critical in neovascularization in wound healing, lower bFGF and higher PDGF-BB may also indicate dysregulated angiogenesis in mesh complications [42].

Finally, IGFBP-1 was also elevated in mesh explants for exposure but not for pain. This may suggest worsened dysregulated wound healing in exposure complications, with a pro-fibrotic, low T regulatory state preventing appropriate mesh integration and wound healing.

4.4. Strengths, limitations and future directions

This is a multidimensional study correlating patient-reported pain with histologic, immunologic and cytokine analysis of polypropylene mesh complications. By using validated pain scores, we were able to link objective observations to meaningful patient reported outcomes. Immunologic assessment included a ratio of T_{regs} to CD8⁺ effector cells, allowing for a comprehensive look at adaptive immune activation and tolerance. In addition, we supported histologic analysis with assessment of profibrotic cytokines and provided a possible mechanism for our TGF- β 1 findings using pathway analysis.

Limitations include lack of a true control group, as mesh is rarely removed from women without complications. The use of matched biopsies here from women with prolapse or stress urinary incontinence provides a control group that most closely mimics women who would have mesh placed. In addition, the identification of responders and nonresponders provides a clinically meaningful comparison. Animal models also provide supporting control data, and stronger pro-inflammatory responses are seen in animals implanted with stiffer mesh materials with smaller pores, consistent with pro-inflammatory and pro-fibrotic responses seen here [21]. Although meshes removed here could not be easily grouped by biomechanical properties due to the variety of mesh removed, categorizing into sling and prolapse mesh takes into account their different mesh burden and stiffness. This is a problem inherent in human studies as so many different mesh types have been on the market. Another limitation is the and the overall small sample size and the small amount of tissue for some samples which limited the numbers for each analysis. Luminex was used in this study to help overcome this, but due to different concentration and bead region requirements, TGF- β 1 and bFGF could not be performed concurrently with the other analytes through the Luminex platform. However, even with small numbers, we were able to see meaningful differences due to large effect sizes. Further studies into dysregulated wound healing in mesh complications will help confirm these findings. Finally, it is possible that patients may falsely elevate pain scores in order to receive mesh removal or for litigation purposes. However, the VAS score has been widely validated and additional measures to assess pain such as the validated PFDI-20 questions on the presence of pelvic pain and bother were reassuringly consistent.

Future studies should explore the role of myofibroblasts, which may be responsible for pathologic collagen deposition, as well as nerve fiber disruption which may contribute to response to removal. This study also suggests a potential therapeutic role of T cells which are in early clinical trials for other diseases [38,43]. These findings should inform future studies of new biomaterials and provide a potential for therapeutic manipulation of the immune response after implantation. Further studies are needed to determine if nonresponders have low T_{reg} levels prior to mesh implantation. Indeed, phenotyping nonresponders is an important pathway moving forward so that these women can be identified ahead of time and counseled and treated appropriately.

4.5. Conclusions

Fibrosis was correlated with patient-reported pain, and higher T_{reg} concentrations and lower levels of TGF- β 1 were associated with improvement after mesh removal. Increased PDGF-BB and decreased bFGF may drive an increase in TGF- β 1 seen in mesh complications. This improved understanding of the mechanism by which mesh can lead to pain moves us closer to the ultimate goal of preventing mesh complications.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Statement of Significance

Polypropylene mesh improves anatomical outcomes in urogynecologic procedures, but is associated with complications, including pain and exposure through the vaginal epithelium. Mesh-induced pain is difficult to treat, and it is unclear why only half of women experience pain improvement after mesh removal. In this study, patient pain correlated with the presence of fibrosis and women with more T regulatory cells and lower TGF- β 1 were more likely to have pain improvement following mesh removal. These findings implicate fibrosis as a mechanism of pain complications and suggest that the adaptive immune response may be responsible for prevention of complication and recovery. This improved understanding of how mesh can lead to pain moves us closer to the ultimate goal of preventing mesh complications.

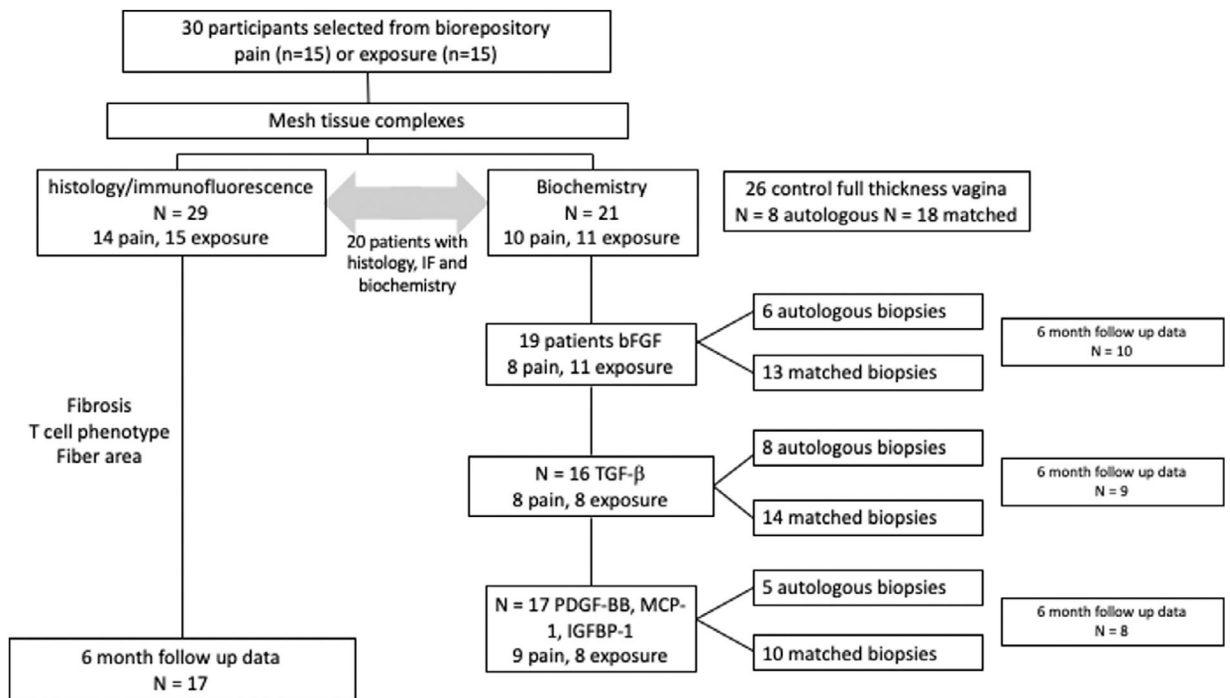


Fig. 1. Flow chart demonstrating subgroups of patients in each analysis. Due to limited tissue for each sample and variable concentrations required for each assay, not every assay could be performed on each patient in this cohort.

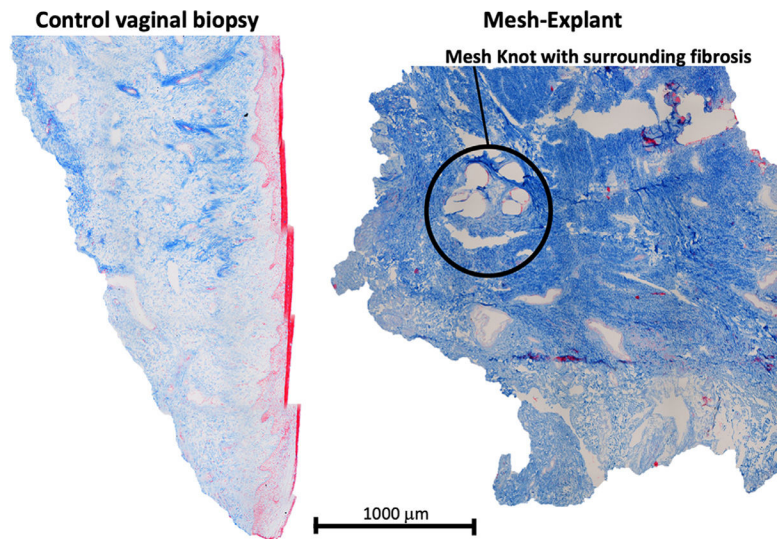


Fig. 2. Representative trichrome images of vaginal control biopsy and mesh-explant. Pathologic fibrosis was more than 2.5 standard deviations higher than controls.

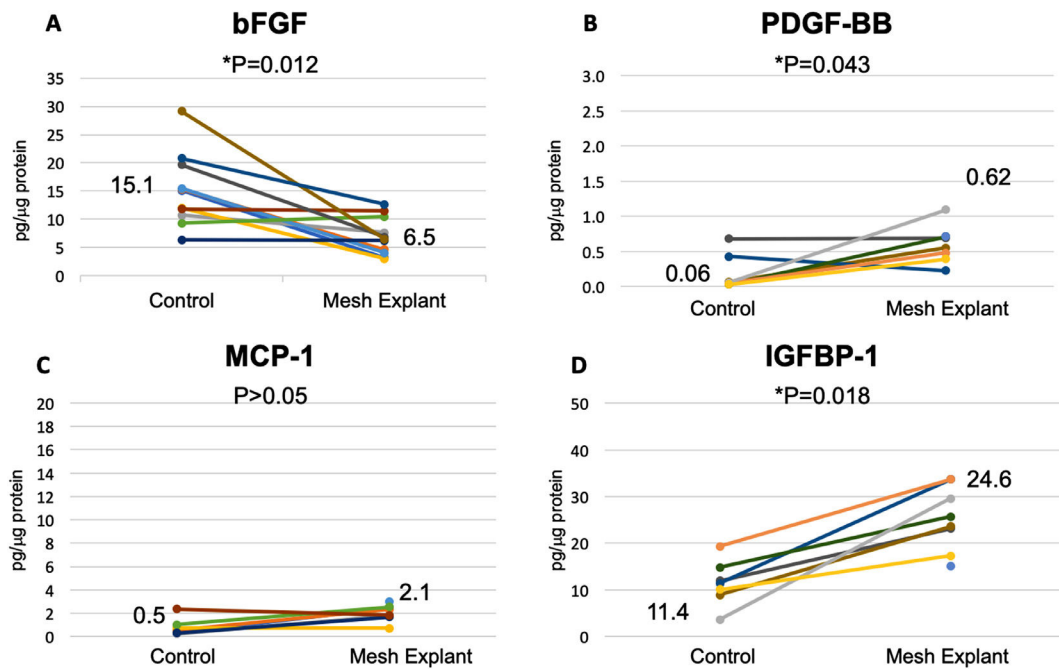


Fig. 3. Profibrotic cytokines in mesh-tissue complexes removed for exposure compared to vaginal control biopsies. Results are given in pg/μg protein and include both autologous and matched controls. Median values are shown. *P* values represent Wilcoxon signed ranks given the wide variation.

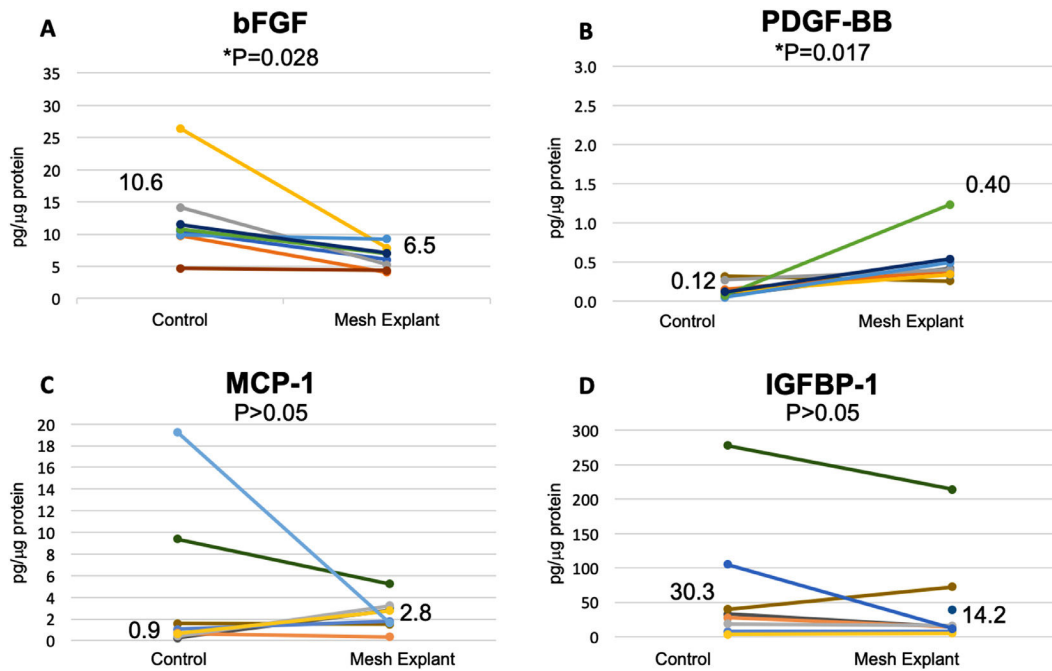


Fig. 4. Profibrotic cytokines in mesh-tissue complexes removed for pain compared to vaginal control biopsies. Results are given in pg/μg protein and include both autologous and matched controls. Median values are shown. *P* values represent Wilcoxon signed ranks given the wide variation.

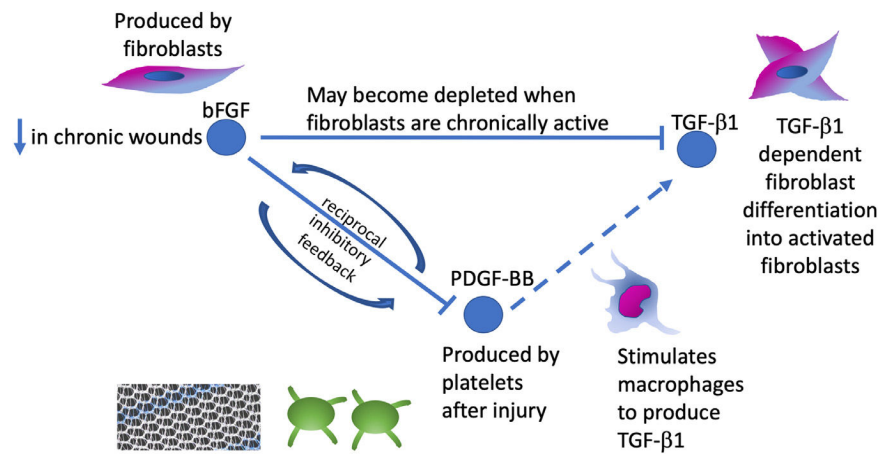


Fig. 5. Pathway analysis of bFGF with factors TGF- β 1 and PDGF-BB, which we have shown to be increased in urogynecologic mesh complications. Pathway analysis revealed bFGF regulation of TGF- β 1 directly and via PDGF-BB.

Table 1

Demographics of women with mesh complications by exposure or pain in the absence of exposure.

	Pain (n = 15)	Exposure (n = 15)	P value
Age (y)	56 (42–59)	56 (54–68)	0.23
BMI (kg/m ²)	28.0 (25.1–32.2)	30.0 (24.7–32.4)	0.80
Median parity (range)	3 (2–5)	3 (1–5)	0.90
Midurethral sling	10 (66.7%)	7 (46.7%)	0.46
Prolapse mesh	5 (33.3%)	8 (53.3%)	0.46
Duration implantation (mo)	36.6 (15–56)	58.5 (22–73)	0.57
Postmenopausal	11 (73.3%)	13 (86.7%)	0.65
Hormone therapy use*	8 (53.3%)	8 (53.3%)	0.99
Prior hysterectomy	11 (73.3%)	9 (60.0%)	0.70
Current or smoking history [^]	8 (53.3%)	8 (53.3%)	0.99
Diabetes	3 (20.0%)	1 (6.7%)	0.60
Chronic steroid use	0 (0%)	1 (6.7%)	0.99
Baseline VAS Pelvic Pain	30.5 (23–47)	73 (45–95)	0.046
TGF- β 1 (pg/ μ g protein) ^o	11.8 (8.2–14.4)	9.1 (6.9–13.3)	0.83
bFGF (pg/ μ g protein) ⁺	6.5 (4.6–7.6)	6.5 (4.0–10.4)	0.99
PDGF-BB (pg/ μ g protein) [‡]	0.4 (0.3–0.5)	0.6 (0.4–0.7)	0.18
MCP-1 (pg/ μ g protein) [‡]	2.8 (1.6–3.0)	2.1 (1.7–2.5)	0.63
IGFBP-1 (pg/ μ g protein) [‡]	14.2 (11.9–38.6)	24.6 (20.2–31.6)	0.29

Median (interquartile range) is listed for continuous variables, frequencies (%) for categorical variables. *P* values represent Mann–Whitney *U* for continuous variables and or Fishers Exact for categorical variables. VAS indicates visual analog scale.

* includes vaginal estrogen.

[^] smoking information missing on one patient.

^o TGF- β 1 samples include 8 patients with pain and 8 with exposure.

⁺ FGF analysis included 7 patients with pain and 11 with exposure.

[‡] Luminex analysis included 9 patients with pain and 8 with exposure complications (see Fig. 1).

Table 2

Characteristics of Responders and Non-responders to Mesh Removal.

	Responders (n = 9)	Non-responders (n = 8)	P value
Age (y)	57 (52.5–65.0)	55.5 (48.8–68.0)	0.82
BMI (kg/m ²)	28.2 (26.2–31.6)	29.6 (23.5–32.4)	0.96
Duration implantation (months)	22.0 (10.5–61.3)	55.5(35.0–58.9)	0.17
Baseline VAS Pelvic Pain	45.0 (33–76)	45.5(16–84.5)	0.53
Fibrosis level (intensity)	54.6 (38.5–72.1)	66.2 (35.8–82.2)	0.89
Treg concentration (cells/mm ²)	7.8 (1.3–10.2)	0.3 (0.5–3.4)	0.036
CD8 (cells/sample)	562.0 (289.5–829.0)	291.0 (100.8–544.8)	0.17
Treg/CD8 ratio (Treg /100 CD8)	2.7 (1.1–7.3)	1.1 (0.1–1.8)	0.07
TGF- β 1 (pg/ μ g protein)*	6.6 (5.7–10.2)	12.8 (10.0–13.3)	0.032
bFGF (pg/ μ g protein) *	5.0 (3.81–8.37)	7.3 (6.03–10.43)	0.29
PDGF-BB (pg/ μ g protein)*	0.3 (0.3–0.6)	0.4 (0.4–0.7)	0.46
MCP-1 (pg/ μ g protein) *	2.3 (1.5–3.0)	2.5 (1.8–2.8)	0.88
IGFBP-1 (pg/ μ g protein)*	38.6 (23.6–72.3)	17.3 (14.2–23.1)	0.05

Data is presented as median (interquartile range). P values represent Mann–Whitney U tests.

* Tissue for TGF- β 1 quantification was available for 4 responders and 5 non-responders; for FGF 4 responders and 6 non-responders, for PDGF-BB, MCP-1 and IGFBP-1 included 3 responders and 5 non-responders.