

**ULTRAFILTRATION DYSFUNCTION IN PERITONEAL DIALYSIS: THE ROLE OF
INFLAMMATION, FIBROSIS, AND ANGIOGENESIS.**

By

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ABSTRACT

Background: Peritoneal dialysis (PD) is a valuable therapy for end-stage renal disease. Changes in the peritoneal transport properties, including increased solute transport and ultrafiltration (UF) dysfunction are main clinical limitations of PD. The capillary wall of peritoneal blood vessels plays a key role as a barrier to solute transport. Therefore, neovascularization of the peritoneal tissues is a main determinant in the functional changes in the peritoneal membrane. **I hypothesize that ultrafiltration dysfunction is caused by increased peritoneal vascularization with a concomitant increase in glucose transport from the peritoneal cavity and rapid loss of the ultrafiltration gradient. Further, I hypothesize that this increased vascular surface area is induced by profibrotic and inflammatory cytokines such as transforming growth factor (TGF) β , interleukin (IL) 1β , and tumour necrosis factor (TNF) α through upregulation of angiogenic cytokines such as vascular endothelial growth factor (VEGF).**

Methods: I used adenovirus mediated gene transfer of cytokines such as TGF β 1, IL-1 β , and TNF α to the peritoneal of rats and studied the effects on histology, angiogenesis, gene regulation, solute transport, and UF. I also used an animal model of daily peritoneal exposure to dialysate solution. The angiogenic and fibrogenic response of the peritoneum to this model was detailed, along with the effect of overexpression of angiostatin and decorin by adenovirus mediated gene transfer.

Results: Adenovirus mediated gene transfer of TGF β 1 led to increased peritoneal fibrosis and angiogenesis which persisted to 28 days. These histologic changes were associated with increased solute transport of glucose and decreased UF. In vitro and in

vivo, TGF β 1 appeared to up regulate expression of VEGF. Inflammatory cytokines such as TNF α and IL-1 β both induced fibrosis, angiogenesis, and peritoneal membrane dysfunction. The kinetics of the response was quite different. TNF α induced a very transient response with a complete resolution of changes by 21 days after adenovirus infection. The response induced by IL-1 β was more sustained. I hypothesize that the strong expression of tissue inhibitor of metalloproteinase after IL-1 β treatment may explain the prolonged fibrogenic response after IL-1 β . Daily exposure to dialysate also induced a strong fibrogenic and angiogenic response in the peritoneum. The anti-angiogenic agent angiostatin, when overexpressed using adenovirus mediated gene transfer, reduced the vascularization and improved the ultrafiltration dysfunction. Decorin, a proteoglycan that binds and inactivates TGF β , reduced peritoneal fibrosis, but did not alter the angiogenic response, nor did it improve peritoneal membrane function.

Discussion: These experiments have clarified the role of angiogenesis in UF dysfunction. The correlation between blood vessel density and ultrafiltration, along with the improvement in UF after treatment with angiostatin, is compelling evidence that vascularization of the peritoneal membrane causes UF dysfunction. There is a close association between fibrosis and angiogenesis in the peritoneum. Fibrosis appears to be necessary for a prolonged angiogenic response. Further work is required to identify the factors leading to the interaction between fibrosis and angiogenesis. The role of the interstitium and lymphatics in solute transport and UF dysfunction needs to be better defined. Finally, anti-angiogenic therapies need to be studied in these models of peritoneal membrane dysfunction and eventually applied to patients on PD to improve the quality and duration of this therapy.

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LIST OF ABBREVIATIONS

AGE	- advanced glycation endproducts
ALK	- activin receptor-like kinase
ANG	- angiopoietin
CTGF	- connective tissue growth factor
EPS	- encapsulating sclerosing peritonitis
ESRD	- end stage renal disease
FGF	- fibroblast growth factor
GDP	- glucose degradation products
HIF	- hypoxia inducible factor
IL	- interleukin
LPS	- lipopolysaccharide
MMP	- matrix metalloproteinase
NOS	- nitric oxide synthase
PAI	- plasminogen activator inhibitor
PD	- peritoneal dialysis
PDGF	- platelet derived growth factor
RRD	- residual renal function
TGF β	- transforming growth factor β
TGF β R	- transforming growth factor β receptor
TIMP	- tissue inhibitor of metalloproteinase
TNF	- tumor necrosis factor
UF	- ultrafiltration
VEGF	- vascular endothelial growth factor
VSMC	- vascular smooth muscle cell

TABLE OF CONTENTS

TITLE PAGE -----	ii
ABSTRACT -----	iii
ACKNOWLEDGEMENTS -----	v
LIST OF ABBREVIATIONS -----	vi
TABLE OF CONTENTS -----	vii
CHAPTER 1: INTRODUCTION	
Epidemiology -----	1
Outcomes -----	2
Encapsulating Peritoneal Sclerosis -----	5
Pathophysiology -----	5
Increased Vascular Surface Area -----	6
Aquaporin Dysfunction -----	7
Lymphatic and Tissue Reabsorption -----	8
Etiology -----	8
Bioincompatible Fluids -----	8
Peritonitis -----	10
Uremia and Systemic Inflammation -----	13
Treatment of Ultrafiltration Dysfunction -----	14
Icodextrin -----	14
Biocompatible Solutions -----	15
Alteration of the Interstitial Compartment -----	16
Anti-angiogenic Therapy -----	16
Peritoneal Fibrosis and the Role of TGF β -----	18
Angiogenesis in Fibrotic Peritoneal Tissue -----	20
Animal Models of PD -----	24
Summary -----	26
CHAPTER 2: GENE TRANSFER OF TRANSFORMING GROWTH FACTOR β TO THE RAT PERITONEUM: EFFECTS ON MEMBRANE FUNCTION -----	28
CHAPTER 3: INFLAMMATORY CYTOKINES, ANGIOGENESIS, AND FIBROSIS IN THE RAT PERITONEUM -----	41
CHAPTER 4: A CHRONIC INFLAMMATORY INFUSION MODEL OF PERITONEAL DIALYSIS IN RATS -----	53
CHAPTER 5: ANTIANGIOGENIC AND ANTIFIBROTIC GENE THERAPY IN A CHRONIC INFUSION MODEL OF PERITONEAL DIALYSIS IN RATS -----	60
CHAPTER 6: DISCUSSION -----	69
REFERENCES -----	76

CHAPTER 1: INTRODUCTION

Peritoneal dialysis (PD) is a valuable therapy for end-stage renal disease (ESRD). It is used by over 100,000 patients world wide which accounts for approximately 15% of the dialysis population ¹. A lack of randomized clinical trials comparing dialysis modalities has meant that the relative value of PD is still uncertain and contested. The clinical experience with PD suggests that it is a useful early therapy in ESRD and is a necessary third modality to complement hemodialysis and transplantation.

Along with peritonitis and inadequate removal of uremic toxins, ultrafiltration (UF) dysfunction is a main clinical limitation of PD. Ultrafiltration refers to the net fluid removal across the peritoneal membrane. Over time, PD patients develop an increased transport of small molecular weight solutes across the peritoneal membrane with a concomitant decrease in ultrafiltration ². Solute, such as urea, creatinine, phosphate, sodium, and potassium, travel from the blood to the peritoneal dialysis fluid. On the other hand, glucose travels from the dialysate solution to the blood. The capillary wall of peritoneal blood vessels plays a key role as a barrier to solute transport ³. Therefore, neovascularization of the peritoneal tissues is a main determinant in the functional changes in the peritoneal membrane. This increased vascular surface area leads to an increase in glucose transport from the peritoneal cavity. Glucose is the main osmotic agent used in PD, and the rapid loss of glucose leads to a rapid loss of the UF gradient. The role of increased fluid absorption from the peritoneum by lymphatics and tissue ⁴, along with dysfunction of aquaporins in the peritoneal vessel walls ⁵, is less well defined.

Epidemiology

In various studies, the prevalence of UF dysfunction leading to technique failure has been reported between 1.7% ⁶ and 13.7% ⁷. These data are confounded by the difficulty in accurately diagnosing the cause of technique failure. Using a clinical definition of UF dysfunction, prevalence rates have been reported between 15.3% ⁸ to 30.9% ⁹. Ho-dac-Pannekeet and colleagues used the recommended 4 hour high osmolarity PET and found the prevalence of UF dysfunction to be 23% in a group of 68 stable PD patients ¹⁰.

Changes in the peritoneal membrane function over time have been documented in several studies and recently reviewed ¹¹. The majority of longitudinal studies of peritoneal membrane function find an increase in small molecular weight solute transport with time on dialysis and a coincident decrease in ultrafiltration ^{2;12}. The onset of UF dysfunction appears between 2 ¹³ and 4 ² years after the initiation of PD. The inverse correlation between UF and solute transport has been clearly demonstrated in these studies. Lo and colleagues demonstrated a centripetal effect of membrane function over time ¹⁴. Among 55 PD patients followed for a mean of 22 months, there was an increase in solute transport in patients with initial low transport peritoneal membranes, and a decrease in solute transport in high transporters over time. One possible explanation is a selective censoring by technique failure or death of high transport patients while the low transport group experience a slow increase in solute transport over time ¹¹. This may also explain to some extent the apparent stability of peritoneal membrane function over the first 2–4 years of PD.

Outcomes

Several features at the initiation of PD predict subsequent survival. These features include high transport status peritoneal membrane¹⁵⁻¹⁸, low serum albumin^{15;19}, age^{15;20}, comorbid disease (including diabetes)^{21;22}, and residual renal function (RRF)^{19;23}. High transport status at the initiation of dialysis has also been shown to be associated with decreased UF function and poor outcome¹⁵. Acquired UF dysfunction is associated with decreased survival in several studies. Wang and colleagues measured the peritoneal membrane function in a cross section of 46 established PD patients and found increased solute transport to be inversely associated with fluid removal and survival¹⁷. In a study of 125 incident PD patients, Ates and colleagues demonstrated the impact of total fluid removal and sodium removal (including peritoneal and renal) on survival²³. From a large longitudinal database, Davies and colleagues compared survivors and non-survivors and demonstrated that non-survivors had a progressive increase in small solute transport with time on dialysis¹⁶.

UF dysfunction is also associated with other poor prognostic indicators such as clinical volume expansion²⁴ and diastolic hypertension²⁵. Volume expansion in PD patients is also associated with hypoalbuminemia²⁶ and adjusting dialysis prescription to increase UF can reduce volume expansion and increase serum albumin²⁷.

In summary, there is evidence that both intrinsic and acquired UF dysfunction is associated with decreased survival in PD patients. UF dysfunction is associated with volume expansion and this may be associated, through hypertension, left ventricular hypertrophy, pre-existing co-morbid disease, systemic inflammation, and

hypoalbuminemia, with increased mortality and morbidity, primarily through cardiovascular outcomes (Figure 1).

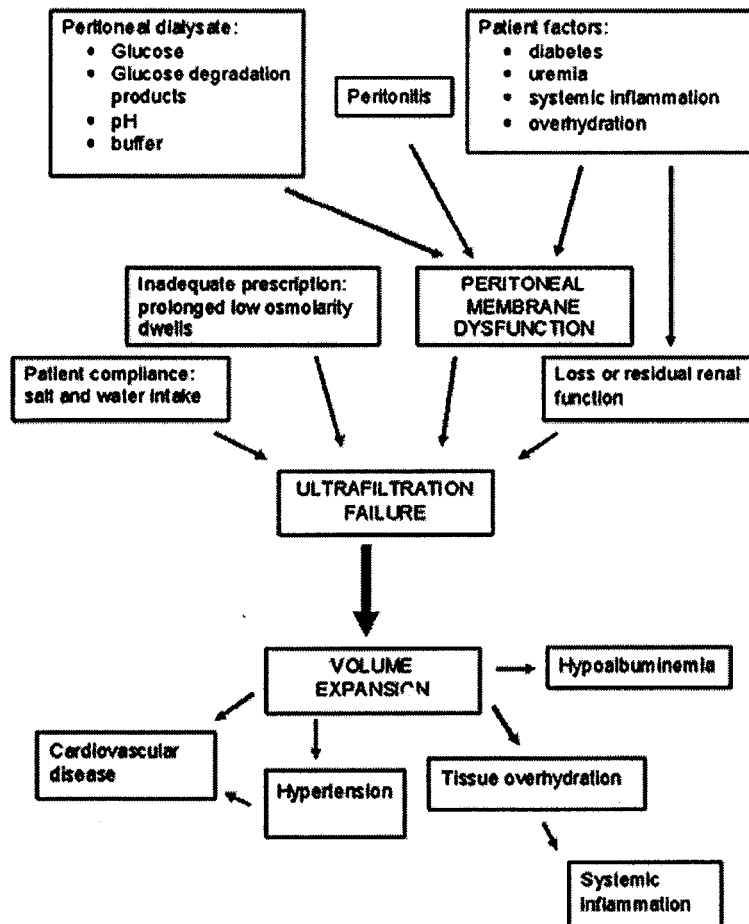


Figure 1. Components of peritoneal membrane dysfunction, ultrafiltration dysfunction, and outcomes of ultrafiltration failure

Encapsulating Peritoneal Sclerosis

In the peritoneum, progressive fibrosis alone can lead to a serious disorder known as encapsulating peritoneal sclerosis (EPS). This rare, poorly understood disorder occurs in less than 1% of PD patients²⁸. Patients present with hemorrhagic peritoneal effluent, abdominal pain, and intermittent bowel obstruction²⁹. Risk factors for this condition include severe unresolving peritonitis, chlorhexidine disinfectant, acetate buffered dialysis fluid, and certain medications³⁰. The histologic changes observed in EPS includes a loss of mesothelial cells, a thick, acellular, collagen rind which encloses the loops of small bowel with severe intra-abdominal adhesions³⁰. The involvement of inflammatory cells in this process has been controversial³¹. Treatment includes surgical debridement, cessation of PD, and immunosuppression with corticosteroids and other agents. Despite treatment, the outcome from this rare condition is generally very poor²⁹.

Pathophysiology

Ultrafiltration across the peritoneal membrane is the net effect of the osmotic force pulling fluid into the peritoneal cavity, and hydrostatic forces pushing fluid into the interstitium and into lymphatic reabsorption³². The 'membrane' across which the osmotic force acts has been described as a theoretical 3 pore model³³. Ultrasmall pores correspond to aquaporin channels located on peritoneal venules and capillaries⁵ and usually supply 40% of the net ultrafiltration³⁴. Small sized pores are the major conduit for solute and a major source of fluid ultrafiltration and large pores are few in number and allow for macromolecular transport. The major determinant of the prevalence of the small pores is the vascular surface area. This explains why increased vascularization of

the peritoneal tissues will lead to rapid glucose transport, loss of osmotic gradient, and loss of ultrafiltration³.

The peritoneal 'membrane' is comprised of an outer mesothelial cell layer covering a basement membrane which in turn overlays a thin interstitial layer comprised of connective tissue and blood vessels. Several biopsy studies have reported on the histologic changes induced by peritoneal dialysis. In the largest study reported to date, Williams and colleagues analyzed biopsy samples from 212 patients including normal, uremic, and peritoneal dialysis patients³⁵. They identified an increase in the thickness of the submesothelial tissue with uremia and further increase with peritoneal dialysis. They also identified a vascular sclerosis which has been recognized in other biopsy studies^{36;37} and correlated with time on peritoneal dialysis. When they analyzed the number of blood vessels in the submesothelial tissue, they did not find a correlation with time on dialysis, but did recognize a significant increase in vascularization in patients whose biopsy was performed at catheter removal due to 'membrane failure'. Several smaller studies have noted an increased vascularization of the submesothelial tissue in patients on peritoneal dialysis³⁷⁻³⁹. These studies have also noted a correlation between vascularization and interstitial fibrosis³⁸.

Increased vascular surface area

We hypothesize that increased vascularization of the peritoneal tissue is the primary mechanism of ultrafiltration failure. Between 50 and 75% of patients with UF dysfunction will have increased vascular area measured by increased small solute transport on a standard PET^{9;10}. In animal models of peritoneal dialysate exposure, we

have seen a similar increase in submesothelial vascularization which correlates with glucose absorption and correlates inversely with net UF⁴⁰. Furthermore, we were able to therapeutically reduce peritoneal vasculature using anti-angiogenic therapy, and this led to an improvement in UF function. Human biopsy studies confirm the association between submesothelial vascularization and UF dysfunction^{35,37-39}. Also, careful studies of the peritoneal effluent have revealed an association between locally produced vascular endothelial growth factor (VEGF) and peritoneal solute transport⁴¹.

Aquaporin dysfunction

Aquaporin dysfunction is a presumed mechanism leading to UF failure; however, the evidence is still somewhat contradictory. Monquill and colleagues identified patients with UF dysfunction but normal solute transport and low reabsorption rate. They concluded that impaired transcellular water transport (aquaporin dysfunction) was the mechanism for UF failure. Hyperosmolar PD solutions (3.86% glucose) did not improve ultrafiltration, and 5 of 6 patients required transfer to hemodialysis⁴². Aquaporin dysfunction as a cause of UF failure is also supported by animal studies of aquaporin 1 knock out mice⁴³. These mice displayed significantly impaired water transport using a modified peritoneal function test.

The ability to accurately measure aquaporin function using a standard peritoneal function test using hyperosmolar solution (3.86% glucose) has been questioned based on computer simulation³⁴. Despite the application of a correction for solute transport⁴⁴, low UF or high solute transport alone may blunt the effect of sodium dilution which is taken to be indicative of aquaporin dysfunction. Careful immunohistochemical analysis of

peritoneal tissue has led to the observation that aquaporin expression does not change in the face of acute peritonitis in an animal model ⁴⁵. Finally, normal aquaporin expression was found in a patient with presumed aquaporin dysfunction ⁴⁶. Alteration of the function of the aquaporin molecule could occur, but more work needs to be carried out in this area.

Lymphatic and tissue reabsorption

The importance of fluid reabsorption from the peritoneal cavity as a mechanism of ultrafiltration failure has been documented ^{9;47;48}. The interstitium of the peritoneal tissue consists mainly of collagen fibers and glycosaminoglycans such as hyaluronan. It is likely that alterations in the constitution of the interstitium will lead to alteration in hydrodynamic properties and macromolecular transport. Hydration will tend to expand the extracellular matrix and increase fluid conductance ⁴⁸. Hyaluronan may act to restrict fluid transport ⁴⁸ and this may have therapeutic importance ⁴⁹. Risk factors for increased tissue and lymphatic reabsorption have not been identified, but recently, Fussholler and colleagues identified an association between increased fluid reabsorption and time on dialysis ⁵⁰.

Etiology

Bioincompatible fluids

The observation that UF dysfunction increases with time on PD ¹² has led to the conclusion that elements of the PD fluid are bioincompatible and damage the peritoneal membrane. Several studies have documented progressive UF dysfunction in patients in

the absence of peritonitis ^{2;51}. Animal models have confirmed the detrimental effect of peritoneal dialysate on the peritoneum ^{52;53}. Putative bioincompatible components of standard PD fluid include glucose, glucose degradation products (GDP), lactate buffer, or acidic pH. Observational ⁵⁴ and animal studies ⁵⁵ have attempted to better define suspect dialysis components.

Prolonged exposure to high concentrations of glucose in the peritoneal dialysate has long been suspected to damage the peritoneal membrane. Davies and colleagues have shown that increased exposure to hyperosmolar glucose dialysate predated by 2 years the onset of UF dysfunction ⁵⁴. In this case control study, patients with increasingly permeable peritoneal membranes also had an early loss of RRF which probably necessitated the increasing use of glucose to replace renal ultrafiltration. Selgas and colleagues identified a similar effect with increased glucose exposure and diabetes both predicting for early UF dysfunction ⁵⁶. In a mechanistic study, Zweers and colleagues noted a decrease in locally produced VEGF in patients switched to non-glucose PD solutions ⁵⁷.

GDPs, such as methylglyoxal, glyoxal, and 3-deoxyglucosone, are reactive carbonyl compounds produced in peritoneal dialysate during the process of heat sterilization ⁵⁸. They have been shown to be more potent inducers of advanced glycation end-products (AGE) than glucose alone ⁵⁹. GDPs have been shown to alter mesothelial cell function and proliferation ⁶⁰ and to induce VEGF production in vitro ⁶¹. In animal models, conventional dialysate fluid induced vasodilation and capillary recruitment in the peritoneal tissues, whereas GDP-reduced PD fluid did not induce these changes ^{55;62}.

AGEs are irreversibly altered proteins (eg. pentosidine, carboxymethyllysine) which have a wide range of vascular and non-vascular⁶³ activity in the peritoneal tissue, signaling through a specific AGE receptor⁶⁴. Several studies have identified the appearance of AGE products such as pentosidine in the peritoneal effluent of patients on dialysis^{65;66}. These are felt to be both transported from the blood and produced and cleared in the peritoneum. Effluent pentosidine levels also correlate with time on dialysis and the use of non-glucose dialysate appears to reduce effluent pentoside presumably through a 'washout' mechanism⁶⁷.

Biopsy studies have confirmed the deposition of AGE in the peritoneal tissues of patients on peritoneal dialysis^{39;68-70}. AGEs are deposited in the interstitium and vessel walls and correlate with fibrosis⁶⁹, solute transport and ultrafiltration dysfunction⁷⁰, and co-localizes with VEGF in the peritoneal vasculature^{39;61}.

Conventional peritoneal dialysis fluid has a pH around 5 and uses lactate as a buffer. To date, the bioincompatible nature of these components is not clear, but appears to be less important than glucose or GDPs⁵⁵. Previous clinical experience with acetate buffered PD fluid demonstrated increased rates of UF dysfunction⁷¹.

Peritonitis

Aside from the bioincompatible nature of conventional dialysate solutions, peritonitis is a major cause of peritoneal membrane dysfunction. There is an acute effect of peritonitis following the infectious event that clinically appears as decreased UF with increased solute transport⁷². The effect of a single episode of peritonitis on long term peritoneal membrane function is less certain. Ates and colleagues have suggested a subtle

longer term UF dysfunction after a single episode of peritonitis ⁷². Others suggest that one episode of peritonitis has little long term effect ^{2;50;73}. Features of peritonitis which predict long term UF dysfunction include later infective episodes (>36 months after initiation of PD) ⁷⁴, multiple events or increased inflammatory response ^{2;75}, severe infections ⁷³, or gram negative infections ⁷⁶.

The response to acute infection involves resident peritoneal mesothelial cells and macrophages which work in a coordinated fashion to recruit other inflammatory cells including lymphocytes and neutrophils ⁷⁷. Heat killed bacterial solutions can induce peritoneal mesothelial cells to produce the inflammatory mediators interleukin (IL) -6 and prostaglandins, along with chemokines IL-8, RANTES, and monocyte chemotactic protein - 1 ⁷⁸. IL-1 and tumor necrosis factor (TNF) α are induced by bacterial products and endotoxins from peritoneal macrophages and impact on mesothelial cells and endothelium with the up-regulation of vascular adhesion molecules and increased vascular permeability.

The mechanism of long term changes after peritonitis episodes is not clear. There is an acute inflammatory event with upregulation of inflammatory cytokines ⁷⁹. Profibrotic cytokines such as transforming growth factor β (TGF β) and fibroblast growth factor (FGF) -2 ^{79;80} have also been shown to be up-regulated after peritonitis episodes. In an animal model of peritonitis, Combet and colleagues demonstrated increased expression of endothelial nitric oxide synthase (NOS) in the peritoneal tissues ⁴⁵. This was associated with increased vascularization, increased protein loss, and UF failure. Nitric oxide has been identified as an important and necessary cofactor in peritoneal angiogenesis ³⁹. Using adenovirus mediated overexpression of inflammatory cytokines

IL-1 β and TNF α , we have shown that longer term effects on vascularization and fibrosis can be induced in the peritoneum after transient cytokine overexpression and IL-1 β appears to be more potent in inducing these changes ⁸¹. This finding is supported by other research in peritoneal inflammation. In vitro experiments suggest that IL-1 can lead to the upregulation of collagen, fibronectin ⁸², and PAI-1 ⁸³ gene expression in peritoneal mesothelial cells. Peritoneal effluent from patients with peritonitis leads to increased mesothelial cell expression of the anti-inflammatory glycosaminoglycan hyaluronan and this effect was blocked with antibodies to IL-1 ⁸⁴. Several studies have noted that IL-1 production from peritoneal mesothelial cells is dependent on the species of infectious agent. *S. aureus* supernatant induces higher IL-1 expression than *S. epidermidis*, which is higher than *E. coli* ^{83;85}. Finally, there has been a demonstrated direct effect of IL-1 on the induction of VEGF and angiogenesis ⁸⁶⁻⁸⁸.

Therefore, there are pathways from acute inflammation to longer term fibrosis and angiogenesis in the peritoneum which may explain the association between peritonitis and UF dysfunction seen in several observational studies. The increased fibrogenic and angiogenic nature of IL-1, combined with the observation that variability in inflammatory response depends on bacterial species, suggests a mechanism whereby different infectious episodes could have different long term effects on peritoneal membrane function. This also suggests that early identification and treatment with antibiotics, combined with an IL-1 inhibiting agent ⁸⁹, may be the optimal therapy for PD related peritonitis.

Uremia and systemic inflammation

The uremic environment has been associated with an increased expression of AGE proteins ⁹⁰. In rats made uremic with subtotal nephrectomy, significant changes in the peritoneal tissues occurred including increased AGE deposition, increased vascularization with increased expression of NOS, VEGF, and FGF-2 ⁹¹. The effect of uremia on the peritoneal tissues have been confirmed in two human peritoneal biopsy studies ^{35;38} that demonstrated increased submesothelial thickening in pre-dialysis patients.

Systemic inflammation has been a controversial topic in chronic renal failure and in peritoneal dialysis. The observed association between markers of malnutrition, inflammation, and atherosclerosis suggests systemic inflammation is a common underlying condition in patients with renal failure ⁹². In PD patients, the association between peritoneal membrane transport properties and markers of systemic inflammation has been contradictory. Wang and colleagues did not find any association in numerous markers of inflammation assessed ⁹³. In subsequent work, patients with signs of inflammation, malnutrition, and atherosclerosis had higher peritoneal transport at the initiation of dialysis ⁹⁴. In a second study, high C-reactive protein and declining RRF were both associated with increasing peritoneal membrane transport status from initiation to 1 year on dialysis ⁹⁵. In a retrospective study, we found patients with a high transport peritoneal membrane had low serum albumin before the initiation of PD and suggested this was the result of systemic factors such as inflammation or overhydration ⁹⁶.

Overhydration is likely a result of a high transport status peritoneal membrane with concomitant UF dysfunction, but it may also impact on the peritoneal membrane

function⁹⁷. Theoretic considerations suggest that peritoneal tissue hydration leads to dilution of extracellular matrix components which increases fluid conductivity and may thus lead to increased tissue and lymphatic reabsorption⁴⁸. An older intervention study suggests that a decrease in extracellular fluid volume in PD patients, either through aggressive diuresis or hemofiltration, improved the UF properties of the peritoneal membrane⁹⁸.

Finally, two studies have demonstrated an association between diabetes and high transport status early in peritoneal dialysis^{15,56}. The assumption is that high glucose levels on either side of the peritoneal membrane will lead to histologic changes with associated functional changes.

Treatment of Ultrafiltration Dysfunction

The treatment of UF dysfunction in PD patients involves three steps. First, the problem of UF dysfunction must be recognized through regular clinical volume assessment and routine measurement of peritoneal membrane function including net UF⁹⁹. Prevention, with protection of the peritoneal membrane from injury due to peritonitis and bioincompatible dialysis solutions, is an important second step. Finally, treatment strategies for established UF dysfunction are limited at present, but should expand with further research in this area.

Icodextrin

Icodextrin is a promising dialysate solution for improving UF function. Icodextrin is a colloid and appears to promote UF through small pores in the peritoneal membrane

and is therefore effective in PD patients with increased peritoneal vascular area or in the setting of acute peritonitis ¹⁰⁰. In large randomized studies in which icodextrin was used in the night time dwell in continuous ambulatory PD patients ¹⁰¹ or the daytime dwell in automated PD patients ¹⁰², a similar UF rate was observed to that obtained with a hyperosmolar glucose solution (3.86% glucose). Two studies have assessed the value of icodextrin in patients with UF failure. The first prospective study took 17 patients with impending technique failure due to UF dysfunction and prescribed icodextrin ¹⁰³. Net daily UF increased by 570 ml, blood pressure improved, and technique survival was prolonged by one year. Patients who benefited the most from icodextrin were those who initially had very low UF.

Biocompatible solutions

Biocompatible solutions are produced in two chambered bags so that glucose is separated from the electrolytes at a very low pH. This prevents the formation of GDPs during heat sterilization ⁵⁹ and allows for adjustment of final pH and buffer.

Conventional PD solutions have detrimental effects on cellular proliferation, function and inflammation which are reversed by more physiological solutions ^{60;104}. Animal studies using more physiological solutions have demonstrated reduced AGE deposition and preserved UF ¹⁰⁵.

In a six month trial of 106 PD patients randomized to conventional or physiological solution, Tranaeus and colleagues found an increase in UF (150 ml/day) with stable peritoneal membrane characteristics ¹⁰⁶. In a voluntary 6 month extension of this study, significantly lower peritonitis rates were reported in the physiological solution

group. A common finding among clinical trials of physiological solutions is an increase in effluent CA125 suggesting improved mesothelial cell viability, and a decrease in effluent hyaluronan concentration, which is interpreted as a decrease in peritoneal inflammation ^{107;108}

Alteration of the interstitial compartment

Hyaluronan is a glycosaminoglycan which is present in the interstitium of the peritoneum. In animal models, hyaluronan has been shown to increase net ultrafiltration and decrease markers of inflammation ^{109;110}. These animal studies have also shown a decrease in protein transport into the peritoneal fluid. Aside from demonstrated anti-inflammatory properties, hyaluronan is felt to effect the interstitium and fluid absorption from the peritoneum ⁴⁹.

Small observational studies in humans have been carried out to study the surface active phospholipid phosphatidylcholine and have demonstrated an increase net UF when delivered orally ¹¹¹ or intraperitoneal ¹¹².

Anti-angiogenic therapy

Our results demonstrate that anti-angiogenic therapy using gene transfer of angiostatin improves net UF in an animal model of peritoneal dialysis ⁴⁰. There are several advantages and disadvantages to using adenoviral mediated gene transfer in the peritoneum in human disease. The type 5 adenovirus used in our experiments is readily taken up by mesothelial cells in the naïve peritoneum ¹¹³, and in peritoneal tissue exposed to chronic peritoneal dialysis fluid ⁵². The adenovirus is readily purified to high titre ¹¹⁴.

The negative aspects of peritoneal adenovirus gene therapy is the acute inflammatory reaction, the possibility that the adenovirus will be transported to the liver or systemically, and the transient nature of the expression. In our experiments, we found the null adenovirus had a minimal impact on the peritoneum. There was a transient inflammatory response which resolved by 48 hours after infection ⁸¹. At very high doses, we did see expression of the virus in liver parenchyma, which suggested the adenovirus did transport out of the peritoneum, probably via lymphatic or hematologic routes.

The transient nature of gene expression with adenovirus gene transfer is likely the most limiting problem. The fibrosis and angiogenesis seen in patients on PD occur over a 2 to 4 year time period. Gene therapy for this problem would likely have to persist for considerably more time than the 10 to 14 days we saw with adenovirus mediated gene transfer ¹¹³. On the other hand, we have investigated strategies for prolonging adenovirus gene transfer effects using repeated application with immunosuppression to prevent the host acute immune response ¹¹⁵. In conjunction with biocompatible solutions, one could imagine a therapy whereby adenovirus mediated gene transfer is used to acutely decrease the vascularization of the peritoneal tissues, and biocompatible solutions are used to prevent recurrent of vascularization.

Ex-vivo gene transfer has been used in several animal models. In this technique, mesothelial cells are cultured and stably transfected with a gene of interest, the peritoneum is denuded using a gelfilm application, and the transfected mesothelial cells are re-inserted ¹¹⁶. This leads to a more prolonged transgene expression, however, natural turn-over of mesothelial cells leads to eventual removal of the transfected cells.

The observation that decreasing the vasculature leads to improvement in ultrafiltration function suggests that non-gene therapy approaches should also be effective. Several anti-angiogenic agents, included thalidomide ¹¹⁷, TNP-470 ¹¹⁸, and VEGF receptor tyrosine kinase inhibitors ¹¹⁹ are available and have been used in animal and human studies and may be suitable for IP or oral administration for patients on PD with UF dysfunction.

Peritoneal fibrosis and the role of TGF β

The response of the peritoneal tissues to systemic and local toxic and inflammatory events is similar to the process of wound healing and tissue repair in other organs. This 'response to injury' process is a ubiquitous physiological event which occurs with slight variations in the majority of living organ systems ¹²⁰. Normal wound healing returns the damaged tissue close to the pre-injury state. With severe or prolonged injury, residual scarring and some loss of function are probable. In aberrant wound healing, the process becomes dysregulated and collagen deposition occurs with minimal ongoing inflammation. This collagen deposition leads to progressive fibrosis and loss of organ function. Fibrosis is the end result of multiple injurious and inflammatory diseases and affects virtually every organ system. There is a high cost, mortality, and morbidity associated with progressive organ fibrosis ¹²¹⁻¹²³.

The fibrogenic response involves a coordinated interaction between cellular and cytokine components. The initial insult leads to an upregulation of fibrogenic cytokines released from platelets during clot formation in wounded tissue ¹²⁰. Profibrotic cytokines can also be directly stimulated by high glucose, dialysis fluid, or inflammation, in the

peritoneum^{80;124-126}. The initial tissue insult may also lead to an influx of inflammatory cells that are an alternate source for profibrotic cytokines¹²⁰. Along with TGF β 1, other cytokines such as TGF β 2, TGF β 3, platelet derived growth factor (PDGF), FGFs, and connective tissue growth factor (CTGF)¹²⁷, are involved in the initiation of the fibrogenic process. Plasminogen activator inhibitor (PAI) -1 is likely involved at an early stage in peritoneal fibrosis with the inhibition of fibrinolytic activity on the surface of the peritoneal tissue and the subsequent accumulation of fibrin tissue which forms the initial scaffold for subsequent peritoneal fibrosis¹²⁸.

Fibroblasts enter the site of tissue damage drawn by chemotactic factors such as PDGF. They transform into myofibroblasts under the influence of mechanical stress and TGF β ¹²⁹. The hallmark of early wound healing is the elaboration of collagen, new blood vessels, and nerves, in a temporary healing structure called granulation tissue¹²⁰. The progression and maintenance of subsequent fibrosis is likely the result of a balance between collagenolytic enzymes (matrix metalloproteinases (MMP)) and their inhibitors (tissue inhibitor of metalloproteinase (TIMP)). The resolution of the fibrogenic process also involves apoptosis of myofibroblasts¹²⁹.

TGF β 1 is a multifunctional cytokine involved with many wound healing processes; fibroblast activation¹²³, collagen deposition¹³⁰, inhibition of fibrinolysis through PAI-1¹²⁸, and angiogenesis^{113;130-133}. TGF β 1 is also involved in cell cycle regulation through cyclin dependent kinase inhibitor p21¹³⁴ inhibition and has immune modulating activity. Specifically, TGF β inhibits T-helper cell differentiation and regulates lymphocyte proliferation¹³⁵. TGF β also prevents T-cell activation in the setting of apoptosis and is therefore potentially crucial in the maintenance of self-tolerance¹³⁶.

TGF β is a member of a family of cytokines including activins and bone morphogenic proteins, which bind to the receptor TGF β RII that then recruits and activates TGF β RI. TGF β RI initiates a cellular signaling cascade through its serine/threonine kinase activity ¹³⁴. Non-signaling membrane associated proteins (TGF β RIII (biglycan) and endoglin ¹³⁷) modify the interaction between TGF β and its receptor complex. TGF β and related cytokines signal predominantly through Smad signal transduction proteins ¹³⁸. Non-Smad signaling after TGF β RI receptor ligation has also been demonstrated.

TGF β 1 exists in tissues in an inactive form bound to latency associated protein. It is activated through cleavage by integrin α V β 6, thrombospondin, plasmin, or cathepsin. The activity of unbound TGF β 1 is also modulated by matrix associated proteoglycans such as decorin and biglycan ¹³⁹. CTGF, a TGF β 1 induced protein, also binds to and increases the activity of TGF β 1 ¹⁴⁰.

Angiogenesis in fibrotic peritoneal tissue

We have demonstrated that overexpression of TGF β 1 in the peritoneum using an adenovirus vector leads to fibrosis, angiogenesis, and similar functional changes as is seen in patients on long term peritoneal dialysis ¹¹³. Human peritoneal biopsy studies have demonstrated a progression of fibrosis ³⁵ and angiogenesis ³⁹ with time on dialysis, and also support a direct correlation between peritoneal fibrosis and angiogenesis ¹⁴¹. Taken together, these data suggests that angiogenesis is a direct result of the fibrotic process and occurs in response to profibrotic cytokines. However, certain components of dialysis fluid, such as GDPs, can directly upregulate VEGF ⁶¹, and AGE deposition

occurs in vascular walls and co-localizes with VEGF in the peritoneal vasculature ^{39;61}. These data suggests that there are fibrosis independent pathways to angiogenesis in the peritoneum and may explain the observation that peritoneal fibrosis is ubiquitous, but UF dysfunction due to increased peritoneal vascularization occurs in a smaller proportion of PD patients.

Angiogenesis is a complex and regulated process ¹⁴². It has been studied in detail because of its role in cancer and ischemic heart disease. The process of angiogenesis in the setting of wound healing and tissue repair is likely similar in general to angiogenesis in malignancy, however certain features are probably unique ¹⁴³. In general, three distinct processes are involved in new blood vessel growth. Vasculogenesis is the appearance of new vessels from endothelial stem cells. Angiogenesis describes the sprouting of new vessels from existing vessels. Arteriogenesis is the maturation process whereby new blood vessels become less permeable and recruit vascular smooth muscle cells (VSMC) ¹⁴⁴.

In pathologic situations such as malignancy and wound healing, angiogenesis is the primary mechanism of new blood vessel growth. Existing vascular structures undergo a series of changes at the initiation of angiogenesis. Vessels demonstrate increased permeability caused by VEGF, angiopoietin (ANG) -2 ¹⁴⁵, and placental growth factor ¹⁴⁶. Other factors involved in early vascular permeability include inflammatory markers such as TNF α ⁸¹, nitric oxide ¹⁴⁷, and prostaglandins ¹⁴⁸. Next is the detachment of endothelial cells and the elaboration of basement membrane degradation enzymes such as MMP and PAI-1, which allow the migration of endothelial cells into interstitial tissue ¹⁴⁴. Integrins such as α v β 3 are integral in endothelial cell migration and survival ¹⁴⁹.

Endothelial cells proliferate and form primitive tubules. These structures recruit vascular smooth muscle cells (VSMC) through cytokines such as PDGF¹⁵⁰. Endothelial cells mature and become resistant to apoptosis through the vascular stabilization factor ANG-1¹⁵¹.

VEGF is the key angiogenic and vasculogenic cytokine and induces endothelial cell proliferation, migration¹⁵², and proteolytic activity¹⁵³. VEGF exists as a series of isoforms with differing angiogenic and heparin binding properties¹⁵³. There are several VEGF related proteins including placental growth factor, VEGF-B, VEGF-C, and VEGF-D, which have differential angiogenic and lymphangiogenic activities¹⁵². VEGF acts through its receptors VEGFR-1 (Flt-1), VEGFR-2 (Flk-1/KDR), and VEGFR-3¹⁵³. VEGFR-2 is likely the dominant receptor which induces signaling pathways. VEGFR-1 may have some signaling properties and may also act as a decoy molecule. VEGFR-3 binds preferentially to VEGF-C and has a role in lymphangiogenesis. A fourth receptor, neuropilin-1, has been recently described and increases the activity of VEGFR-2¹⁵². VEGF was initially identified as vascular permeability factor and has a potent effect in increasing macromolecular transport across the vessel wall¹⁵². VEGF is up-regulated in the setting of hypoxia through a cofactor, hypoxia inducible factor (HIF) -1¹⁴⁴. Recently, it has been identified that VEGF is modified in vitro by CTGF which binds to and inhibits the protein activity¹⁵⁴.

PDGF and FGF-2 are other growth factors independently associated with new blood vessel growth¹⁴³. Not only can these two growth factors induce endothelial cell mitogenesis, but they also appear to induce the recruitment of VSMCs and thus complete

the blood vessel maturation process ¹⁵⁰. VEGF and FGF-2 have synergistic effects on blood vessel growth ¹⁵⁵.

Other families of cytokines, the angiopoietins and ephrins, are involved in the modification of established or newly formed blood vessels ^{156;157}. ANG-1 and ANG-2 both interact with the Tie-2 receptor on endothelial cells with ANG-1 likely promoting receptor activation and ANG-2 acting to block this kinase activity. ANG-1 is secreted by perivascular cells ¹⁵⁸ and is possibly upregulated in the fibrotic setting ^{159;160}. ANG-2 is secreted by endothelial cells and its production is induced by VEGF, FGF, and hypoxia ¹⁶¹. ANG-1 induces a more mature phenotype in endothelial cells with decreased intracellular macromolecular transport ¹⁴⁵ and resistance to apoptosis ¹⁶².

Fibrosis and angiogenesis are integrated processes ¹⁴³. Fibrosis could be viewed as a benign tumor, and the cells forming the fibrotic tissue require metabolic support through angiogenesis ¹⁶³. With this view, interruption of angiogenesis may impair the fibrotic process and several experiments have been carried out to demonstrate this ¹⁶⁴. Agents involved in the fibrotic process also appear to induce angiogenesis. FGF-2 and PDGF are involved in both fibrogenesis and angiogenesis. Nitric oxide is a necessary cofactor with VEGF in angiogenesis ¹⁶⁵ and nitric oxide synthase is upregulated in the peritoneal tissues in patients on PD ³⁹. The injured or fibrotic tissue may suffer from hypoxia due to inadequate vascularization ¹⁶³, and this may induce angiogenesis through HIF-1 α . The inflammatory cytokine IL-1 is both strongly pro-fibrotic ¹⁶⁶ and induces a sustained angiogenesis in peritoneal tissues ^{81;167}.

The role of TGF β in angiogenesis has been somewhat controversial. In cell culture, TGF β has a biphasic effect on the growth of endothelial cells; facilitating at low

dose but inhibitory at high dose ¹³³. TGF β synergistically acts with HIF-1 α ^{131;168} and high glucose ¹³⁰ in upregulating VEGF. TGF β also directly upregulates VEGF in vitro and in vivo ^{113;169}. TGF β appears to induce expression ANG-1 and may lead to the stabilization of blood vessels in fibrogenesis ^{160;170}. Subclasses of TGF β RI, activin receptor-like kinase (ALK) -1, may be involved in signaling involved in vascular maturation ¹⁷¹. Mutations of endoglin, a TGF β receptor complex related molecule, leads to the human disease hereditary hemorrhagic telangiectasia ^{137;172}. TGF β also has a role in the maturation of VSMCs after their recruitment by PDGF ¹⁷³. Finally, TGF β knock out mice have lethal defects in blood vessel maturation and hematopoiesis ¹⁷⁴. Therefore, TGF β 1 has putative angiogenic roles both in the initiation phase through VEGF, and in the maturation phase.

Angiogenesis appears in the setting of fibrosis and is related to pro-fibrotic factors. There is preliminary evidence that angiogenic factors may have a direct role in collagen expression and fibrosis. Hypoxia upregulates VEGF and collagen 1 gene expression in rat model of liver fibrosis ¹⁷⁵. In a transgenic mouse model, combined overexpression of VEGF and ANG-2 lead to significant cardiac fibrosis ¹⁷⁶.

Animal Models of PD

Animal models are required in PD research in order to study changes in the complex interaction between solute transport, fibrosis, and angiogenesis. Rats have been used most commonly as they are large enough to manipulate for accurate intraperitoneal injection and indwelling catheter placement but are small enough to be reasonably economical. Mice, rabbits, and dogs have been used in the past ¹⁷⁷.

Two main models are utilized; the first is an acute model where fluid is placed by intraperitoneal injection. Some form of functional peritoneal test is performed before tissue is harvested for analysis. We have used a simplified PET where net UF, glucose, creatinine, and albumin transport is measured after a 4 hour dwell with 2.5% glucose dialysis solution ¹¹³. Others have used a more extensive evaluation with frequent intraperitoneal samples taken and a volume marker such as ¹³¹I labeled albumin used as a volume marker ¹⁷⁸. This allows the measurement of UF, and solute transport, and allows the estimation of the relative contribution of aquaporin transport, increased vascular surface area, and interstitial / lymphatic uptake to net fluid flux across the peritoneal membrane.

We have used this acute model to study the effect of overexpression of cytokines using adenovirus mediated gene transfer ^{81;113}, which are effectively transferred to the peritoneal tissues. The effect of pharmacologic agents such as hyaluronan ¹⁰⁹ or nitric oxide inhibitor ¹⁷⁹ has been studied with these acute models. The effect of infectious episodes on peritoneal structure and function has also been studied with this acute model ⁴⁵. Combined with intravital microscopy, the effect of different dialysis solutions on vascular recruitment and leukocyte adhesion can be evaluated ⁵⁵.

Chronic models involve the surgical placement of an indwelling catheter with daily infusion of dialysis solution. This model has been used extensively to study the effect of standard PD fluid on peritoneal tissue ^{52;180}, to study biocompatible PD solutions ¹⁸¹, and to study the effects of novel therapies on peritoneal function ⁴⁰. These models have also been used to study the relative roles of systemic factors such as uremia, and local effects of PD solutions, on the peritoneum ⁹¹.

Summary

The main finding in peritoneal biopsies of patients on long term peritonitis is fibrosis, submesothelial thickening, and loss of the mesothelial cell layer³⁵. This observation was difficult to reconcile with the hypothesized cause of loss of ultrafiltration function which appeared to be mainly due to increased vascularization of the peritoneal tissues³. Subsequent biopsy studies have confirmed an association between vascularization of the peritoneum, and ultrafiltration dysfunction^{35;38}.

I hypothesize that ultrafiltration dysfunction is caused by increased peritoneal vascularization with a concomitant increase in glucose transport from the peritoneal cavity and rapid loss of the ultrafiltration gradient. Further, I hypothesize that this increased vascular surface area is induced by profibrotic and inflammatory cytokines such as TGF β , IL-1 β , and TNF α through upregulation of angiogenic cytokines such as vascular endothelial growth factor (VEGF).

Our work in this area has confirmed the association between angiogenesis and UF failure⁵² and demonstrated a causality by experimentally reducing the vasculature using adenovirus mediated gene transfer of angiostatin with subsequent improvement in ultrafiltration⁴⁰. We also further explored the causes of angiogenesis in the fibrotic environment and identified that TGF β ¹¹³ and IL-1 β ⁸¹ are both strongly pro-fibrotic and angiogenic in the peritoneum. These observations have several implications. The goal of improving the peritoneal membrane for patients on peritoneal dialysis should now focus on anti-angiogenic therapy, along with methods to prevent injury to the peritoneal membrane. We have demonstrated that IL-1 has a prominent role in inducing peritoneal damage. It is possible that therapy aimed at IL-1 may prevent progressive peritoneal

membrane damage in the setting of acute peritonitis. Finally, we have created a new model to study angiogenesis in the setting of fibrosis. This will allow further evaluation of interventions, and further experiments to understand the role of such agents as the angiopoietins, CTGF, TGF β 3, matrix metalloproteinases, and their inhibitors in the process of angiogenesis and fibrogenesis.

CHAPTER 2: GENE TRANSFER OF TRANSFORMING GROWTH FACTOR β TO THE RAT PERITONEUM: EFFECTS ON MEMBRANE FUNCTION

The following paper is published in the Journal of American Society of Nephrology 2001; 12: 2029-2039. The goal of this work was to investigate the effect of transient overexpression of TGF β 1 in the peritoneum. The adenovirus construct, AdTGF β 1 was created by Dr. Patricia Sime. It was made with standard co-transfection techniques using the porcine TGF β 1 gene plasmid (supplied by Dr. Anita Roberts). This gene sequence is mutated so that a serine replaces a cysteine at the 223 and 225 locations. This mutation impairs the ability of TGF β 1 to bind to its latency associated protein, and it is therefore secreted in the active form.

We demonstrate expression of the adenovirus vector transgene between 4 and 14 days after infection. Using an adenovirus construct expressing the reporter gene β -galactosidase, we demonstrated virtually 100% adenovirus uptake by mesothelial cells. We also identified a lack of significant inflammatory reaction to our control adenovirus AdDL70. AdTGF β 1 had a dramatic effect on the structure and function of the peritoneum. There was submesothelial thickening with persisting angiogenesis. There was a significant ultrafiltration dysfunction that persisted until 28 days after adenoviral infection. This was associated with increased transport of glucose out of the peritoneal fluid, and albumin into the peritoneal fluid.

We also identified an increased presence of VEGF in the peritoneal effluent of rats exposed to AdTGF β 1. We explored this further in cell culture experiments, and

demonstrated the production of VEGF by primary mesothelial cells exposed to recombinant TGF β 1.

These experiments, therefore, suggested that TGF β 1 up-regulation may explain the changes seen in patients on long term peritoneal dialysis.

The concept for this paper was developed by Dr. Gauldie, Dr. Cathy Hoff, and me. I designed the experiments with suggestions on the animal model from Dr. Hoff and Dr. Gauldie. I developed the concept for, and the experiments to measure, angiogenesis and VEGF. The animal experiments were carried out by Dr. Kolb, Tom Galt, and me. I carried out all benchwork including ELISA, hydroxyproline, and cell culture. Histology support was carried out by Mary Jo Smith; I performed the immunohistochemistry. I analyzed the data and wrote the paper. The manuscript was reviewed by Dr. Gauldie, Dr. Hoff, and Dr. Shockley.

Gene Transfer of Transforming Growth Factor- β 1 to the Rat Peritoneum: Effects on Membrane Function

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Abstract. Long-term peritoneal dialysis is limited by physiologic changes in the peritoneum that lead to ultrafiltration failure. To determine the role of profibrotic cytokines in the alteration of peritoneal transport, a rodent model of transforming growth factor- β (TGF- β)-mediated peritoneal fibrosis was established. An adenoviral vector driving the active form of TGF- β 1 (AdTGF β 1) was administered intraperitoneally, and peritoneal structure and function were evaluated for 28 d after infection. Seven days after AdTGF β 1 infection, thickening of the peritoneum, with cellular proliferation and increased vascularization, was noted. By day 28, there was persistent thickening and extensive collagen deposition. The mesenteric collagen content was significantly elevated, compared with control adenovirus-treated animals, 21 d after infection (2.9 versus 1.8 mg hydroxyproline/g tissue, $P = 0.006$). Blood

vessel density, as measured using factor VIII immunohistochemical analyses, was significantly increased from day 4 to day 21 but decreased by day 28. Animals infected with AdTGF β 1 demonstrated increased transport of solutes and decreased net ultrafiltration, which was maximal on day 7 and returned to baseline levels by day 28. It was demonstrated *in vitro* and *in vivo* that TGF- β 1 induced production of vascular endothelial growth factor. Overexpression of TGF- β 1 after adenovirus-mediated gene transfer causes peritoneal fibrosis, neoangiogenesis, and increased peritoneal membrane solute transport. This model should allow further delineation of the relative contributions of profibrotic and angiogenic cytokines to changes in peritoneal function and may lead to potential new interventions for peritoneal membrane failure.

Peritoneal dialysis is an attractive option for the treatment of patients with end-stage renal disease. However, recurrent peritonitis and alterations in the peritoneal membrane transport of solutes complicate long-term peritoneal dialysis (1). The most common functional change in the peritoneal membrane with long-term peritoneal dialysis is ultrafiltration failure, which is the cause of technique failure for 9 to 27% of patients undergoing peritoneal dialysis (2). The relationship between the duration of peritoneal dialysis and ultrafiltration failure has been confirmed (1,3–5). In population-based studies, ultrafiltration was inversely correlated with increased peritoneal membrane transport of solutes (1,5,6).

The pathophysiologic features of the changes in peritoneal membrane transport with time are not well understood. The main barrier to peritoneal transport of solutes seems to be the vascular wall (7,8). The observed changes in peritoneal membrane function with time are therefore attributable to increases in the permeability of individual blood vessels or increases in

the total vascular surface of the peritoneum (9,10). Recent studies of human peritoneal biopsy samples demonstrated increases in the density of the peritoneal microvasculature with dialysis time (11,12). Human (13,14) and animal (15) biopsy samples revealed a generalized thickening of the submesothelial collagenous zone of the peritoneum, with connective tissue deposition and microvascular sclerosis, corresponding to the duration of dialysis. α -Smooth muscle actin (α -SMA)-positive cells, suggesting myofibroblast differentiation induced by fibrogenic cytokines (16), were identified in the peritoneum and were correlated with the degree of fibrosis and the density of blood vessels (13).

The role of increased peritoneal vascular area in ultrafiltration failure suggests that angiogenic cytokines such as vascular endothelial growth factor (VEGF) are upregulated in the process of peritoneal dialysis. VEGF has been identified, by both immunohistochemical and immunoblot analyses, in biopsy samples from patients undergoing long-term dialysis (11). VEGF concentrations in dialysate fluid have been correlated with measures of solute transport among patients undergoing chronic peritoneal dialysis who are in stable condition (17).

VEGF may be induced directly in the peritoneum of patients undergoing peritoneal dialysis. Glucose degradation products, formed by heat-sterilization of dialysate, have been demonstrated to be toxic to mesothelial cells (18) but also enhance the production of VEGF by those cells (19). Rats with streptozocin-induced diabetes mellitus exhibit changes in peritoneal membrane transport similar to those of patients undergoing

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long-term dialysis, and this effect can be reversed with the intraperitoneal administration of anti-VEGF antibodies (20).

Renal failure and dialysis expose the peritoneum to inflammatory mediators (21), and this exposure may induce the expression of profibrotic cytokines such as transforming growth factor- β (TGF- β). Peritoneal mesothelial cells may play a role in this process, because TGF- β can be induced in mesothelial cells by high-concentration glucose solutions and spent dialysate (22). Two studies noted increased dialysate TGF- β concentrations with increased duration of dialysis (23,24), and dialysate TGF- β concentrations were correlated with peritoneal membrane solute transport in a population of peritoneal dialysis patients in stable condition (25). In addition to local factors, membrane transport at the initiation of peritoneal dialysis is variable and predictive of outcome (26), which suggests that systemic processes, such as diabetes mellitus, atherosclerosis, systemic inflammation, and malnutrition, may also play a role in membrane dysfunction (27).

TGF- β is a multifunctional cytokine that is central in the process of fibrogenesis (28). By inducing fibroblasts to increase production of collagen I and III and increasing expression of the tissue inhibitor of metalloproteinase (29), TGF- β promotes the formation of extracellular matrix. As part of the fibroproliferative response, TGF- β has been demonstrated to enhance the expression of VEGF in several cell lines (30–32). In turn, VEGF is a mitogen for endothelial cells and induces angiogenesis (33). VEGF is also a vascular permeability factor and may directly induce endothelial cells to increase cellular fenestrations (34).

To study the relationship between TGF- β and changes in the structure and function of the peritoneum, we used adenovirus-mediated gene transfer of active TGF- β 1 to the peritoneum in a rat model. This adenoviral construct (AdTGF β 1) was previously used to transfer this gene to the rat lung, resulting in transient overexpression of TGF- β 1, with subsequent progressive pulmonary fibrosis (35).

We demonstrated that transient overexpression of TGF- β 1 after intraperitoneal administration leads to histologic changes consistent with peritoneal fibrosis, including thickening of the submesothelial zone and increased interstitial collagen deposition. There was a TGF- β 1-mediated increase in dialysate VEGF concentrations and increased vascularization. We observed changes in solute transport induced by TGF- β that mimicked those observed in patients undergoing long-term peritoneal dialysis but were transient and resolved with the resolution of the increased peritoneal vascularity.

Materials and Methods

Adenoviral Construct

A replication-deficient recombinant adenovirus expressing the biologically active form of porcine TGF- β 1 was used in these studies. The construction of AdTGF β 1 was previously described (35). In this construct, the TGF- β 1 gene is mutated at residues 223 and 225, and the resulting protein does not bind to its latency-associated protein, allowing production of active TGF- β 1. An adenovirus construct coding for β -galactosidase (AdLacZ) (36) and an empty (null) virus

(AdDL70) (36) were also used. Virus preparations were expanded, purified, and plaque-titered as described previously (37).

Animal Studies

All animal studies were performed according to the Canadian Council on Animal Care guidelines (38). Food and water were provided *ad libitum*. Animals were anesthetized with isoflurane (MTC Pharmaceuticals, Cambridge, Canada). Sprague-Dawley rats (225 to 275 g; Harlan, Indianapolis, IN) were given intraperitoneal injections of AdTGF β 1, AdDL70, or phosphate-buffered saline (PBS) (control). To ensure intraperitoneal delivery, an intravenous catheter over a 25-gauge needle (Becton-Dickson, Franklin Lakes, NJ) was introduced through the abdominal wall of anesthetized animals. The needle was removed, the catheter was fully advanced into the peritoneal cavity, and the substance was administered. The animals received 2×10^9 plaque-forming units of adenovirus, diluted in 50 μ l of PBS, or PBS alone on day 0. Three to six animals per group were euthanized on days 4, 7, 21, and 28. The day 21 and 28 groups were repeated. Until euthanasia, a survival peritoneal equilibration test (PET) was performed in each animal on days 4, 7, 14, and 21. This test involved the intraperitoneal administration of a 0.09 ml/g dose of 2.5% Baxter Dianeal (Baxter Healthcare, Round Lake, IL). Four hours later, a tail blood sample and a peritoneal fluid sample were obtained. A PET was performed before euthanasia, and an accurate ultrafiltration measurement was made. One group was euthanized on day 0 for baseline assessments of histologic features, ultrafiltration, and membrane transport. Two animals were administered AdLacZ and euthanized on day 4 for β -galactosidase analysis.

Solute Transport

Whole blood was centrifuged at 5000 rpm for 10 min, and the serum was removed. Peritoneal fluid samples obtained at the time of PET were centrifuged at 1500 rpm for 5 min, and the supernatant was removed for analysis. Samples were analyzed with an Hitachi 917 automated chemistry analyzer (Roche Diagnostics, Laval, Canada). Creatinine levels were measured using the picric acid-modified Jaffé method, and peritoneal samples were corrected for glucose levels [corrected creatinine concentration = measured creatinine concentration - (0.137 \times measured glucose concentration)]. Albumin levels were measured by using the bromocresol green method, and glucose levels were measured by using a standard enzymatic test. The following measures of peritoneal transport were calculated: creatinine and albumin dialysate/plasma concentration ratios after a 4-h peritoneal dwell time and the glucose 4-h/initial dialysate concentration ratio. Solute clearance or mass transfer was also measured at time points when the animals were euthanized and an accurate ultrafiltrate volume could be assessed. Because the initial solute concentration for creatinine and albumin was 0, the 4-h clearance was calculated as follows: final Dianeal solute concentration \times final volume/plasma concentration. Mass transfer of glucose out of the peritoneum was calculated as follows: (initial dialysate glucose concentration \times initial volume) - (final dialysate glucose concentration \times final volume). These values were corrected for animal weights at the time of euthanasia.

Cell Counts and Differential Determinations

The cell pellets remaining after centrifugation of the peritoneal fluid samples were resuspended in PBS. The numbers of cells were counted by using a hemocytometer. The cells were centrifuged onto a glass slide (Shandon, Pittsburgh, PA) and stained (Biochemical Science, Swedesboro, NJ). Cell differentials were determined with at least 300 cells.

Cell Culture

Rat peritoneal mesothelial cells were collected from a whole section of anterior abdominal wall placed in an isolation device (39). The parietal peritoneal surface was washed once with PBS and once with 0.25% trypsin (Life Technologies, Burlington, Canada) for 5 min and was exposed to fresh trypsin for 30 min at 37°C. The surface was gently scraped, and the cells were recovered. After centrifugation at 1000 rpm at 4°C for 5 min, the cell pellet was resuspended in Dulbecco's modified Eagle's medium/F-12 medium with 10% fetal bovine serum (Sigma Chemical Co., St. Louis, MO), 1% penicillin/streptomycin (Life Technologies), and 0.08% Fungizone (Life Technologies), and cells were plated in 40-cm² flasks (Corning, Cambridge, MA). Mesothelial cells in culture exhibited a cobblestone appearance at confluence and were vimentin-positive, with the eccentric nuclei characteristic of mesothelial cells (data not shown). Cells were used between passage 4 and passage 8.

Cells were plated on 100-mm plates (Corning) and grown to confluence. Plated cells were rested in 1% fetal bovine serum-containing medium for 24 h and were then exposed to recombinant human active TGF- β 1 (R&D Systems, Minneapolis, MN), at concentrations of 1 or 10 ng/ml, for 12 h. The cells were then washed three times with PBS, and fresh medium with 1% fetal bovine serum was added. Supernatant samples were obtained at 12, 24, 48, and 72 h after exposure to recombinant TGF- β 1.

Histologic Assessments

Cytochemical staining for β -galactosidase was performed on samples obtained from animals 4 d after intraperitoneal infection with AdLacZ. Fresh tissue samples were fixed in 2% formaldehyde/0.2% glutaraldehyde for 3 h. Fixation was followed by staining for 6 h in a solution containing potassium ferrous cyanide, potassium ferric cyanide, and magnesium chloride (Sigma), with Triton X-100 and 5-bromo-4-chloro-3-indolyl- β -D-galactopyranoside (Boehringer Mannheim, Indianapolis, IN). The samples were stored in 70% ethanol and then paraffin-processed and -embedded, and 5- μ m sections were cut. These sections were counterstained with nuclear fast red.

Tissue samples were obtained from the anterior abdominal wall, liver, and mesentery of AdTGF β 1- and control virus-treated animals on days 4, 7, 21, and 28 and were fixed in a sufficient amount of 4% phosphate-buffered formaldehyde for 24 h. The tissue samples were then paraffin-processed and -embedded, and 3- μ m sections were cut. Cut sections were stained with hematoxylin and eosin and with Masson's trichrome stain.

Immunohistochemical Analyses

Using the same 4-, 7-, 21-, and 28-d samples, 3- μ m sections were cut on aptex-coated slides and immunohistochemical analyses were performed with antibodies to α -SMA, vimentin, and von Willebrand factor/factor VIII-related antigen (Dako Corp., Carpinteria, CA). Negative control sections were assayed in parallel, with nonimmune mouse or rabbit Ig diluted to the same concentrations as the primary antibodies. All sections were deparaffinized in xylene, followed by 100% ethanol, and were then placed in a freshly prepared methanol/H₂O₂ solution for 30 min, for blockade of endogenous peroxidase activity. After hydration to water with graded alcohols, the sections were placed in 0.05 M Tris-buffered saline (TBS) (pH 7.6). The sections for α -SMA assays were blocked for 20 min with 1% normal swine serum (NSS) in TBS, followed by overnight incubation with mouse anti-human α -SMA (1:100) in 1% NSS. Sections were then incubated with a biotinylated rabbit anti-mouse Ig (1:300; Dako) for 1 h, followed by a 45-min incubation with a streptavidin-peroxidase

conjugate (1:600; Dako). Sections for vimentin and factor VIII were pretreated before primary incubations. For vimentin assays, microwave antigen retrieval was performed with hot 1 mM ethylenediaminetetraacetate for 10 min, and sections were allowed to reach room temperature before subsequent procedures were performed. Sections were then blocked with 1% NSS for 20 min, followed by overnight incubation with mouse anti-human vimentin (1:300; Dako) in antibody-diluting fluid. Sections were incubated for 1 h with biotinylated rabbit anti-mouse Ig (1:300) in antibody-diluting fluid, followed by a 45-min incubation with a streptavidin-peroxidase conjugate (1:600; Dako) in peroxidase-stabilizing buffer. For factor VIII, the sections were digested for 17 min with 0.05% Pronase (Sigma) in TBS with calcium chloride, at room temperature. Sections were blocked with 5% normal goat serum, followed by a 1-h incubation with rabbit anti-human factor VIII (1:500) in 1% normal goat serum. Sections were then incubated with a prediluted, biotinylated, goat anti-rabbit Ig preparation, followed by a streptavidin-peroxidase conjugate (Zymed Laboratories, San Francisco, CA), according to the instructions provided by the manufacturer. All immunohistochemical incubations were performed at room temperature, and all sections were washed with 0.05 M TBS (pH 7.6) for 3 \times 5 min between incubations, except before the addition of primary antibodies. All sections were rinsed with 0.05 M acetate buffer (pH 5.0) before development with an aminoethylcarbazole chromogen substrate for 15 min. All sections were counterstained with Mayer's hematoxylin for 2 min before being mounted with glycerin gelatin.

Blood Vessel Analysis

Sections of the anterior abdominal wall that had been immunostained for factor VIII were studied in a blinded manner, by using a standardized microscope grid to count peritoneum-associated blood vessels and to measure the thickness of the peritoneum. The area examined included all tissue from the abdominal wall muscle to the peritoneal cavity. Slides contained one to four separate tissue pieces, and each slide was examined in 10 to 12 randomly selected, high-power views. All measurements were performed at the same magnification. The number of blood vessels counted per grid was standardized to a count per 1-mm length of peritoneum.

Cytokine Analyses

Peritoneal fluid and cell culture supernatants were collected and centrifuged at 1500 rpm for 5 min. The samples were analyzed using a human TGF- β 1 enzyme-linked immunosorbent assay (ELISA) kit, a mouse VEGF ELISA kit, or a rat tumor necrosis factor- α (TNF- α) ELISA kit (R&D Systems). The sensitivities of these assays were 7.0, 3.0, and 5.0 pg/ml, respectively. Samples were assayed for active TGF- β 1 concentrations without treatment; samples were also treated with HCl, pH-adjusted with NaOH, and then assayed for total (active and latent) TGF- β 1.

Hydroxyproline Assay

A portion of mesentery was frozen for an hydroxyproline assay, which was modified from the method described by Woessner (40). Tissues were weighed, homogenized in water, and centrifuged at 1000 rpm for 5 min, and the superficial fatty material was removed by vacuum suction. Solid material was precipitated with TCA, with centrifugation at 1500 rpm for 15 min at 4°C. Samples were hydrolyzed overnight in 6 N HCl at 110°C. Hydroxyproline contents were quantified with Erlich's reagent (Sigma) and were assayed by measurement of the OD at 557 nm. Standard hydroxyproline samples (Sigma) were used to create a standard curve.

Statistical Analyses

Data are presented as mean \pm SEM unless otherwise noted. Comparisons between groups were performed by *t* test. We used linear regression analyses to compare Dianeal cytokine levels and solute transport. ANOVA was used to examine the mean clearances of creatinine and albumin and the mass transfer of glucose in the treated and control animals during the course of the experiments.

Results

Adenoviral Gene Transfer to Peritoneal Mesothelial Cells

Adenovirus is highly infective for mesothelial cells, as demonstrated by widespread expression of β -galactosidase in the peritoneal cavity 4 d after AdLacZ administration (Figure 1). At the dose used, adenoviral transgene product was not observed in the liver, spleen, skeletal muscle tissue, or submesothelial interstitium. We searched for evidence of inflammation attributable to the adenovirus (Table 1). Specifically, we did not observe a significant difference between AdTGF β 1-

and AdDL70-treated animals with respect to inflammatory cells in the peritoneal fluid. The day 4 TNF- α concentrations in the dialysate were low and were not different between groups. This finding suggests that the inflammatory response to the adenovirus was limited and was similar for control and active virus. Previous work identified a mild early inflammatory response to adenovirus (as indicated by neutrophilia and increased TNF- α concentrations), which resolved by 48 h (41).

The animals that received AdTGF β 1 demonstrated significant elevations in active TGF- β 1 levels in the dialysate from day 4 to day 14 (Figure 2). Levels peaked on day 4 (1920 versus 87 pg/ml for AdDL70, $P < 0.001$). There was a significant secondary increase in total TGF- β 1 concentrations that was maximal on day 7 (16.4 versus 0.3 ng/ml, $P < 0.001$). Biologically active TGF- β 1 is the form produced after infection with AdTGF β 1, and the total (active plus latent) amount likely represents autoinduction of latent TGF- β 1 by resident peritoneal cells (42). The time course of transient expression of gene product after AdTGF β 1 infection was similar to that demonstrated in the rat lung model (35).

Effects of TGF- β 1 on the Structure of the Peritoneum

Animals were euthanized on days 4, 7, 21, and 28, and parietal and visceral peritoneal surfaces were evaluated by microscopic and immunohistochemical techniques. By day 4, there were significant changes in the peritoneum of AdTGF β 1-treated animals. There was thickening of the peritoneum, with increased cellularity and vascularization (Figure 3). By day 7, these effects were even more prominent. On day 7, increased collagen amounts were present in the peritoneum. The more superficial cells in the thickened peritoneum were predominantly vimentin-positive (data not shown). Immunohistochemical analyses demonstrated the accumulation of α -SMA-positive interstitial cells (Figure 4). Vascular proliferation was prominent in factor VIII-stained sections (Figure 4). By day 21, the proliferative response had diminished but there was still generalized thickening of the submesothelial collagenous zone, with increased vascularity. By day 28, the submesothelial zone remained thickened but the degree of vascularity had decreased.

We noted similar changes in visceral peritoneal sections from animals exposed to TGF- β 1. Specifically, mesenteric tissue demonstrated increased collagen deposition and vascularity. Also, in the day 7 samples from animals exposed to TGF- β 1, there seemed to be increased blood vessel density extending into the superficial muscle layer of the abdominal peritoneal wall. These changes were not quantified. No histologic changes were detected in the control adenovirus-treated groups, compared with PBS-treated animals (Figure 3).

Mesenteric Collagen Content

Hydroxyproline is a component of collagen, and its concentration is directly correlated with the collagen content. The hydroxyproline contents of mesenteric samples from AdTGF β 1-treated animals were unchanged, compared with control adenovirus-treated animals, on day 4 but were elevated



Figure 1. Adenovirus-mediated gene transfer to the peritoneal mesothelium. Four days after delivery of AdLacZ, the peritoneum was fixed and stained for β -galactosidase activity. (A) Whole section of the anterior abdominal wall, demonstrating extensive uniform expression (blue staining) of adenoviral gene product. (B) Section of the anterior abdominal wall, demonstrating that β -galactosidase staining was limited to the mesothelial cell monolayer. The section was counterstained with nuclear fast red. Magnification, $\times 100$.

Table 1. Inflammatory changes after adenoviral administration, measured as cellular responses and TNF- α concentrations in peritoneal fluid^a

Parameter	PBS		AdTGF β 1		AdDL70	
	Day 4	Day 7	Day 4	Day 7	Day 4	Day 7
Total cells ^b	8.3 \pm 5.3	19.2 \pm 9	5.3 \pm 2.1	10.9 \pm 3.5	13.0 \pm 2.9	20.8 \pm 10.2
Lymphocytes (%) ^c	30.7 \pm 2	20.6 \pm 7.7	32.4 \pm 12.3	22.7 \pm 11.4	17.6 \pm 8.8	29.4 \pm 20.6
Macrophages (%) ^c	56.0 \pm 8.5	44.4 \pm 10.6	61.1 \pm 10.1	73.4 \pm 11.3	69.8 \pm 6.1	43.7 \pm 29.1
Neutrophils (%) ^c	0.1 \pm 0.2	7.6 \pm 2.5	2.6 \pm 2.4	3.2 \pm 2.9	6.1 \pm 7.5	0.4 \pm 0.5
TNF- α (pg/ml)	ND	ND	32.2 \pm 17.1	42.4 \pm 22.5	25.8 \pm 3.8	35.9 \pm 13.3

PBS, phosphate-buffered saline; TNF- α , tumor necrosis factor- α ; ND, not determined.

$P > 0.05$ for comparisons between AdTGF β 1 and AdDL70 groups at each time point.

^a All values are mean \pm SD.

^b Total cell count (total cells in peritoneal fluid $\times 10^6$).

^c Differential counts (percentage of total cells).

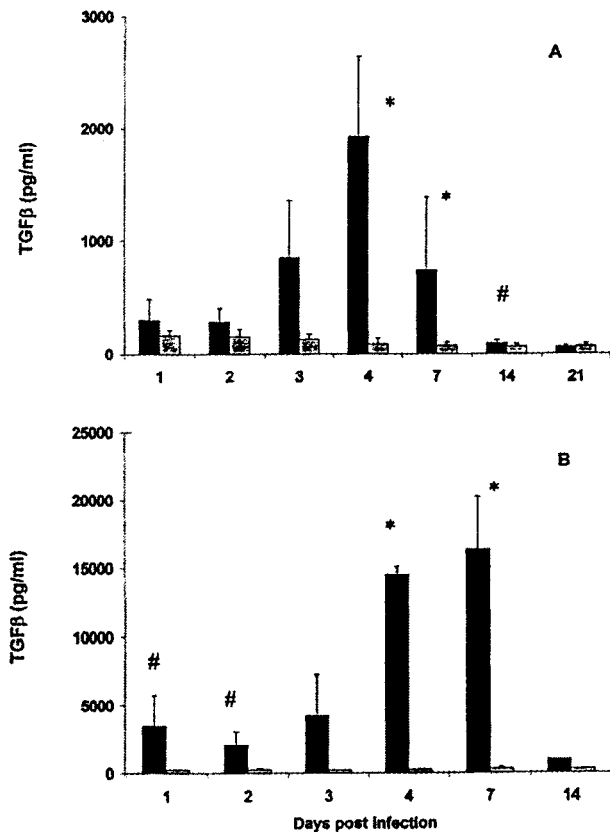


Figure 2. Transforming growth factor- β (TGF- β) concentrations in peritoneal dialysate after adenoviral infection. (A) Active TGF- β 1 concentrations in AdTGF β 1-treated animals (■) were significantly elevated, compared with AdDL70-treated animals (▨), from day 4 to day 14. (B) Total TGF- β dialysate concentrations were elevated in AdTGF β 1-treated animals on days 1 to 2 and days 4 and 7. * $P < 0.001$, # $P < 0.05$. Data are mean \pm SD ($n = 3$ or 4 for days 1 to 3, $n = 5$ to 22 for days 4 to 14).

on days 7, 21, and 28 (data not shown). The difference reached statistical significance on day 21 (2.9 ± 0.7 versus 1.8 ± 0.3 mg/g tissue for control AdDL70, $P = 0.006$).

Quantitative Morphologic Assessments

Quantitative morphologic assessments performed using a standardized microscope grid and factor VIII-stained sections demonstrated significantly greater average thicknesses of the peritoneum for AdTGF β 1-treated animals on days 4, 7, and 21 (Figure 5). With resolution of the fibroproliferative phase on day 7, the thickness of the peritoneum decreased to $67 \mu\text{m}$ by day 21. This change persisted to day 28 ($64 \mu\text{m}$), but the difference was no longer significant. The number of vessels per 1-mm length of peritoneum was maximal on day 7 (11.4 vessels/mm) and then progressively decreased from day 21 (8.9 vessels/mm) to day 28 (4.4 vessels/mm). No differences were noted between control virus-treated animals and day 0 PBS-treated animals with respect to thickness (29.8 versus $32.2 \mu\text{m}$ for day 21 control virus, $P = 0.43$) or number of blood vessels (1.2 versus 1.2 vessels/mm for day 21 control virus, $P = 0.99$) (data not shown).

Effects of TGF- β 1 on the Function of the Peritoneal Membrane

The transport characteristics of the peritoneal membrane in AdTGF β 1-, AdDL70-, or PBS-treated animals were evaluated using a modified PET (Table 2). Animals treated with AdTGF β 1 exhibited significant reductions in ultrafiltration from day 4 to day 21 ($P < 0.05$). The ultrafiltration impairment resolved by day 28 (Figure 6).

We used both the standard dialysate/plasma solute ratio and clearance (for creatinine and albumin) or mass transfer (for glucose) to analyze the alterations in transport properties after transient exposure to TGF- β 1. Because of the significant differences in net ultrafiltration identified, solute clearance or mass transfer better estimates the diffusive transport nature of the peritoneum (10). Using these measures, we observed no significant change in the clearance of creatinine (Table 2). We did, however, note a significant increase in the clearance of albumin ($P = 0.006$, compared with AdDL70, by ANOVA) and in the mass transfer of glucose ($P < 0.001$) during the course of the experiment. These differences were most apparent on days 4 and 7, and values returned to baseline levels by day 28.

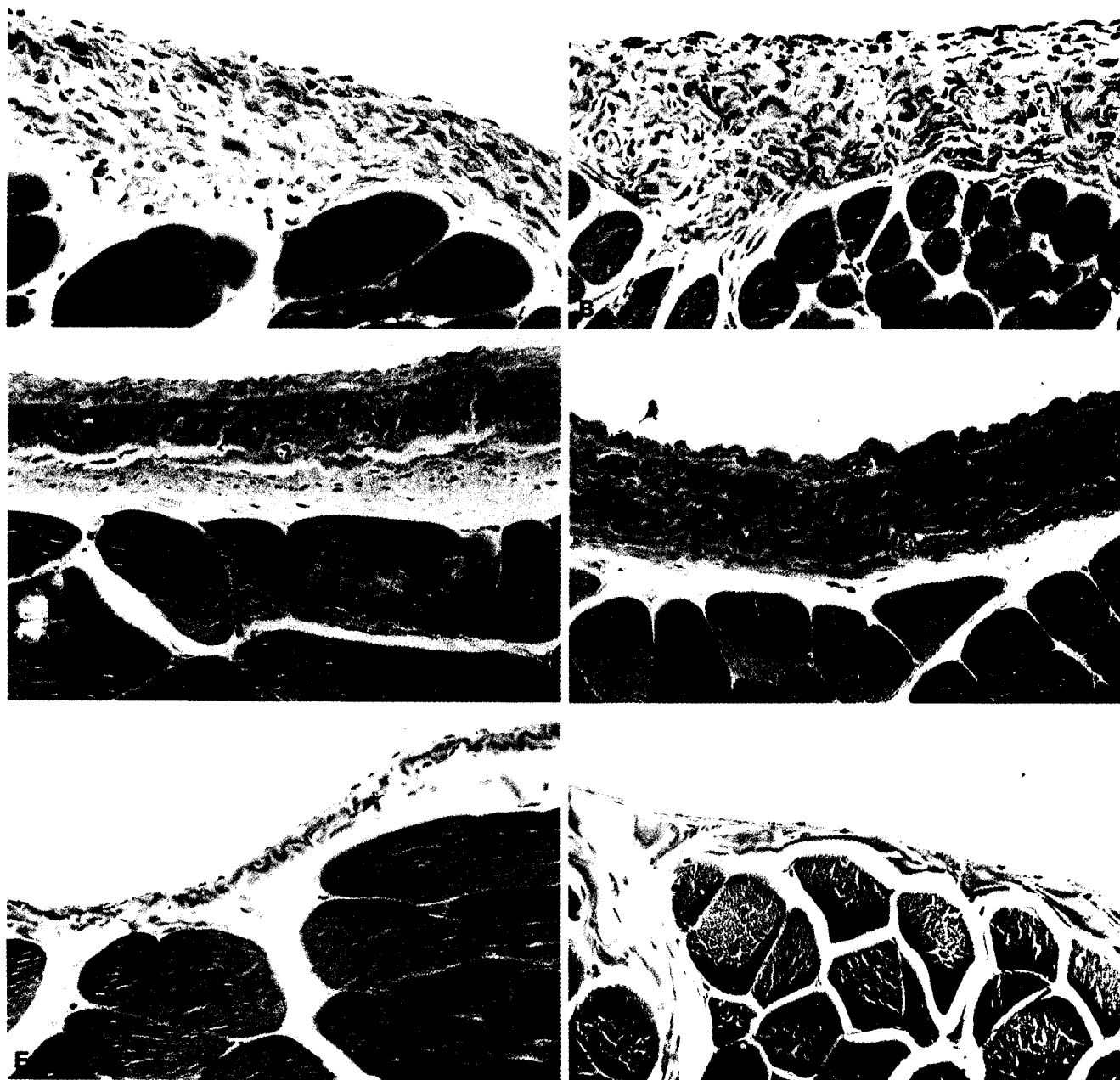


Figure 3. Histologic features of the anterior abdominal wall after exposure to AdTGF β 1. (A) On day 4 after infection, there is a fibroproliferative response, with increased thickness, cellularity, and vascularization. (B) On day 7, these changes are exaggerated, with some increase in collagen deposition. (C) By day 21, the cellularity of the peritoneal membrane has decreased but there is ongoing vascularization and extensive collagen deposition. (D) Twenty-eight days after infection, the collagen deposition remains but vascularity is decreased. (E) Four days after control virus (AdDL70) infection, the peritoneum exhibits no acute virus-related changes. (F) Normal rat peritoneum on day 21 after phosphate-buffered saline (PBS) treatment is also shown. All sections, Masson's trichrome stain. Magnification, $\times 200$.

Role of VEGF and Increased Peritoneal Membrane Transport

Primary mesothelial cells in culture produce VEGF in response to exposure to recombinant TGF- β 1 (Figure 7). Forty-eight hours after exposure to 10 ng/ml TGF- β 1, mesothelial cells produced 2.4 times the amount of VEGF measured in the supernatant by ELISA ($P = 0.001$).

The dialysate concentration of VEGF was significantly in-

creased for AdTGF β 1-treated rats, compared with control rats, on days 4 and 7 (1512 versus 92 pg/ml for AdDL70-treated animals on day 7; $P < 0.001$) (Figure 7). The VEGF dialysate concentration decreased by day 14 but remained significantly elevated, compared with dialysate from AdDL70-treated animals, until day 21 (118 versus 64 pg/ml, $P = 0.04$).

There was a correlation between TGF- β 1 exposure and VEGF expression. On day 4, the dialysate TGF- β 1 concentra-

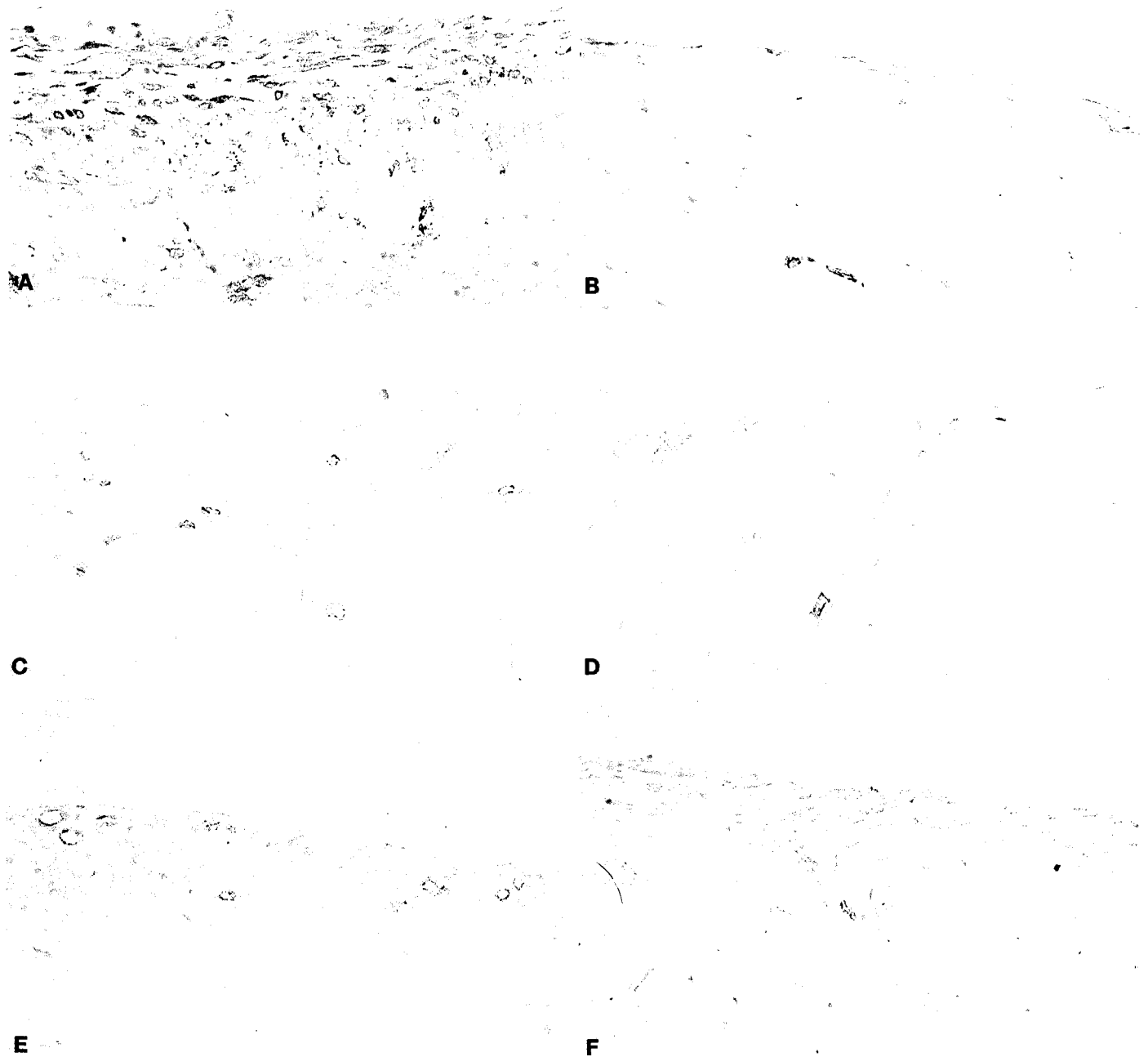


Figure 4. Immunohistochemical analyses of the anterior abdominal wall. (A) α -Smooth muscle actin (α -SMA) staining 7 d after AdTGF β 1 infection demonstrates hypercellularity of the peritoneal membrane, with staining of myofibroblasts and smooth muscle of small arterioles. (B) α -SMA staining 7 d after AdDL70 control virus infection demonstrates a normal thin peritoneal membrane, with staining of smooth muscle cells in occasional deep arterioles. (C) Factor VIII staining 7 d after exposure to AdTGF β 1 demonstrates a thickened membrane with markedly increased vascularity, which is not evident in the control virus-treated animal (D). (E) Factor VIII staining on day 21 demonstrates the persistence of increased vascularity, which is not evident in AdDL70-treated animals on day 21 (F). Magnification (all sections), $\times 200$.

tion was correlated with the VEGF concentration ($r^2 = 0.240$, $P = 0.03$) (Figure 7).

Discussion

In these experiments, we demonstrated that adenovirus-mediated gene transfer to the peritoneum is a useful technique for

studying the effects of cytokines on peritoneal membrane structure and function. Using a β -galactosidase adenoviral vector in mouse peritoneum, Osada *et al.* (43) demonstrated the effectiveness of adenoviral gene transfer to mesothelial cells and the lack of acute virus-induced changes. Similarly, we observed efficient infection of rat mesothelial cells by replica-

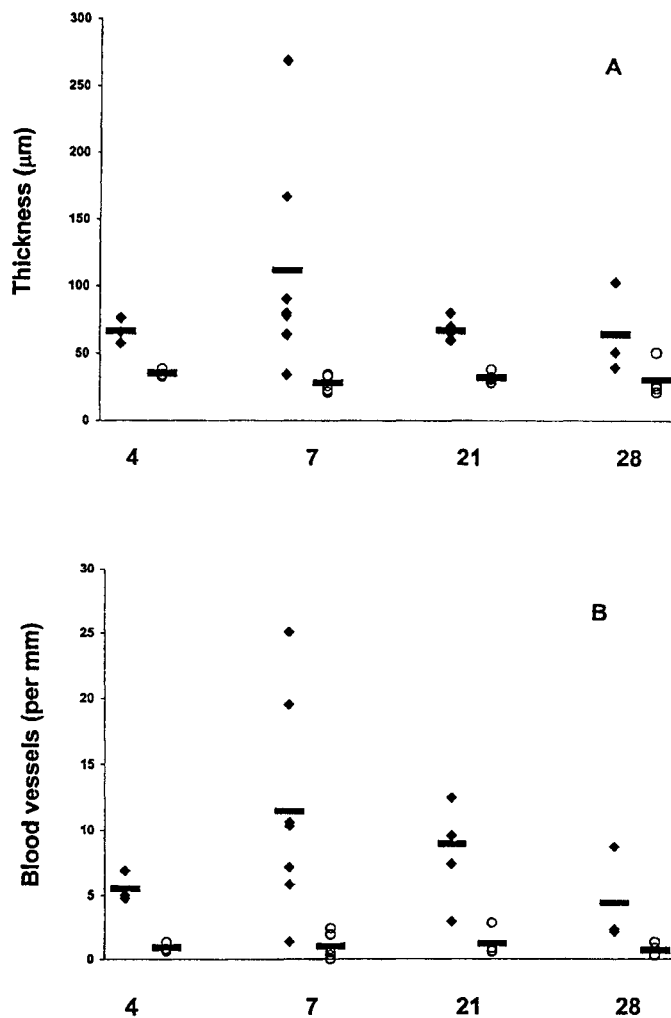


Figure 5. AdTGF β 1-mediated changes in membrane thickness and vascularity, as assessed by quantitative immunohistochemical analysis of factor VIII-stained sections of parietal peritoneum. All data points are presented with the black bars representing mean values. (A) Thickness of the peritoneal membrane in AdTGF β 1-treated animals (\blacklozenge) and control (AdDL70-treated) animals (\circ). (B) Number of blood vessels counted per millimeter length of peritoneal membrane in AdTGF β 1-treated animals (\blacklozenge) and control animals (\circ). Ten to 12 fields were counted for each section. $P < 0.05$ for comparison between AdTGF β 1- and AdDL70-treated animals (days 4 to 21) for both thickness and vessels per millimeter.

tion-deficient adenoviral vectors. In comparisons of control virus- and PBS-treated animals, we observed no demonstrable, acute, virus-induced effects on morphologic features or peritoneal membrane transport at the vector dose used. In comparisons of the AdDL70 control and AdTGF β 1 virus, we observed no difference in markers of acute inflammation after 4 d.

There are similarities between our animal model of AdTGF β 1-induced peritoneal fibrosis and the peritoneum of patients undergoing long-term dialysis. As has been observed in biopsy samples from patients undergoing long-term peritoneal dialysis (11,12,14), we noted increased thickness of the submesothelial collagenous zone, with increased α -SMA staining and increased vascularity. The solute transport properties

of the AdTGF β 1-treated peritoneum also mimic the changes observed for long-term dialysis patients, with increased transport of glucose and albumin and decreased ultrafiltration capacity. Some of the changes we observed in our model were transitory. The early fibrocellular proliferation and increased vascularization declined after 7 and 21 d, respectively. The changes in membrane transport and ultrafiltration also normalized with the decline in vascularization.

Variability was noted in the histologic changes observed in the peritoneum of animals treated with AdTGF β 1. These changes were fairly uniform among sections from individual animals, suggesting that inadequate sampling did not affect these results. It is more likely that there was variability in the level of TGF- β 1 expressed in each animal, despite delivery of similar adenoviral doses, and this range of exposure explains the variable histologic presentation.

Previous studies in the lung with AdTGF β 1 demonstrated a progressive fibrogenic response (35). In the peritoneum, there seems to be sustained expansion of the extracellular matrix, as indicated by both a sustained increase in the thickness of the submesothelial collagenous zone and increased hydroxyproline contents in the mesenteric tissue. Our findings are therefore in agreement with a maintained “fibrotic” response to transient overexpression of TGF- β 1. Of importance is the identified dissociation between “fibrosis,” as indicated by extracellular matrix deposition, and the vascular component.

There is a narrow concentration range in which TGF- β 1 produces effects in the peritoneum. To identify a suitable dose of AdTGF β 1 to use in this study, we initially performed a dose-response study. We observed that doses of AdTGF β 1 of $<1 \times 10^9$ plaque-forming units led to transient TGF- β 1 levels of <1000 pg/ml, with no histologic or functional effects. At higher doses, excessive morbidity resulted from a striking early fibroproliferative response, which led to stricture and bowel obstruction. This resistance to the effects of TGF- β 1 could be explained by the upregulation of natural inhibitors of TGF- β 1, such as the proteoglycan decorin, or by inhibitors of the fi-

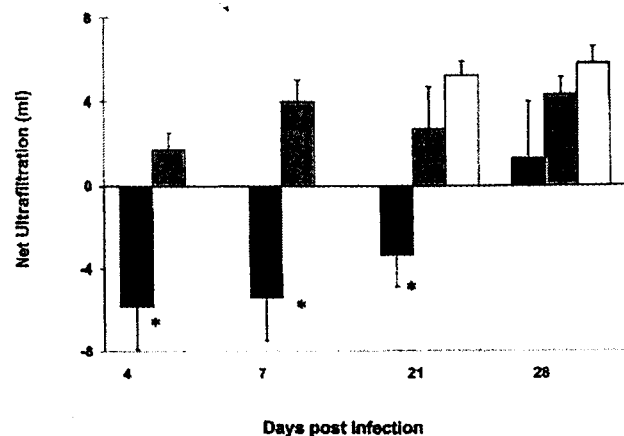


Figure 6. Net ultrafiltration after adenovirus infection. AdTGF β 1 treatment led to temporary ultrafiltration failure. AdTGF β 1-treated animals (\blacksquare) were compared with AdDL70-treated (\square) and PBS-treated (\square) animals. $*P < 0.05$.

Table 2. Transport properties of the peritoneal membrane after adenovirus-mediated gene transfer of TGF- β 1^a

Day	Group	Creatinine Clearance (ml/kg)	Glucose Transport (mmol/g)	Albumin Clearance (ml/kg)	Creatinine (d/p)	Glucose (d/d ₀)	Albumin (d/p) (10 ⁻³)
4	AdTGF β 1	32 \pm 11	9.8 \pm 0.2	4.7 \pm 2.0	0.59 \pm 0.05 ^b	0.08 \pm 0.01 ^c	65.5 \pm 59
	AdDL70	42 \pm 2	7.4 \pm 0.7	1.1 \pm 1.3	0.43 \pm 0.11	0.29 \pm 0.03	21.4 \pm 26.0
	PBS	ND	ND	ND	0.22 \pm 0.08	0.29 \pm 0.02	0
7	AdTGF β 1	46 \pm 13	8.9 \pm 1.8	10 \pm 7	0.68 \pm 0.17 ^c	0.07 \pm 0.01 ^c	245 \pm 117 ^c
	AdDL70	41 \pm 4	6.2 \pm 0.3	2.1 \pm 0.2	0.41 \pm 0.09	0.25 \pm 0.10	13.7 \pm 24.6
	PBS	ND	ND	ND	0.44 \pm 0.20	0.27 \pm 0.03	0
14	AdTGF β 1	ND	ND	ND	0.42 \pm 0.06	0.12 \pm 0.06 ^b	57.6 \pm 44.4 ^b
	AdDL70	ND	ND	ND	0.47 \pm 0.10	0.25 \pm 0.09	12.8 \pm 28.6
	PBS	ND	ND	ND	0.43 \pm 0.04	0.30 \pm 0.03	0
21	AdTGF β 1	38 \pm 5	8.3 \pm 0.7	2.6 \pm 2.3	0.46 \pm 0.09	0.20 \pm 0.06 ^b	43.7 \pm 58.0
	AdDL70	46 \pm 10	7.4 \pm 1.1	2.3 \pm 1.8	0.38 \pm 0.09	0.26 \pm 0.06	17.9 \pm 17.2
	PBS	49 \pm 2	6.4 \pm 0.8	0	0.42 \pm 0.07	0.24 \pm 0.08	0
28	AdTGF β 1	48 \pm 15	7.3 \pm 0.7	0.6 \pm 1.3	0.40 \pm 0.09	0.27 \pm 0.02	9.1 \pm 20.3
	AdDL70	49 \pm 7	6.8 \pm 0.4	1.0 \pm 1.3	0.39 \pm 0.04	0.28 \pm 0.03	7.2 \pm 11.2
	PBS	52 \pm 9	6.6 \pm 0.4	0	0.40 \pm 0.03	0.27 \pm 0.02	0

^a All data are mean \pm SD. d/p, dialysate/plasma concentration ratio; d/d₀, 4-h/initial concentration ratio; TGF- β 1, transforming growth factor- β 1; ND, not determined.

^b $P < 0.02$ by t test for comparison between AdTGF β 1 and AdDL70 groups.

^c $P < 0.001$.

brotic response, such as the metalloproteinases. The resolution of the vascular component of the fibrotic response suggests that there may be upregulation of natural inhibitors to angiogenesis in response to the transient high levels of TGF- β or a natural decrease in vascularity with the withdrawal of VEGF. If the peritoneum can recover from a transient insult, then perhaps prolonged or repeated exposure to profibrotic cytokines (via recurrent infection or daily dialysis) may induce more lasting changes in the structure and function of the peritoneal membrane. Chronic rat models of peritoneal dialysis have indeed confirmed this hypothesis (44,45).

The time course of neovascularization coincides with the changes in peritoneal membrane transport. This suggests that alteration of blood vessels, through either increased vascular surface area (represented by the number of vessels per unit length of peritoneal membrane in cross-sections) or increased vascular permeability, is responsible for the changes in peritoneal membrane transport we observed. The increased albumin transport observed on day 4, at a time when the dialysate concentration of VEGF was very high and vascularization was only moderately increased, suggests a direct permeability response of existing blood vessels to VEGF at earlier time points. The ongoing ultrafiltration defect at day 21, *i.e.*, a time when VEGF levels had returned to baseline values and blood vessel counts were still above control levels, supports a role for increased vascularization in the demonstrated changes in peritoneal membrane function.

On the basis of our limited *in vitro* observations and recently reported observations that exposure of rat primary mesothelial cells to TGF- β led to an increase in VEGF protein and mRNA production (32), we suggest that mesothelial cells may be responsible for *in vivo* angiogenic cytokine production after

TGF- β 1 exposure. Other resident peritoneal cell populations, such as macrophages, fibroblasts, and endothelial cells, may also participate in this response.

We have established a model of TGF- β 1-mediated peritoneal fibrosis in rats. This model can be used to elucidate the molecular mechanisms and pathobiologic features of peritoneal fibrosis and the consequent effects on solute transport. Our model suggests that TGF- β 1 induces structural alterations in the peritoneum, which lead to increased solute permeability and decreased ultrafiltration. We demonstrate that VEGF expression is dramatically increased in peritoneal membranes exposed to TGF- β 1 and mesothelial cells likely have a role in this process. Changes in peritoneal membrane function seem to be correlated with exposure to VEGF, and the changes observed resolve as the neovascularization declines. Our observation that angiogenic factors play an important role may lead to potential therapeutic interventions for the prevention or treatment of peritoneal membrane failure among patients undergoing long-term peritoneal dialysis. Effective adenoviral gene therapy using antiangiogenic genes such as angiostatin has been demonstrated in tumor models (46). Because of their potential use in cancer therapy, new antiangiogenic compounds are being developed and tested (47,48), and these compounds may be useful in protecting the peritoneal membrane during chronic peritoneal dialysis.

Acknowledgments

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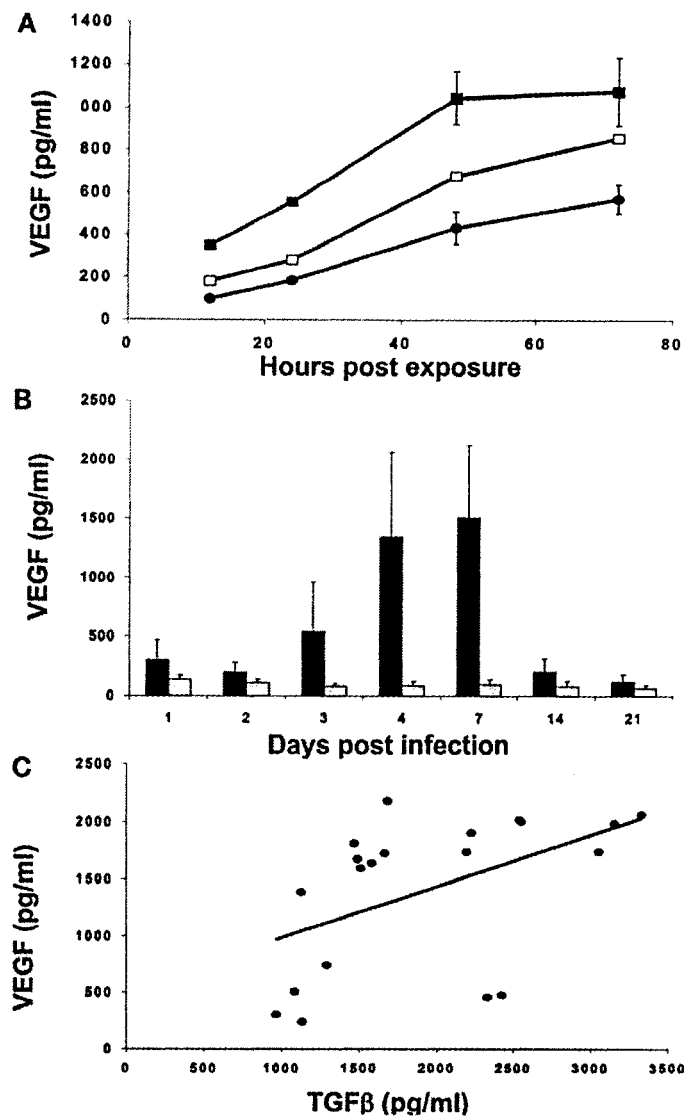


Figure 7. Relationship between TGF- β 1 and vascular endothelial growth factor (VEGF). (A) *In vitro* expression of VEGF in mesothelial cell culture supernatants after exposure for 12 h to 10 ng/ml recombinant TGF- β 1 (■), 1 ng/ml TGF- β (□), or PBS (●). Significant differences ($P < 0.01$) were observed at 48 and 72 h (10 ng/ml versus control). (B) Peritoneal dialysate VEGF concentrations after exposure to AdTGF β 1 (■) or AdDL70 (control) (□). Significant differences ($P < 0.001$) were observed for days 4, 7 ($P < 0.001$), and 14 ($P = 0.03$). Values shown are mean \pm SD ($n = 3$ to 5 for days 1 to 3 and $n = 10$ to 21 for days 4 to 14). (C) Correlation between day 4 TGF- β peritoneal dialysate concentrations and day 4 VEGF concentrations ($r^2 = 0.240$, $P = 0.03$).

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CHAPTER 3: INFLAMMATORY CYTOKINES, ANGIOGENESIS, AND FIBROSIS IN THE RAT PERITONEUM

This paper is published in the American Journal of Pathology 2002; 160: 2285-2294. We used similar techniques as was used in the AdTGF β 1 model described above. This time, we looked specifically at two inflammatory cytokines - TNF α and IL-1 β . We had some suggestions from other work that IL-1 β may be a more fibrogenic cytokine than TNF α ¹⁶⁶. It was therefore of some interest to see the effect on the peritoneal function and angiogenesis.

IL-1 β treated animals had a more potent inflammatory reaction with prolonged weight loss and increased peritoneal effluent markers of acute inflammation. One interesting finding was that adenovirus mediated gene transfer of TNF α led to an up-regulation of IL-1 β , but IL-1 β did not induce any expression of TNF α . This confirms that TNF α is an upstream, early inflammatory mediator.

TNF α induced a potent, but very transient angiogenesis with ultrafiltration dysfunction. There was also a transient increase in peritoneal fibrosis measured by hydroxyproline concentration in peritoneal tissues. IL-1 β , in contrast, induced a very prolonged angiogenesis, ultrafiltration dysfunction, and peritoneal fibrosis. In trying to understand this, we identified a prolonged significant increased expression of TIMP-1 and PAI-1 mRNA expression in the peritoneal tissues of animals exposed to IL-1 β . We felt that this inhibition of metalloproteinases and collagenolysis may explain the persisting fibrosis. Of note was the observation that TGF β 1 mRNA levels were very similar between TNF α and IL-1 β treated animals.

This study identified the potential for angiogenesis, even very transient angiogenesis, to be a mechanism of acute ultrafiltration failure seen in PD patients who develop peritonitis. It also provided a potential link, through IL-1 β , between peritonitis and longer term changes in fibrosis and ultrafiltration dysfunction.

The AdIL-1 β was created by Dr. Daniel Anthony from the University of Southampton. The AdTNF α was created by Dr. Robert Marr who was previously in Dr. Frank Graham's lab.

I developed the concept for this paper. I designed the experiments with some input from Dr. Gauldie and Dr. Hoff. I designed the experiments. All work was carried out by me, with some technical support from Martin Kolb and Lisa Yu. I analyzed the data and wrote the paper. The manuscript was reviewed by Drs. Gauldie, Hoff, and Cliff Holmes.

Animal Model

Inflammatory Cytokines, Angiogenesis, and Fibrosis in the Rat Peritoneum

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Peritonitis, a common complication of peritoneal dialysis, is followed by acute changes in the function of the peritoneum. The role of inflammatory cytokines in these processes is not clearly identified. We used adenoviral-mediated gene transfer to transiently overexpress interleukin (IL)-1 β (AdIL-1 β) or tumor necrosis factor (TNF)- α (AdTNF- α) in the rat peritoneum then used a modified equilibrium test to study the histological and functional changes. Overexpression of IL-1 β or TNF- α led to an acute inflammatory response. Both inflammatory cytokines induced an early expression of the angiogenic cytokine, vascular endothelial growth factor, along with increased expression of the profibrotic cytokine, transforming growth factor- β 1, along with fibronectin expression and collagen deposition in peritoneal tissues. Both inflammatory cytokines induced angiogenesis, increased solute permeability, and ultrafiltration dysfunction at earlier time points. Changes in structure and function seen in AdTNF- α -treated animals returned to normal by 21 days after infection, whereas AdIL-1 β -treated animals had persistently increased vasculature with submesothelial thickening and fibrosis. This was associated with up-regulation TIMP-1. TNF- α or IL-1 β both induce acute changes in the peritoneum that mimic those seen in peritoneal dialysis patients who experience an episode of peritonitis. These functional changes were associated with early angiogenesis that resolved rapidly after exposure to TNF- α . IL-1 β exposure, however, led to a different response with sustained vascularization and fibrosis. IL-1 β inhibition may be a therapeutic goal in acute

peritonitis to prevent peritoneal damage. (*Am J Pathol* 2002, 160:2285–2294)

Peritonitis is a common complication of peritoneal dialysis.¹ The response of the peritoneum to infective organisms involves the inflammatory cytokines and the interaction between resident cell populations: macrophages, mesothelial cells, and fibroblasts.^{2,3} The earliest response involves tumor necrosis factor (TNF)- α and interleukin (IL)-1³ derived from macrophages in response to bacterial products.⁴ What follows is a complex interaction between inflammatory cytokines, such as IL-6, chemokines and subsequent leukocyte transmigration, prostaglandins, nitric oxide, and adhesion molecules. There is also an anti-inflammatory response consisting of soluble receptors to TNF- α ,⁵ IL-1 receptor antagonist,⁶ and repair molecules such as hyaluronan.⁷

In the first month after peritonitis, patients develop an acute dysfunction of the peritoneum as a dialysis membrane.⁸ This is characteristically associated with increased transport of small molecular weight solutes and ultrafiltration failure. The mechanism for this is not well established, but vascular effects of nitric oxide⁹ or prostaglandins¹⁰ have been hypothesized to play a role.

Recurrent peritonitis has been shown, in some studies, to be associated with long-term peritoneal membrane changes.^{11,12} As in the acute changes after peritonitis, long-term changes are associated with increased solute transport and decreased ultrafiltration. In this setting, human biopsy studies^{13–15} and animal experiments¹⁶ have identified an increase in the peritoneal-associated vasculature, which seems to be the primary cause of increased solute transport. A fibrogenic response and

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associated vasculopathy have also been identified in patients on long-term peritoneal dialysis.¹⁷

The association between peritoneal inflammation, angiogenesis, and fibrosis is not clear. IL-1 β has been identified *in vivo* as having strong fibrogenic properties through up-regulation of transforming growth factor (TGF)- β .¹⁸ This has been confirmed in *in vitro* mesothelial cell culture work.¹⁹ Both IL-1 β and TNF- α have angiogenic properties as has been demonstrated in studies using rat mesenteric window assays^{20,21} and TNF- α has been associated with up-regulation of the angiogenic cytokine vascular endothelial growth factor (VEGF) in cell culture.²²

We have previously described the use of adenovirus-mediated gene transfer of cytokines and growth factors as an effective tool in elucidating functional and morphological changes in the peritoneum.²³ In the following experiments, we compared the effects of transient overexpression of TNF- α or IL-1 β on the peritoneum of rats using adenovirus-mediated gene transfer (AdTNF- α or AdIL-1 β) and controlled with a null adenovirus (AdDL70). Four days after infection with either AdTNF- α or AdIL-1 β we saw increased vascularity of the peritoneum, increased solute transport, and decreased ultrafiltration. By day 21, the animals treated with AdTNF- α returned to normal peritoneal morphology and function, but the AdIL-1 β -treated animals remained quite abnormal with peritoneal fibrosis and persistent vascularization.

Materials and Methods

Adenovirus

The construction of the adenovirus vectors AdIL-1 β ,¹⁸ AdTNF- α ,²⁴ and AdDL70²⁵ have been previously described. Adenovirus preparations were purified by CsCl gradient centrifugation and PD-10 Sephadex chromatography (Amersham Pharmacia, Baie d'Urfe, Canada) and plaque-titered on 293 cells as previously described.²⁶

Animals

All animal studies were performed according to the Canadian Council on Animal Care Guidelines. Three groups of female Sprague-Dawley rats (Harlan, Indianapolis, IN), 200 to 250 g (corresponding to 6 to 8 weeks of age²⁷) were studied. The first group ($n = 27$) received an intraperitoneal injection of AdIL-1 β . The second group ($n = 20$) received AdTNF- α and the third group ($n = 28$) received a null control virus (AdDL70). All were administered adenovirus at a dose of 1.5 to 2.0×10^9 plaque-forming units/ml diluted to $100 \mu\text{l}$ in phosphate-buffered saline on day 0. Animals did not receive anti-inflammatory agents during the experimental protocol. Animals from each group were sacrificed on days 4, 7, 21, and 28 after adenovirus administration. Before sacrifice, 0.09 ml/g of 2.5% Dianeal (Baxter Health Care, McGaw Park, IL) was administered intraperitoneally. Four hours later, the peritoneum was opened and the entire fluid content removed for accurate ultrafiltration measurement. Net ultrafiltration

was the volume of fluid removed after 4 hours minus the volume of fluid administered. Blood samples were drawn and the entire anterior abdominal wall was removed after skin and subcutaneous tissue was removed. The lower portion of this tissue was stored in formalin and the upper portion taken for RNA extraction. Mesenteric tissue was taken and frozen in liquid nitrogen.

Whole blood was centrifuged at 5000 rpm for 10 minutes and the serum removed. Peritoneal fluid samples were centrifuged at 1500 rpm for 5 minutes. Samples were analyzed on a Hitachi 917 automated chemistry analyzer (Roche Diagnostics, Laval, Canada) for glucose and albumin. Mass transport of glucose out of the peritoneum was calculated as (initial dialysate glucose \times initial volume) - (final dialysate glucose \times final volume). Albumin clearance was calculated as mass transport divided by the serum solute concentration. All values were corrected for animal weight at sacrifice.

Cell number in the peritoneal fluid was counted with a hemocytometer. The fluid was plated to a glass slide by cytopsin (Shandon Inc., Pittsburgh, PA) and stained (Biochemical Science Inc., Swedesboro, NJ). Cell differential was counted on at least 300 cells.

Histology

Tissue samples at sacrifice from the lower anterior abdominal wall or omentum were taken and fixed in a sufficient amount of 4% phosphate-buffered formaldehyde for 24 hours. The tissue samples were then paraffin-processed, embedded, and $5\text{-}\mu\text{m}$ sections cut. Cut sections were then stained for Masson's trichrome and immunohistochemistry was performed with antibodies to Factor VIII-von Willebrand factor (vWF) antigen (DAKO Corp., Carpinteria, CA). Negative control sections were run in parallel with nonimmune mouse or rabbit serum. All sections were deparaffinized in xylene followed by 100% ethanol and then placed in a methanol H_2O_2 solution for 30 minutes to block endogenous peroxidase activity. After hydration to water with graded alcohols, the sections were placed in 0.05 mol/L Tris-buffered saline, pH 7.6. The sections were digested with 0.05% Pronase (Sigma Chemical Co., St. Louis, MO) in Tris-buffered saline with calcium chloride for 17 minutes at room temperature then blocked in 5% normal goat serum followed by a 1-hour incubation in the 1:500 rabbit anti-human factor VIII in 1% normal goat serum. Sections were then incubated in a prediluted kit of a biotinylated goat anti-rabbit followed by a streptavidin/peroxidase conjugate (Zymed Labs, San Francisco, CA) as per the manufacturer's instructions. Immunohistochemistry incubations were performed at room temperature and sections were washed in between incubations 3×5 minutes with 0.05 mol/L Tris-buffered saline, pH 7.6, except before the addition of primary antibody. Sections were rinsed in 0.05 mol/L acetate buffer, pH 5.0, before development in an 3-amino-9-ethylcarbazole (AEC) chromogen substrate for 15 minutes. Sections were counterstained in Mayer's hematoxylin for 2 minutes before mounting with glycerin gelatin.

Quantitative Immunohistochemistry

Low-power fields from factor VIII-immunostained sections of the anterior abdominal wall were digitized using a Leica DMR microscope (Leica Microsystems, Wetzlar, Germany). All sections were analyzed by the same image-processing algorithm using Leica Qwin Image Processing Software (Leica Imaging Systems, Cambridge, England). Results are reported as number of vessels/mm² of peritoneal tissue. We were able to estimate the total vessel cross-section area in each digitized image and we could therefore calculate an average cross-sectional area per vessel for each slide analyzed.

Hydroxyproline Assay

A portion of mesentery was taken and frozen for a hydroxyproline assay, modified from Woessner's method.²⁸ Tissues were weighed, homogenized in water, centrifuged at 1000 rpm for 5 minutes, and the superficial fatty material removed by vacuum suction. Solid material was precipitated with trichloroacetic acid with centrifugation at 1500 rpm for 15 minutes at 4°C. Samples were hydrolyzed overnight in 6 N HCl at 110°C. Hydroxyproline content is quantified by Erlich's reagent (Sigma) and assayed by measuring the optical density at 557 nm. A hydroxyproline standard sample (Sigma) was used to create a standard curve.

Cytokine Analysis

Peritoneal fluid taken at sacrifice was analyzed for the following cytokines using enzyme-linked immunosorbent assay (ELISA) (all R&D Systems, Minneapolis, MN) as directed by the manufacturer: human TGF- β (cross-reactive with rat), murine VEGF (cross-reactive with rat), rat IL-6, rat TNF- α (cross-reactive with murine), human IL-1 β (not cross-reactive with rat), and murine TNF-soluble receptor II (TNFsrII, cross-reactive with rat). To measure total TGF- β , samples were first activated using 1 mol/L HCl for 10 minutes then normalized with 1 N NaOH to dissociate TGF- β from its latency-associated binding protein.

Frozen omental tissue was taken and homogenized in Trizol reagent (Life Technologies, Burlington, Ontario, Canada). Protein was extracted from the phenol layer after centrifugation according to the manufacturer's protocol. Total protein in this extract was assayed using a standard protein assay method (BioRad DC Protein Assay; BioRad Laboratories, Mississauga, Canada). Equal quantities of protein were then assayed using murine VEGF ELISA (R&D Systems).

RNase Protection Assay

The peritoneal surfaces of the anterior abdominal wall sections were immersed for 15 minutes in Trizol reagent. The parietal peritoneum was gently scraped and the Trizol collected and processed according to the manufacturer's instruction for isolation of RNA. The concentra-

tion of RNA resuspended in RNase-free water was measured by optical density at 260 nm. RNA (8 μ g) was then hybridized overnight with a custom probe set (Pharmin-gen, Mississauga, ON) labeled with α -³²P-UTP (New England Nuclear, Boston, MA). The custom probe set contains different length probes for fibronectin, TGF- β 1, tissue inhibitor of metalloproteinase-1 (TIMP-1), and housekeeping genes GAPDH and L32 for loading control. The hybridized samples were extracted using phenol/chloroform and acetate precipitation and then washed with ethanol. The extracted bound RNA was then run on a 5% polyacrylamide gel, transferred to blotting paper, dried, and exposed for 4 days to film (Eastman Kodak, Rochester, NY).

The images obtained were digitized and analyzed for band density using Scion Image software (Scion Corp, Frederick, MD). Densities were standardized to L32.

Statistics

Data are presented \pm SEM unless otherwise noted. Comparison between groups was made by *t*-test. We used regression analysis to compare the cytokine concentration on day 4 in the peritoneal fluid and subsequent vascularization of the peritoneal membrane and combined all three treatment groups (AdIL-1 β , AdTNF- α , AdDL70) in these results.

Results

Adenoviral Response

In previous work, we have identified an early (48 hour) inflammatory response to AdDL70 in similar dosage used in these experiments.²⁹ There were no adenoviral effects identified by 4 days,²⁹ the earliest time point used in our experiments. We therefore did not include a nonadenoviral control group in these experiments.

We have previously shown that adenovirus is effectively taken up, and the transgene product expressed, by mesothelial cells in the peritoneum.²³ The peak of expression is 4 to 7 days after infection, and the total duration of expression is 10 to 14 days. In these experiments, we confirmed the high levels of expression of the transgene product by analysis of the peritoneal dialysis fluid with the appropriate ELISA. Four days after infection peritoneal fluid was taken after a 4-hour dwell and analyzed by ELISA. Animals treated with AdIL-1 β had 326 \pm 89 pg/ml of human IL-1 β and animals treated with AdTNF- α had 2076 \pm 1131 pg/ml of rodent TNF- α measured in the peritoneal fluid (Table 1).

Inflammatory Response

The inflammatory response after intraperitoneal administration of AdTNF- α , AdIL-1 β , or control adenovirus was measured from various markers in the peritoneal fluid and the results are shown in Table 2. The cellular response to overexpression of IL-1 β and TNF- α was dramatic with a significant increase in the total number of cells and a

Table 1. Cytokine Concentration in the Peritoneal Fluid and Tissue 4, 7, and 21 Days after Infection with Control Adenovirus, AdTNF- α , or AdIL-1 β

	Day 4			Day 7			Day 21		
	AdDL70	AdTNF- α	AdIL-1 β	AdDL70	AdTNF- α	AdIL-1 β	AdDL70	AdTNF- α	AdIL-1 β
Human IL-1 β (pg/ml)	<10	<10	326 \pm 89* [†]	<10	<10	77 \pm 105	<10	<10	<10
Rat IL1 β (pg/ml)	32 \pm 10	302 \pm 255*	310 \pm 186*	31 \pm 10	114 \pm 92	638 \pm 800	46 \pm 21	28 \pm 2	251 \pm 301
TNF- α (pg/ml)	69 \pm 8	2076 \pm 1131*	69 \pm 7 [†]	58 \pm 14	365 \pm 482*	83 \pm 12 [†]	nd	66 \pm 11	69 \pm 9
Total TGF- β (pg/ml)	131 \pm 80	186 \pm 82	756 \pm 441* [†]	88 \pm 50	306 \pm 137*	1056 \pm 610* [†]	63 \pm 2	64 \pm 18	166 \pm 101
VEGF (pg/ml)	75 \pm 44	172 \pm 130*	210 \pm 173*	85 \pm 49	205 \pm 133*	262 \pm 157*	63 \pm 24	30 \pm 4	47 \pm 17
Tissue VEGF (pg/ml)	43 \pm 10	72 \pm 40	76 \pm 16	47 \pm 14	45 \pm 21	78 \pm 8*	nd	nd	nd

**P* < 0.05 compared with AdDL70 control.
[†]*P* < 0.05 compared with AdTNF- α .
 Data shown \pm SD.
 nd, No data.

disproportionate increase in neutrophils measured in the peritoneal fluid. There was also an associated weight loss that was more dramatic in the AdIL-1 β -treated animals compared to control adenovirus-treated animals. Rodent IL-6 was measured by ELISA in peritoneal fluid and was elevated after exposure to IL-1 β or TNF- α (Table 1). Finally, we measured rodent TNFsrlI by ELISA in peritoneal fluid and this was significantly elevated 4 and 7 days after infection with AdIL-1 β and AdTNF- α (Table 1).

We also noted a quantitative difference between the inflammatory response to AdIL-1 β or AdTNF- α . Specifically the response to transient overexpression of IL-1 β was more intense, as measured by total cell count and percentage of neutrophils, especially 7 days after infection. Also, two animals in the AdIL-1 β group became very ill in the first week after treatment and required euthanasia. The response to IL-1 β was more prolonged, with levels of TNFsrlI remaining significantly elevated in AdIL-1 β -treated animals 21 days after infection.

Cytokine Response after Transient Overexpression of TNF- α or IL-1 β

We analyzed the peritoneal fluid after a 4-hour dwell for rat IL-1 β , TNF- α , TGF- β , and VEGF using ELISA (Table 1). Both AdTNF- α and AdIL-1 β treatment induced the production of endogenous IL-1 β measured using a rodent-specific assay. Interestingly, AdIL-1 β -treated animals did not show an increase in peritoneal concentration of TNF- α . Overexpression of both TNF- α and IL-1 β in-

duced a significant increase in the peritoneal fluid concentration of total TGF- β 1 and VEGF compared to AdDL70-treated animals 7 days after infection. We identified a threefold increase in peritoneal concentration of TGF- β 1 7 days after AdIL-1 β treatment compared to AdTNF- α . Increases in peritoneal VEGF concentrations were similar after exposure to IL-1 β and TNF- α .

To confirm that VEGF was present in the tissues after adenovirus infection, we isolated protein from omental homogenates and assayed for VEGF. We noted an increase in tissue VEGF that was significantly elevated 7 days after treatment with AdIL-1 β (Table 1).

We isolated RNA from parietal peritoneal tissue of animals 4, 7, and 21 days after receiving adenovirus and analyzed it using an RNase protection assay (Figure 1). Quantitative density analysis indexed to L32 and referenced to AdDL70-treated animals revealed that at 4 and 7 days after infection, both TNF- α and IL-1 β led to an increase in mRNA signal for fibronectin and TGF- β 1. These changes declined to baseline by day 21 except for a small persistent elevation in fibronectin mRNA in AdIL-1 β -treated animals at day 21. Most striking was the progressive increase in TIMP-1 mRNA expression seen in AdIL-1 β -treated animals at day 21 (Figure 1).

Changes in Peritoneal Membrane Morphology

Transient overexpression of both IL-1 β and TNF- α led to early, significant histological changes in the parietal peritoneum compared to animals treated with the control

Table 2. Inflammatory Changes Measured 4 and 7 Days after Infection with Control Adenovirus, AdTNF- α , AdIL-1 β

	Day 4			Day 7		
	AdDL70	AdTNF- α	AdIL-1 β	AdDL70	AdTNF- α	AdIL-1 β
White blood cells (10 ⁴ /ml)	77 \pm 69	158 \pm 81*	952 \pm 275* [†]	132 \pm 164	262 \pm 148	763 \pm 541* [†]
Monocyte, %	19.4	15.3	8.0*	28.9	20.3	13.2
Macrophage, %	46.2	41.4	30.7	50.5	49.4	27.1
Neutrophil, %	4.2	31.7*	48.3*	3.5	25.9*	58.3* [†]
Eosinophil, %	27.4	9.8*	11.3*	19.4	3.8	1.1
Change in weight from day 0 (g)	1.3 \pm 5.3	-2.1 \pm 2.8	-5.6 \pm 4.1	2.8 \pm 6.1	2.2 \pm 3.9	-5.9 \pm 4.2
IL-6 (pg/ml)	59.1 \pm 0.3	988 \pm 857	1451 \pm 497	66 \pm 2	74 \pm 8	1788 \pm 746
TNFsrlI (pg/ml)	34 \pm 16	604 \pm 420*	751 \pm 364*	63 \pm 11	1156 \pm 275*	917 \pm 388*

**P* < 0.05 compared with AdDL70 control.
[†]*P* < 0.05 compared with AdTNF- α .
 Data shown \pm SD.

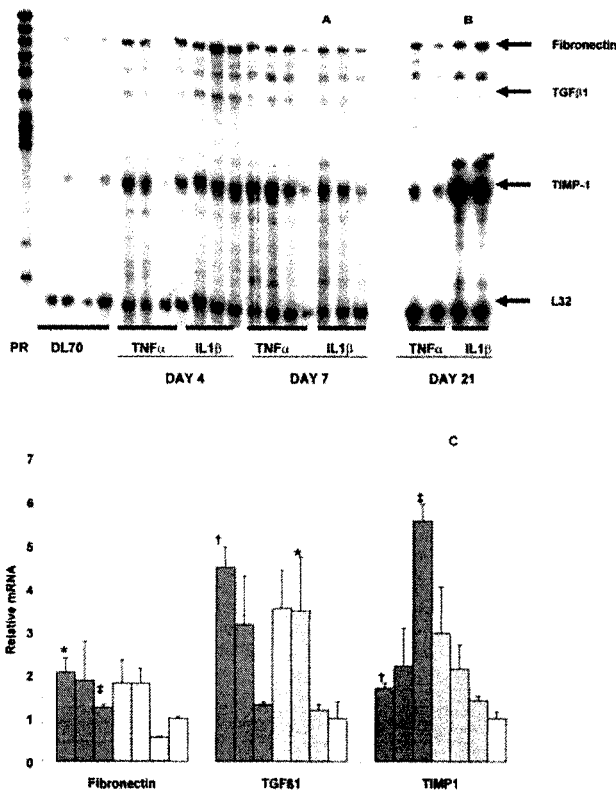


Figure 1. RNase protection assay. RNA was extracted from parietal peritoneal tissue and hybridized with a radiolabeled probe set. Samples were then RNase-digested, purified, and separated by polyacrylamide gel electrophoresis. **A:** PR is unprotected probe. Samples are from animals treated with AdDL70, AdTNF- α , or AdIL-1 β 4 and 7 days after infection. **B:** Day 21 samples run on a separate gel. **Arrows** indicate signal from hybridized mRNA for fibronectin, TGF- β 1, TIMP-1, and housekeeping gene L32 for loading control. **C:** Quantification of band density. Images were digitized and band density analyzed. Density was corrected for loading by L32 and then normalized to AdDL70-treated animals. **Dark gray bars**, AdIL-1 β -treated animals; **light gray bars**, AdTNF- α -treated animals; **open bars**, AdDL70-treated animals. The three bars for each treatment group represent days 4, 7, and 21 after adenovirus administration. AdDL70 animals are day 4 after adenovirus infection. There was a similar up-regulation of fibronectin and TGF- β 1 in AdIL-1 β - and AdTNF- α -treated animals at days 4 and 7. TIMP-1 was strongly expressed at day 21 in AdIL-1 β but not AdTNF- α animals. Data represents three to four animals at each data point. *, $P < 0.01$ compared with AdDL70 control; †, $P < 0.05$ compared with AdDL70; ‡, $P < 0.01$ for AdIL-1 β compared with AdTNF- α -treated animals.

adenovirus, AdDL70 (Figure 2). The mesothelial cells, which normally are flattened against the basement membrane, became rounded up with enlarged nuclei. The submesothelial zone became substantially thickened with inflammatory cells and edematous changes. These changes were evident in both parietal and visceral peritoneal tissue as seen in omental tissue 7 days after infection (Figure 3).

Seven days after infection with AdIL-1 β and AdTNF- α there was a dramatic angiogenesis that occurred in the peritoneum as evidenced by factor VIII-stained sections (Figure 2). Peritoneal-associated vessels were evaluated using digital image analysis from these sections (Figure 4). Overexpression of IL-1 β or TNF- α led to significantly different kinetics of angiogenesis. Animals treated with AdIL-1 β had increased peritoneal-associated blood vessels at each time point studied compared with AdDL70-treated animals. AdTNF- α -treated animals had an in-

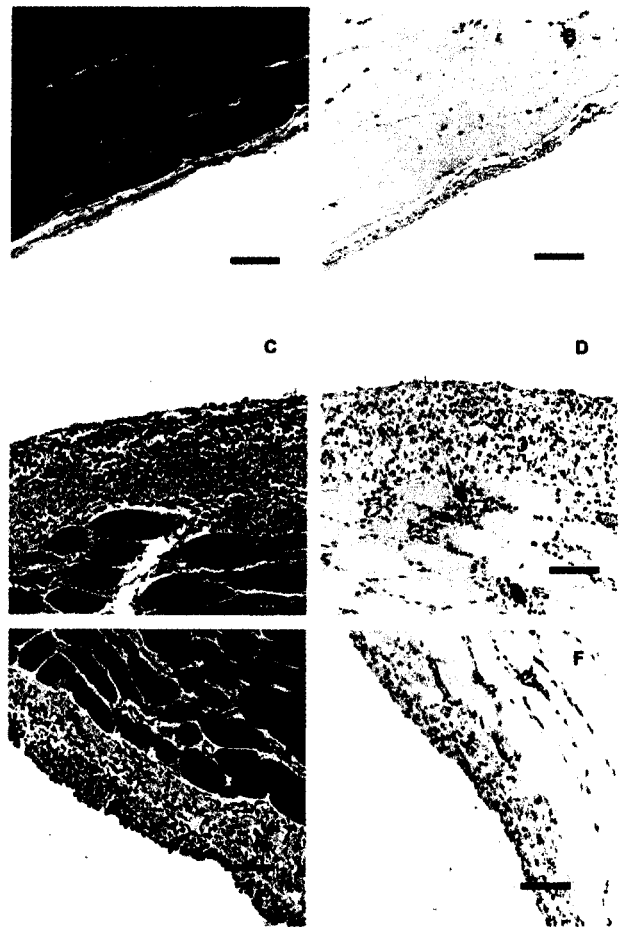


Figure 2. Histology of the anterior abdominal wall 7 days after infection with AdDL70 (**A** and **B**), AdIL-1 β (**C** and **D**), or AdTNF- α (**E** and **F**). After control adenovirus infection, the parietal peritoneum has normal histology (**A**) with thin submesothelial zone with few cells, intact mesothelial cell layer, and few blood vessels (**B**). Seven days after infection with AdIL-1 β or AdTNF- α , similar changes are observed. There is an increase in the submesothelial thickness, collagen deposition, and markedly increased cellularity (**C** and **E**). There is an associated increase in neovascularization (**D** and **F**). Scale bars, 100 μ m. Masson's trichrome (**A**, **C**, and **E**) and Factor VIII-vWF immunohistochemistry (**B**, **D**, and **F**).

crease in peritoneal vasculature that was significantly elevated over AdDL70-treated animals 4 and 7 days after adenovirus infection, but by day 21, the peritoneal-associated vasculature in the AdTNF- α -treated animals had returned to baseline.

There was an alteration in morphology of blood vessels after exposure to TNF- α and IL-1 β . We observed an increase in the average cross-sectional area of the blood vessels after exposure to inflammatory cytokines (IL-1 β , 146 ± 18 ; TNF- α , 136 ± 17 versus AdDL70, $73 \pm 23 \mu\text{m}^2$ 4 days after infection). This dilation of blood vessels persisted to 21 days after infection.

In both AdIL-1 β - and AdTNF- α -treated animals there was an early accumulation of collagen evident by day 7 (Figures 2 and 3). We confirmed the histological evidence of collagen accumulation by analyzing mesenteric tissue for hydroxyproline content. As shown in Figure 5, AdIL-1 β -treated animals had persistent elevation in hydroxyproline, and therefore, collagen content, of mesenteric tissue at each time point evaluated. AdTNF- α -

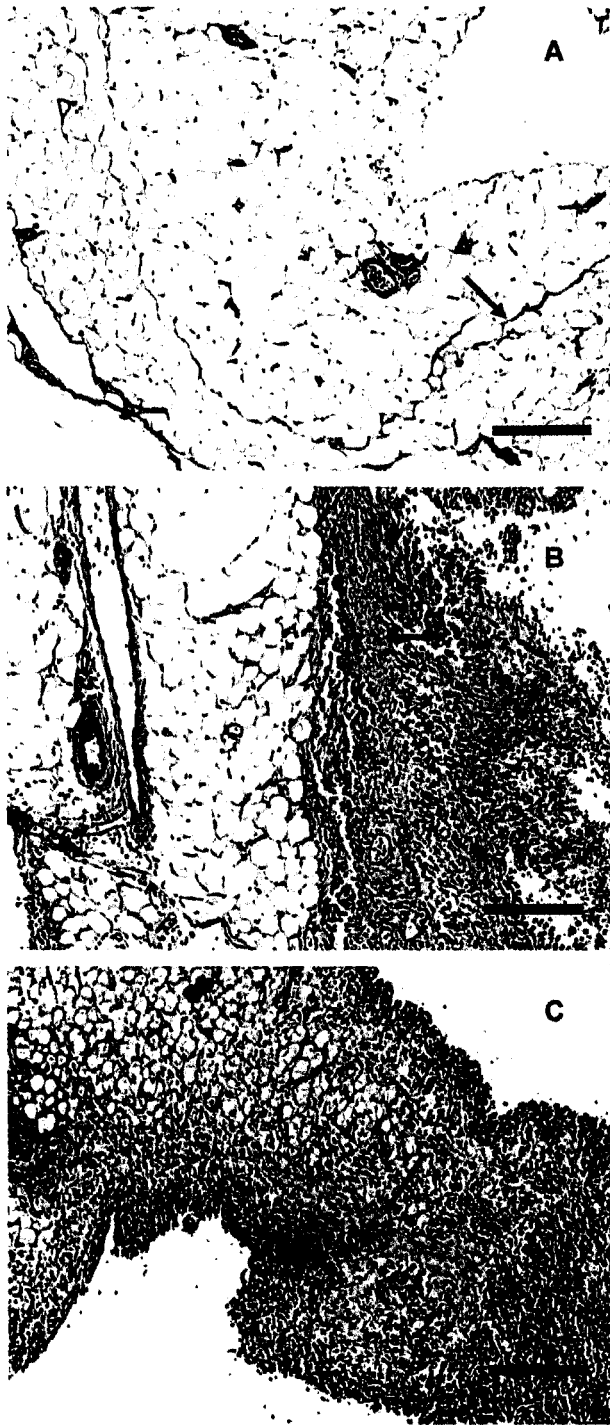


Figure 3. Histology of the visceral peritoneum (omentum) 7 days after infection with AdDL70 (A), AdIL-1 β (B), or AdTNF- α (C). AdIL-1 β and AdTNF- α -treated animals demonstrate increased cellular infiltration in omental tissue with collagen deposition (arrows). Scale bars, 200 μ m. Masson's trichrome.

treated animals showed a significant early accumulation of collagen but by day 21, this had resolved.

The most significant histological difference after exposure to TNF- α or IL-1 β can be seen in the day 21 samples (Figure 6). AdIL-1 β -treated animals had submesothelial thickening, now with collagen deposition (Figure 6A). There was persistent angiogenesis seen in the factor

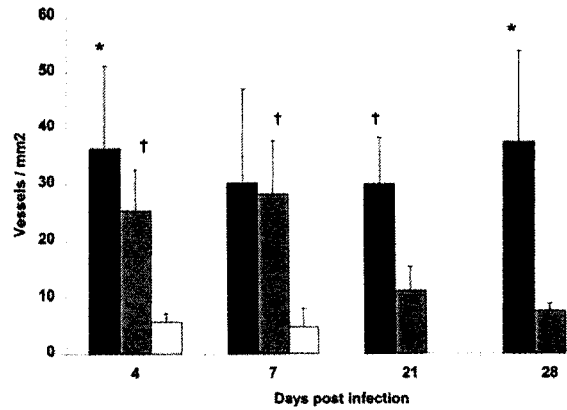


Figure 4. Vascular density in the submesothelial zone of the parietal peritoneum was measured using image analysis from digitized Factor VIII-vWF-stained sections taken from animals exposed to AdIL-1 β (black bars), AdTNF- α (gray bars), or control adenovirus (AdDL70, open bars). Animals treated with AdIL-1 β demonstrated increased vascular density in the submesothelial zone at all time points. In contrast, animals treated with AdTNF- α demonstrated a transient induction of angiogenesis that resolved by 21 days after infection. *, $P < 0.01$ compared with AdDL70-treated animals; †, $P < 0.05$ compared with AdDL70-treated animals. Data represents four to six animals for each data point.

VIII-stained section (Figure 6B). The AdTNF- α -treated animals had peritoneal histology indistinguishable from normal rat peritoneum 21 days after infection and demonstrated thin, compact submesothelial tissue and virtual absence of peritoneal-associated vessels (Figure 6, C and D).

Using regression analysis of all treatment groups combined, we could find a weak but significant correlation between dialysate VEGF concentration 4 days after adenovirus administration and subsequent vascularization (Figure 7A). Of note, there was a stronger correlation between TGF- β peritoneal fluid concentration on day 4 and subsequent vascularization (Figure 7B).

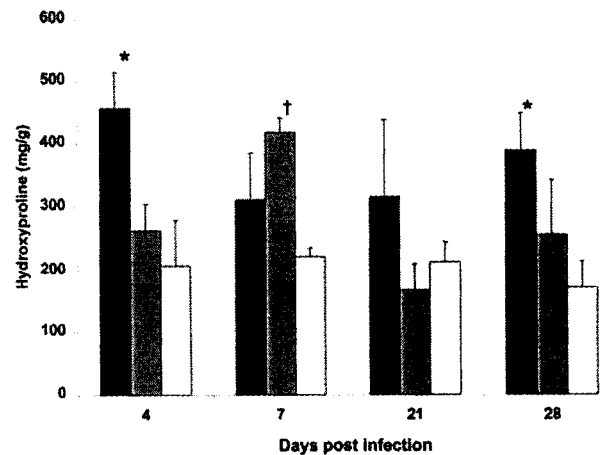


Figure 5. Hydroxyproline analysis of mesenteric tissue after infection with AdIL-1 β (black bars), AdTNF- α (gray bars), or control adenovirus (AdDL70, open bars). AdIL-1 β -treated animals demonstrated sustained fibrosis measured by hydroxyproline content of mesenteric tissue whereas TNF- α -treated animals demonstrated a transient accumulation of collagen by day 7 that disappeared by day 21. *, $P < 0.01$ compared with AdDL70; †, $P < 0.05$ compared with AdDL70. Data represents three to six animals at each time point.

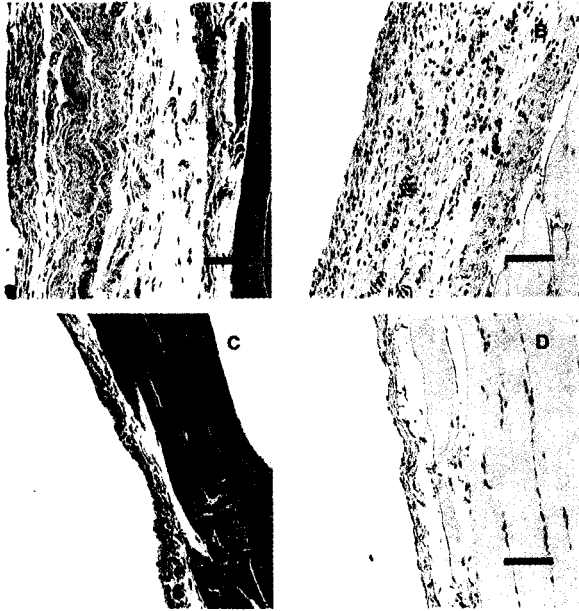


Figure 6. Histology of the anterior abdominal wall 21 days after infection with AdIL-1 β (A and B) or AdTNF- α (C and D). Animals infected with AdIL-1 β show persisting thickening of the submesothelial zone with collagen deposition (A) and persisting vascularization (B). After infection with AdTNF- α , the parietal peritoneum returned to normal histology (C) with absence of vascularization (D). Scale bars, 100 μ m. Masson's trichrome (A and C); Factor VIII-vWF immunohistochemistry (B and D).

Peritoneal Membrane Function

We studied the function of the peritoneum as a dialysis membrane using net ultrafiltration, glucose mass transport, and albumin clearance, after a 4-hour dwell of 2.5% Dianeal. The net ultrafiltration (Figure 8A) reveals a significant impairment 4 days after infection with AdTNF- α and AdIL-1 β compared with AdDL70-treated animals. The ultrafiltration function of the AdTNF- α -treated animals recovered by day 7, but remained impaired in IL-1 β -treated animals until day 28. The glucose transport (Figure 8B) showed a similar trend with early increased glucose transport after overexpression of both inflammatory cytokines, but with increased glucose transport persisting until day 28 for AdIL-1 β -treated animals.

Transient overexpression of TNF- α not only caused significant peritoneal vascularization by day 7, but albumin clearance suggests an increased permeability to macromolecules at this time point. The albumin clearance in AdTNF- α -treated animals, 7 days after adenovirus infection, was 30 ± 23 L/kg, which was significantly elevated over AdDL70-treated animals (3 ± 1 L/kg, $P = 0.03$), and elevated over AdIL-1 β -treated animals (7 ± 1 L/kg, $P = ns$).

Discussion

Adenovirus-mediated gene transfer is a versatile tool to analyze the effects of transient overexpression of cytokines on peritoneal morphology and function.²³ Both inflammatory cytokines IL-1 β and TNF- α induced an early acute inflammatory response not seen in control adeno-

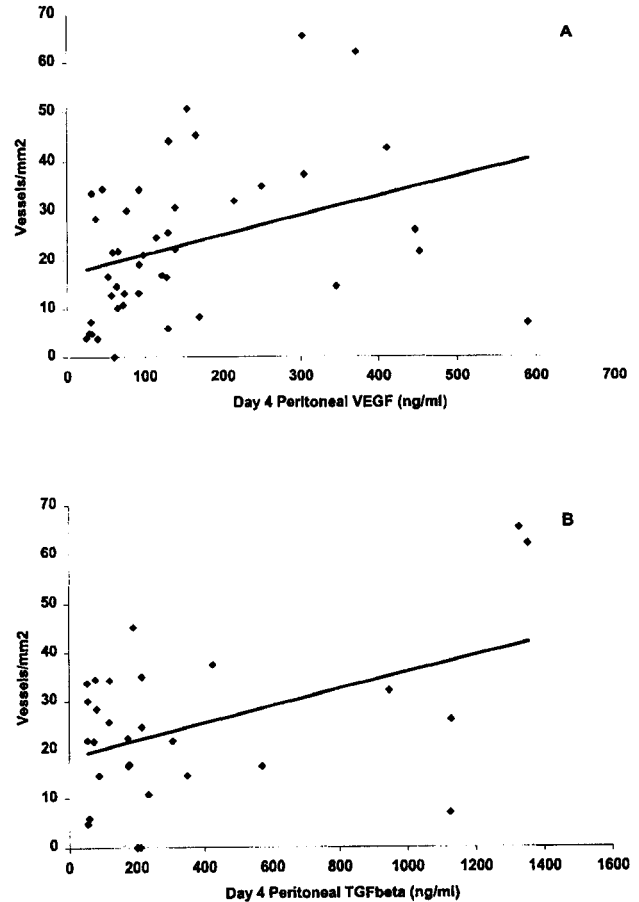


Figure 7. Correlation between peritoneal cytokine concentrations in all three treatment groups and vascularization measured by vessels/mm². **A:** Day 4 peritoneal VEGF concentration correlates weakly with vascularization ($r = 0.35$, $P = 0.02$). **B:** A stronger correlation exists between peritoneal TGF- β 1 concentration and vascularization ($r = 0.45$, $P = 0.01$).

virus-treated animals. This response consisted of increased peritoneal inflammatory cell infiltration, notably by neutrophils. There was a secondary up-regulation of IL-6 and TNFsrII, as has been seen in patients with peritonitis.⁵

IL-1 β and TNF- α also induced functional changes similar to those seen in patients who develop peritonitis. Specifically, we saw an early ultrafiltration failure with increased transport of glucose and loss of albumin. These functional changes corresponded to increased expression of VEGF and TGF- β 1 and increased peritoneal angiogenesis. We, and others, have demonstrated that angiogenesis is a key component in increased glucose transport and ultrafiltration dysfunction^{16,30} and we suggest that, in this model, angiogenesis is an integral factor causing the acute peritoneal dysfunction. Others have observed a role for nitric oxide⁹ or prostaglandins¹⁰ in acute peritoneal dysfunction associated with peritonitis. We did not assay for these compounds in our experiments and they may play a role, but they are likely downstream components and the inflammatory cytokines can alone initiate peritoneal angiogenesis and acute ultrafiltration failure.

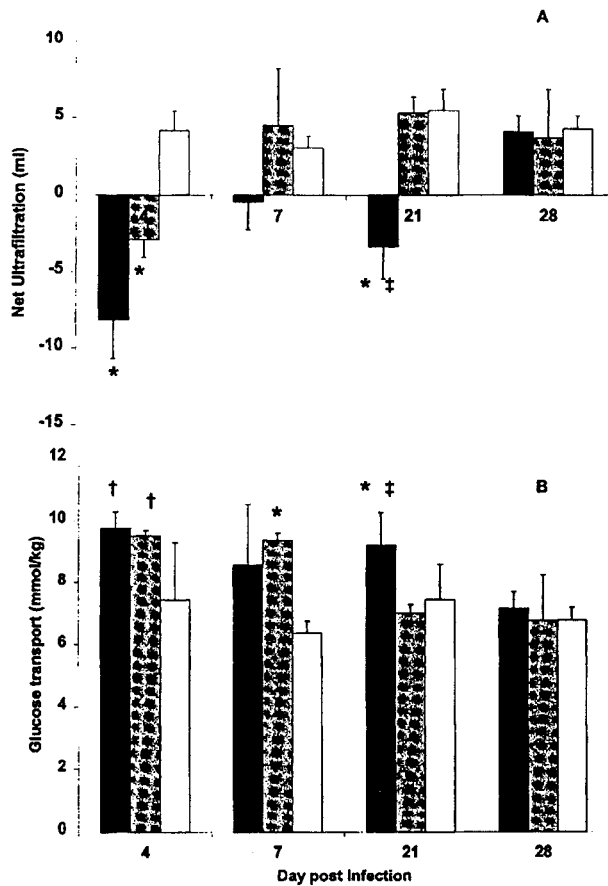


Figure 8. Peritoneal membrane function after adenovirus infection with AdIL-1 β (black bars), AdTNF- α (gray bars), or control adenovirus (AdDL70, open bars) represented by net ultrafiltration (A) and mass transport of glucose transport across the peritoneal membrane (B). AdIL-1 β -treated animals displayed impaired peritoneal function with decreased ultrafiltration and increased solute transport until day 28. AdTNF- α -treated animals demonstrated a more transient membrane dysfunction with resolution of ultrafiltration failure at day 7 and normalization of solute transport by day 21 after infection. *, $P < 0.01$ compared with AdDL70; †, $P < 0.05$ compared with AdDL70; ‡, $P < 0.05$ AdIL-1 β compared with AdTNF- α . Data represents three to six animals for each data point.

The role of inflammatory cytokines in angiogenesis has been studied previously and the results are in agreement with our findings. Specifically, in a series of studies using the rat mesenteric window model, Norrby^{20,21} has shown that both inflammatory cytokines induce angiogenesis, but the angiogenesis induced by IL-1 β is more prolonged.

The most interesting findings from our experiments derive from the differences we observed between IL-1 β and TNF- α . Specifically, TNF- α induced a transient, dramatic angiogenesis with collagen deposition and increased macromolecular permeability. These changes resolved completely by 21 days after adenovirus-mediated gene transfer. In contrast, IL-1 β exposure induced a persisting vasculogenesis with associated up-regulation of profibrotic cytokines and induction of submesothelial fibrosis.

There was a quantitative difference in inflammatory response to IL-1 β and TNF- α with the former leading to a more severe and prolonged inflammation measured by persisting neutrophilia and up-regulation of TNFsrl. We

do not think this quantitative increased inflammatory response alone explains the qualitative fibrotic and angiogenic response seen by days 21 and 28. We saw a similar angiogenic and fibrogenic response between TNF- α and IL-1 β at early time points. This was accompanied by an equal up-regulation of TGF- β and fibronectin mRNA and tissue VEGF. We hypothesize that the different fibrogenic response at later time points is related to the significant up-regulation of TIMP-1 as demonstrated by RNase protection assay (Figure 1). Increasing evidence would suggest that persisting fibrosis is related to imbalance between collagenase proteins and their inhibitors.³¹ The profibrotic nature of IL-1 β has been demonstrated previously in which overexpression in the lung led to progressive fibrosis,¹⁸ however overexpression of TNF- α in the lung led to a patchy, mild fibrosis.³²

One possible explanation for the differential angiogenic response to IL-1 β and TNF- α is the recognized property of TNF- α to induce apoptosis through binding to its receptor p55.³³ IL-1 β is not known to directly induce apoptosis. It is possible that the high peritoneal concentration of TNF- α seen during the first 7 days induces inflammation and angiogenesis and, as the concentration falls, there is a secondary induction of apoptosis and reversion of the membrane to normal histology by day 21.

Another possibility is that some element, or elements, of the profibrotic environment initiated by IL-1 β is responsible for protecting the viability and stability of endothelial cells. Recent experiments have revealed that angiogenesis involves the growth of new immature blood vessels following a stabilization phase.³⁴ This stabilization involves different stimuli such as angiopoietin-1³⁵ along with the recruitment of perivascular cells and these factors may be enhanced in the fibrotic environment.

We cannot rule out a differential response to human IL-1 β , the gene insert in the adenovirus used in these experiments. However, IL-1 β has significant homology across mammalian species³⁶ and the receptor apparatus is known to be remarkably well preserved.³⁷ Only very minor differences in response to different species IL-1 β has been documented previously.³⁸ Therefore, we do not feel that the differential effects between IL-1 β and TNF- α seen in these experiments are because of the presence of human IL-1 β .

We have not clearly identified the major angiogenic cytokine in this model. VEGF concentration was up-regulated in the peritoneal dialysate but this is not an accurate marker of local production.³⁹ Peritoneal cytokine concentration is effected by shifts of water into or out of the peritoneal cavity as well as transport of the cytokine into the peritoneum across permeable blood vessels. The tissue concentration of VEGF was elevated in animals exposed to TNF- α and IL-1 β but the changes from AdDL70-treated animals were not striking (Table 2). It could be that higher tissue VEGF levels are present at earlier time points, as angiogenesis is already present by day 4. It is also possible that other fibrosis-related cytokines such as platelet-derived growth factor or basic fibroblast growth factor are important modulators of angiogenesis in this setting of acute inflammation.⁴⁰ This could explain the observation that peritoneal fluid TGF- β 1

concentration correlates more closely with vascularization than VEGF concentration (Figure 7).

Some discrepancies exist between functional and morphological findings in the peritoneum. Seven days after infection with AdTNF- α , there is persisting angiogenesis and increased glucose transport, but the ultrafiltration has reverted to normal in AdTNF-treated animals (Figure 8). One possible confounding variable is the extremely high albumin transport measured in TNF- α -exposed animals at this time point. The increased osmotic component of the peritoneal protein may counteract the rapid loss of glucose gradient and preservation of ultrafiltration. Functional properties of IL-1 β -treated animals improved at day 28 despite ongoing evidence of angiogenesis and fibrosis. We saw a similar pattern after adenoviral-mediated gene transfer of TGF- β 1.²³ It is possible that there are progressive alterations in the interstitium of the peritoneum in fibrosis, such as increased deposition of hyaluronan, that may improve the peritoneal membrane function despite ongoing angiogenesis.⁴¹ Alternatively, the peritoneum may regulate the blood supply to the peritoneum throughout time so that, despite the histological persistence of blood vessels, the overall volume of blood supply slowly returns to a basal level.

Using adenoviral-mediated gene transfer of the proinflammatory cytokines IL-1 β and TNF- α , we have shown that these cytokines individually can induce peritoneal angiogenesis and functional changes similar to those found in peritoneal dialysis patients who suffer acute peritonitis. More importantly, IL-1 β and TNF- α induce differing fibrogenic and angiogenic responses. These differential responses suggest that IL-1 β may have a greater and more prolonged impact on the peritoneum through induction of sustained vasculogenesis and fibrosis. IL-1 β may be the link between recurrent peritonitis and longer-term changes in the function of the peritoneum. This may explain why some patients have a poor outcome after one or several episodes of peritonitis. Targeted inhibition of IL-1 β , perhaps through its receptor antagonist, may preserve the peritoneum after acute peritonitis. Finally, the further study of the angiogenic response in this inflammatory model may yield insights into the supporting role of fibrosis in angiogenesis and this may have implications beyond maintenance of the peritoneum as a dialysis membrane.

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CHAPTER 4: A CHRONIC INFLAMMATORY INFUSION MODEL OF PERITONEAL DIALYSIS IN RATS

This short paper appeared in *Peritoneal Dialysis International* 2001; 21 (Suppl 3): S368 – S372. In this paper, we describe our initial experience with a chronic daily infusion model of peritoneal dialysis. In this model, we surgically placed an indwelling catheter attached to a subcutaneous port and daily injected the rats with various solutions. We used LPS in this model to induce an acute inflammatory reaction, but subsequent experience demonstrated that the LPS was not necessary.

We saw a small effect on peritoneal morphology of the catheter placement alone with no fluid infusion. Ringer's Lactate or buffered saline solution had a definite effect on peritoneal morphology. We saw an impact on UF, thickness of the submesothelium, and angiogenesis, with this 'physiological saline' solution. This was contrary to what others have observed in longer models¹⁸⁰. I believe the catheter alone induces an inflammatory reaction, and the fluid instillation allows the catheter to move around the peritoneum and extend the affected area.

Dialysis solutions had a more dramatic effect on structure and function of the peritoneum compared with physiological saline infusion. This effect was likely due to the additional injurious effect of high glucose and GDPs. We did see an inverse association between peritoneal vascularization and UF. We also demonstrated the ability to express AdLacZ in the peritoneal tissues in this daily infusion model.

I developed the concept and designed these experiments. The work was carried out by myself, Dr. Kolb, and Lisa Yu. I wrote the paper which was reviewed by Drs Gauldie and Hoff.

A CHRONIC INFLAMMATORY INFUSION MODEL OF PERITONEAL DIALYSIS IN RATS

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◆ **Objectives:** Peritoneal membrane changes are related to daily exposure to non physiologic dialysate and recurrent acute inflammation. We modified a daily infusion and inflammation model and evaluated it for fibrotic and angiogenic features. The feasibility of adenovirus-mediated gene transfer in the model was also assessed.

◆ **Methods:** Peritoneal catheters were implanted in rats. Over a period of 4 weeks, the animals received a daily infusion of Dianeal 4.25% (Baxter Healthcare Corporation, Deerfield, IL, U.S.A.) with an initial three doses of lipopolysaccharide (LPS) or physiologic saline. Peritoneal fluid was assayed for transforming growth factor beta (TGF β) and vascular endothelial growth factor (VEGF). Animals were humanely killed at week 5. Net ultrafiltration was then measured, and tissue samples were immunostained for factor VIII. Mesenteric tissue was assayed for hydroxyproline content. Adenovirus-mediated gene transfer of β -galactosidase was assayed by intraperitoneal administration of the virus, 4 days before the end of the experiment.

◆ **Results:** Animals treated with either Dianeal or physiologic saline showed peritoneal membrane thickening and increased vascularity. Fibrosis was demonstrated by increased hydroxyproline concentration. Ultrafiltration was impaired. We found increased concentrations of VEGF and TGF β in the peritoneal fluid of animals treated with LPS and daily infusion. Adenovirus-mediated gene transfer to the peritoneal membrane was demonstrated in the model.

◆ **Conclusions:** Exposure to LPS and daily Dianeal or physiologic saline leads to peritoneal fibrosis and neo-angiogenesis. Vascularization and glucose transport correlate with ultrafiltration failure. The present animal model mimics changes seen in humans on peritoneal dialysis and may be valuable for evaluating short-term interventions to prevent membrane damage.

KEY WORDS: Fibrosis; angiogenesis; vascular endothelial growth factor; transforming growth factor; ultrafiltration; animal model.

Long-term peritoneal dialysis is characterized by alterations in the peritoneum leading to increased solute transport and decreased ultrafiltration (1). Those effects are likely due to exposure to non physiologic dialysate and recurrent inflammatory events. After long-term dialysis, the peritoneum shows signs of fibrosis with collagen deposition, thickening, and perivascular sclerosis (2). Several biopsy studies have identified increased vascularization in the peritoneum and correlated that finding with membrane dysfunction (3). Profibrotic cytokines such as transforming growth factor beta 1 (TGF β 1) (4) and angiogenic factors such as vascular endothelial growth factor (VEGF) (5) have been identified in peritoneal effluent and correlated with membrane function.

We previously developed an acute model that uses adenovirus-mediated gene transfer to study the effects on the peritoneum of transient exposure to a high concentration of a cytokine (TGF β 1) (6). In later experiments, we adopted a shorter-term chronic model (7) in which we could intervene with adenovirus to alter the environment of the peritoneum and attempt to modify the damaging effects of daily infusion. The original model, described by Kim and colleagues (7) used a combination of twice-daily exposure to dialysate with an initial administration of lipopolysaccharide (LPS). That model induced significant alterations in the peritoneum after 4 weeks. That model therefore seemed appropriate for the introduction of adenovirus-mediated gene transfer to the peritoneum, which has demonstrated transgene expression for a period of 10 – 14 days.

Here we describe our initial experience with the model. We show that significant histologic and functional changes occur in the parietal peritoneum after 4 weeks of daily infusion. The effects include increased

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thickness of the peritoneum, neovascularization, increased glucose transport, and significantly impaired ultrafiltration. Those effects were seen both with physiologic saline solution and with high-concentration glucose dialysate. We also demonstrate that, in the model, adenovirus is effectively taken up and expressed by mesothelial cells.

METHODS

All animal studies were carried out according to the guidelines of the Canadian Council on Animal Care. Sprague-Dawley rats, 200 – 250 g (Harlan, Indianapolis, IN, U.S.A.) were anesthetized with ketamine. A 7-French silicone catheter (Access Systems, Newark, NJ, U.S.A.) was inserted into the peritoneum through a small incision in the abdominal wall. The catheter was tunneled subcutaneously to an implanted port (Access Systems). The animals were allowed to recover for 1 week before daily fluid infusion was initiated. The animals were treated with antibiotics.

Four groups of animals were studied, and all were humanely killed after 5 weeks. The first group ($n = 5$) had no catheter inserted. A second group ($n = 3$) had a catheter inserted, but no fluid instilled. A third group ($n = 8$) had catheters and daily infusion of phosphate-buffered saline or Ringer's lactate from week 1 to week 5. The last group ($n = 11$) had daily instillation of dialysate (Dianeal 4.25%: Baxter Healthcare Corporation, Deerfield, IL, U.S.A.) along with 75 μ g LPS (Sigma-Genosys, Oakville, Canada) on days 8, 10, and 12. Animals having fluid infused received 15 mL for the first 3 days, then 20 mL for 3 days, and then 25 mL for the remainder of the duration of the experiment. At week 3, a peritoneal fluid sample was taken from all animals 4 hours after infusion of 20 mL Dianeal 2.5%. At the end of the experiment, 20 mL of Dianeal 2.5% was infused, and 4 hours later the animals were killed, an accurate ultrafiltration volume was measured, and blood and tissue samples were taken.

Peritoneal fluid samples were centrifuged at 1500 rpm for 5 minutes. Glucose was measured by a standard enzymatic test on a Hitachi 917 automated chemistry analyzer (Roche Diagnostics, Laval, Canada). Mass transfer of glucose out of the peritoneum was calculated using the formula:

$$\begin{aligned} & (\text{initial dialysate glucose} \times \text{initial volume}) \\ & - (\text{final dialysate glucose} \times \text{final volume}) \end{aligned}$$

Values were corrected for animal weight at the time of death.

To demonstrate effective adenoviral gene transfer, a replication-deficient recombinant adenovirus expressing β -galactosidase (AdLacZ) was administered

through the indwelling catheter in 2 animals. Adenovirus at a dose of 2×10^9 plaque-forming units diluted to 200 μ L with PBS was administered 7 days after initiation of the daily dialysate infusion. The animals were killed 4 days later. Cytochemical staining for β -galactosidase was carried out on samples from animals treated with adenovirus. Fresh tissue samples were fixed in 2% formaldehyde, 0.2% glutaraldehyde for 3 hours. Fixation was followed by staining for 6 hours in a potassium ferrous cyanide, potassium ferric cyanide, magnesium chloride solution (Sigma), Triton X100, and X-gal (5-bromo-4-chloro-3-indolyl- β -D-galactopyranoside: Boehringer Mannheim, Indianapolis, IL, U.S.A.). Sections were counterstained with nuclear fast red.

Tissue samples were taken from the lower anterior abdominal wall distant from the site of catheter placement. Sections from both sides of the midline were taken and fixed in a sufficient amount of 4% phosphate-buffered formaldehyde for 24 hours. The tissue samples were then paraffin processed and embedded, and 3 μ m sections were cut. Cut sections were then stained for Masson trichrome and immunohistochemistry was carried out with antibodies to von Willebrand factor-factor VIII-related antigen (Dako Corporation, Carpinteria, CA, U.S.A.).

Sections of the anterior abdominal wall, immunostained for factor VIII, were studied in a blinded fashion using a standardized microscope grid to count peritoneum-associated blood vessels and to measure the thickness of the peritoneum. Slides contained at least 4 separate tissue pieces, and each slide was examined in 10 – 12 random high-power views.

Peritoneal fluid samples were analyzed using a human TGF β 1 ELISA kit or mouse VEGF ELISA kit (R&D Systems, Minneapolis, MN, U.S.A.). Samples were treated with HCl, pH-adjusted with NaOH, and then assayed for total (active and latent) TGF β 1.

A portion of mesentery was taken and frozen for a hydroxyproline assay. Tissues were weighed and homogenized in water. Solid material was precipitated with trichloroacetic acid. Samples were hydrolyzed overnight in 6N HCl at 110°C. Hydroxyproline content was quantified by Erlich reagent (Sigma) and assayed by measuring the optical density at 557 nm.

Data are presented as mean \pm standard deviation unless otherwise noted. Comparisons between groups were made by t-test. Linear regression was used to compare cytokine levels, quantitative immunohistochemistry, solute transport, and net ultrafiltration.

RESULTS

In general, we saw significant changes in the histologic and functional characteristics of the peritoneum in animals exposed to daily physiologic saline

MARGETTS *et al.*

or 4.25% Dianeal as compared to animals in the catheter-only or no-catheter control groups (Table 1). Net ultrafiltration after a 4-hour dwell with Dianeal 2.5% was significantly impaired in animals exposed to daily fluid, and the mass transfer of glucose was significantly increased. We also demonstrated a significant increase in the number of peritoneal vessels and in the thickness of the submesothelial tissue (Table 1, Figure 1). Animals exposed to daily fluid, both isotonic saline and Dianeal 4.25%, showed increased collagen deposition as measured by mesenteric hydroxyproline content. The thickness of the peritoneum in rats in the catheter-only group was increased, with a nonsignificant increase in vascularization. Those changes did not seem to affect functional characteristics such as glucose transport and net ultrafiltration.

The peritoneal effluent taken at week 3 showed a significant elevation in VEGF concentration in animals exposed to daily Dianeal 4.25% (135 pg/mL vs 70 pg/mL in no-catheter control animals, $p = 0.03$).

In the isotonic saline group, we treated 5 animals with Ringer's lactate and 3 with phosphate-buffered saline. We saw no differences between the treatments, and those animals were grouped for analysis. Overall, a nonsignificant trend toward more severe changes in ultrafiltration, in glucose transport, in submesothelial thickness, and in number of vessels was seen in animals treated with daily Dianeal 4.25% as compared with animals treated with isotonic saline (Table 1). Glucose-treated animals had a higher total TGF β dialysate concentration as measured by ELISA (411 pg/mL vs 231 pg/mL, $p = 0.03$).

In animals treated with saline or glucose, the mass transport of glucose was inversely correlated with net

CHRONIC INFUSION MODEL OF PERITONEAL DIALYSIS

ultrafiltration ($r = -0.712$, $p < 0.001$). The number of peritoneum-associated vessels per millimeter of peritoneum was also inversely correlated with net ultrafiltration ($r = -0.524$, $p = 0.03$). Finally, the total TGF β 1 concentration at week 3 was correlated with the thickness of the peritoneum as measured at week 5 ($r = 0.5$, $p = 0.03$).

We were able to identify uptake and expression of adenovirus by staining for β -galactosidase 4 days after transfection with AdLacZ (Figure 1).

DISCUSSION

We further explored the chronic inflammatory infusion model first described by Kim and colleagues (7). We noted impressive fibrosis and neovascularogenesis with associated ultrafiltration failure in animals treated with daily fluid infusion. The concentrations of profibrotic and vasculogenic cytokines in the dialysate were increased. The strong inverse correlation between glucose transport and net ultrafiltration, and between new peritoneal blood vessels and net ultrafiltration suggest that increased vascular surface area and the subsequently increased glucose transport leads to significantly decreased ultrafiltration. The dialysate concentration of TGF β 1 correlated with thickening of the peritoneum, but not with vascularity or transport properties, suggesting an important role for vasculogenesis, as compared with fibrogenesis, in the functional changes noted.

We saw fairly dramatic changes with infusion of isotonic saline. The reason for that finding is not clear, but infection is unlikely, given our extensive use of antibiotics. A foreign body reaction also does not ex-

TABLE 1
Functional and Histologic Changes in the Peritoneum—Daily Infusion Model

	No catheter (<i>n</i> =5)	Catheter only (<i>n</i> =3)	Isotonic saline (<i>n</i> =8)	4.25% Dianeal (<i>n</i> =11)
Net ultrafiltration (mL)	4.3±1.8	3.2±0.1	-5.9±2.4 ^a	-8.0±3.9 ^a
Mass transfer glucose (mmol/kg)	6.8±0.4	6.7±0.8	7.8±0.4 ^a	8.0±0.9 ^a
Vessels (/mm)	0.7±0.5	13±11	25±9 ^a	30±33
Thickness (μ m)	31±14	88±24 ^b	118±50 ^a	205±170
Hydroxyproline (μ g/mg)	1.41±0.4	1.88±0.3	3.37±0.9 ^a	2.83±1.2 ^a
VEGF (pg/mL)				
Week 3	70±36	182±44	135±82	135±50 ^a
Week 5	74±41	122±85	120±87	111±68
Total TGF β (pg/mL)				
Week 3	247±44	221±150	231±91 ^c	411±185 ^c
Week 5	210±21	225±41	260±104	308±184

^a $p < 0.05$ physiologic saline or Dianeal versus no catheter.

^b $p < 0.05$ catheter only versus no catheter.

^c $p < 0.05$ physiologic saline versus Dianeal.

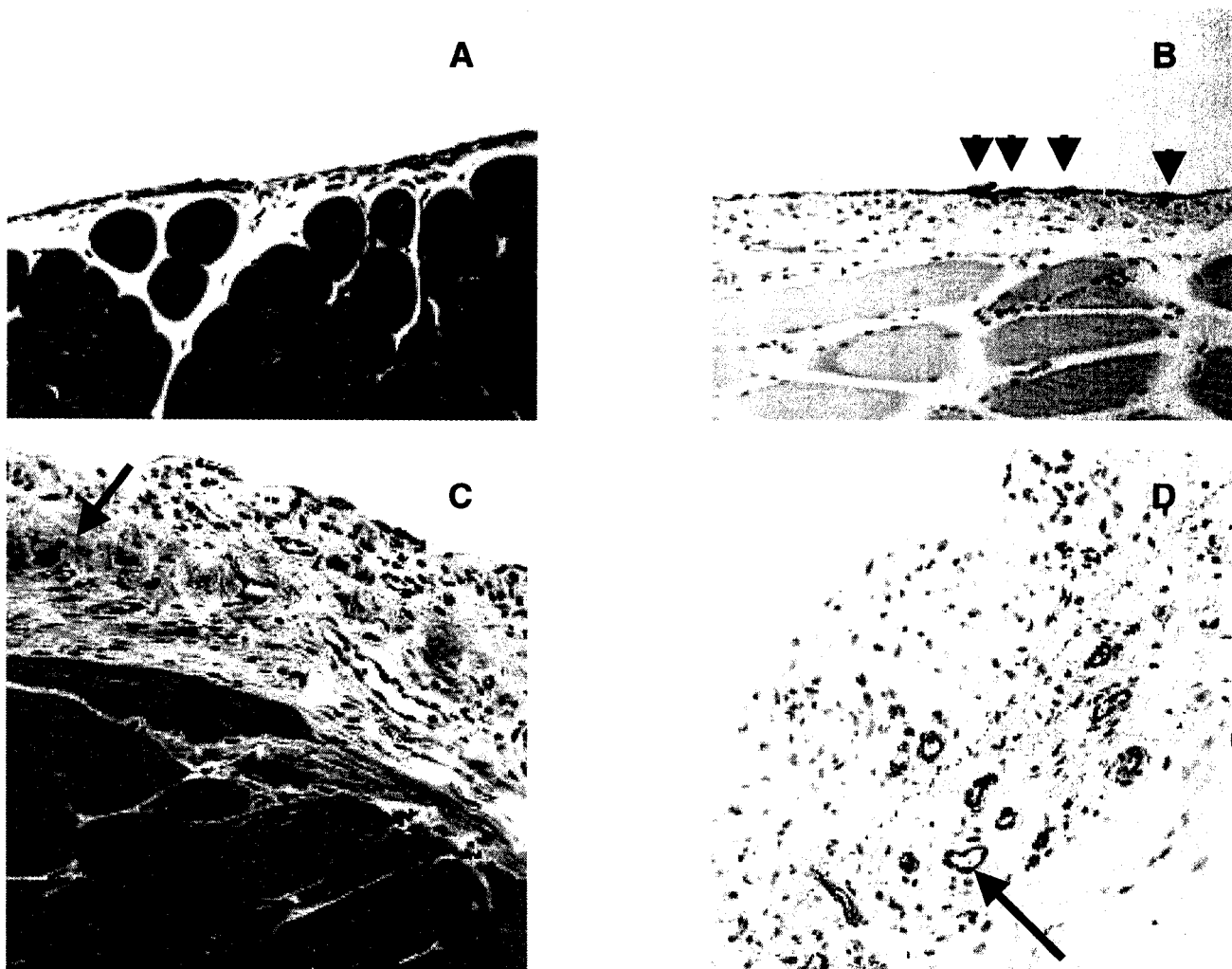


Figure 1 — (A) Normal anterior abdominal wall with peritoneal tissue from “no catheter” control animal. (B) Anterior abdominal wall from animal treated with adenovirus-mediated gene transfer of β -galactosidase and stained with X-gal (5-bromo-4-chloro-3-indolyl- β -D-galactopyranoside). Arrowheads indicate mesothelial cells infected with adenovirus and expressing transgene product. (C) Peritoneum after 4 weeks of daily infusion with 4.25% Dianeal. Tissue is thickened with increased collagen deposition (arrow) and neovascularization. (D) Factor VIII–von Willebrand factor stain of same section as in (C) demonstrates neovascularization (arrow). All sections 125 \times magnification. (A) and (C) use Masson trichrome; (B) is counterstained with nuclear fast red.

plain the entire effect, because changes in the catheter-only animals were minimal.

We hypothesize that the trauma of catheter insertion initiates a fibroproliferative response that is rapidly contained in an otherwise untouched peritoneum. In the presence of daily fluid infusion, the ability of the peritoneum to respond to the insult is delayed, and the overall fibroproliferative effect is prolonged. A trend toward more pronounced effects was seen with Dianeal 4.25%. That finding suggests that glucose or other solution components may add to the already profibrotic state of the peritoneum after catheter insertion. However, the roles of LPS and glucose cannot be separated in the present experiments.

By 5 weeks, the VEGF and TGF β 1 dialysate concentrations decline (Table 1), which suggests that the acute fibroproliferative response would subside if the model were prolonged. Less dramatic, but clearly evident changes have been seen in 16-week to 20-week chronic infusion models (8), leading us to believe that an initial acute fibroproliferative phase occurs, subsides, and is then replaced by a more slowly developing chronic fibrosis and vascular injury.

Finally, we demonstrated that adenovirus-mediated gene transfer is possible in the present model. Adenovirus vectors have been engineered to transfer genes coding anti-inflammatory, anti-angiogenic (9), and anti-fibrotic (10) cytokines. The effect of those

MARGETTS *et al.*

agents can be studied in the present chronic inflammatory infusion model.

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CHAPTER 5: ANTIANGIOGENIC AND ANTIFIBROTIC GENE THERAPY IN A CHRONIC INFUSION MODEL OF PERITONEAL DIALYSIS IN RATS

This final paper was published in the Journal of the American Society of Nephrology 2002; 13: 721-728. In this paper, we used our established daily dialysis model in rats and treated the rats with either antiangiogenic (AdAngiostatin) or antifibrotic (AdDecorin) gene therapy. This paper identified 2 main points. First, AdAngiostatin therapy significantly reduced the peritoneal vasculature. In places, we could see a dramatic apoptosis of endothelial cells. With this decrease in vasculature was an improvement (but not normalization) in UF function. Decorin, on the other hand, significantly reduced fibrosis measured by hydroxyproline concentration, but did not effect vascular density or UF function. As we had previously hypothesized that TGF β 1 is involved in angiogenesis in the peritoneum, this finding was somewhat contradictory. One possible explanation is that fibrogenic and angiogenic functions of TGF β 1 are separated by the type of activin-like kinase (ALK) receptor¹⁷². Decorin may block TGF β 1 signaling through ALK-5 and thus prevent fibrosis, but may allow signaling through ALK-1 and thus may not abrogate the angiogenic effect. This hypothesis requires further work to confirm.

The angiostatin adenovirus was created by Dr. Steve Gyorffy. The decorin adenovirus was created by Dr. Patricia Sime. I developed the concept and designed these experiments. The work was carried out by Lisa Yu with some assistance from Dr. Martin Kolb and myself. The paper was written by myself and was reviewed by Drs Gauldie, Hoff and Clifford Holmes.

Antiangiogenic and Antifibrotic Gene Therapy in a Chronic Infusion Model of Peritoneal Dialysis in Rats

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Abstract. To identify the relative importance of peritoneal fibrosis and angiogenesis in peritoneal membrane dysfunction, adenoviral mediated gene transfer of angiostatin, a recognized angiogenesis inhibitor, and decorin, a transforming growth factor- β -inhibiting proteoglycan, were used in a daily infusion model of peritoneal dialysis. A peritoneal catheter and subcutaneous port were inserted in rats. Five and fourteen d after insertion, adenovirus-expressing angiostatin, decorin, or AdDL70, a null control virus, were administered. Daily infusion of 4.25% Baxter Dianeal was initiated 7 d after catheter insertion and continued until day 35. Three initial doses of lipopolysaccharide were administered on days 8, 10, and 12 to promote an inflammatory response. Net ultrafiltration was used as a measure of membrane function, and peritoneum-associated vasculature and mesenteric collagen content was quantified. Ultrafiltration dysfunction, angiogenesis, and fibrosis

were observed in daily infusion control animals. Animals treated with AdAngiostatin demonstrated an improvement in net ultrafiltration (-3.1 versus -7.8 ml for control animals; $P = 0.0004$) with a significant reduction in vessel density. AdDecorin-treated animals showed a reduction in mesenteric collagen content (1.8 versus 2.9 $\mu\text{g}/\text{mg}$; $P = 0.04$); however, AdDecorin treatment had no effect on net ultrafiltration. In a rodent model of peritoneal membrane failure, net ultrafiltration was significantly improved and peritoneal-associated blood vessels were significantly reduced by using adenovirus-mediated gene transfer of angiostatin. Decorin, a transforming growth factor- β -inhibiting proteoglycan, reduced collagen content but did not affect net ultrafiltration. Improvement in the function of the peritoneum as a dialysis membrane after treatment with angiostatin has implications for treatment of peritoneal membrane dysfunction seen in patients on long-term dialysis.

During long-term peritoneal dialysis, the peritoneum undergoes histologic changes that include increased submesothelial collagen deposition, increased vascularization with vasculopathy, and loss of mesothelial cells (1,2). These changes suggest that both fibrogenic and angiogenic processes are active. Peritoneal concentration of vascular endothelial growth factor (VEGF) (3) and transforming growth factor- β (TGF- β) (4), key angiogenic and fibrogenic growth factors, have also been shown to correlate with peritoneal membrane function. These features strongly suggest that fibrogenic and angiogenic processes in the peritoneum are responsible for the alteration in peritoneal membrane function that is seen in patients on long-term peritoneal dialysis.

In these experiments, we used a short-term daily infusion model of peritoneal dialysis in rats. This model is associated with increased expression of VEGF and TGF- β , thickening of the submesothelial zone of the parietal peritoneum, increased

angiogenesis, ultrafiltration failure, and increased transport of solutes (5). Similar models have previously been used to demonstrate the effect of peritoneal membrane rest (6) and systemic angiotensin-converting enzyme inhibition (7) on peritoneal membrane function.

Adenovirus-mediated gene transfer has been shown to be an effective tool for studying changes induced by transient overexpression of cytokines or growth factors on the peritoneal membrane. We have demonstrated that the adenovirus vector, when administered to the peritoneum, is highly infective for mesothelial cells. The transferred gene product is produced in high quantities and is expressed from approximately day 4 to day 14 after infection. We have previously demonstrated that adenovirus-mediated gene transfer of the active form of TGF- β_1 leads to fibrosis of the peritoneum with increased VEGF expression and vasculogenesis (8). This led to an associated change in function, including increased transport of glucose and decreased net ultrafiltration.

The antiangiogenic and antifibrotic approaches we used in the following experiments have been previously described. Angiostatin is a potent inhibitor of endothelial cell proliferation that was first described for its ability to inhibit cancer growth by inhibiting tumor neovascularization (9). The adenovirus carrying the gene product for angiostatin (AdAngiostatin) has been effective in several studies, including a direct tumor injection model (10).

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Decorin is a proteoglycan that binds and inactivates TGF- β . Decorin has been shown to be effective in direct application to prevent lung fibrosis (11) and as systemic gene therapy to modify renal fibrosis (12). We have shown adenovirus-mediated gene transfer of decorin (AdDecorin) to be an effective treatment in preventing pulmonary fibrosis in a murine bleomycin model (13).

In these experiments, we introduced AdAngiostatin, AdDecorin, or a control adenovirus (AdDL70) into a 4-wk, daily dialysate exposure model in rats. We demonstrate that animals treated with AdAngiostatin showed reduction in the parietal peritoneal vasculature, decreased glucose transport, and improved ultrafiltration. Animals treated with AdDecorin did not demonstrate alteration in peritoneal vasculature or ultrafiltration, but they did show reduced peritoneal fibrosis.

Materials and Methods

Animals

All animal studies were carried out according to the Canadian Council on Animal Care Guidelines. Sprague-Dawley rats, 200 to 250 mg (Harlan, Indianapolis, IN) were anesthetized with ketamine/xylazine. A 7 French silicone catheter (Access Systems, Newark, NJ) was inserted into the peritoneum through a small incision in the abdominal wall. The catheter was tunneled subcutaneously to an implanted port (Access Systems) on the back.

Animals were administered adenovirus (2×10^9 plaque-forming units [pfu] per animal) 5 d after catheter insertion with a repeat administration on day 14. Both administrations were given intraperitoneally via the dialysis catheter. Daily infusion of dialysate (4.25% Dianeal; Baxter Healthcare, McGaw Park, IL) was initiated 7 d after catheter insertion, beginning at 15 ml then quickly increasing to 25 ml daily. Animals also received intraperitoneal lipopolysaccharide (LPS-75 μ g; Sigma, Oakville, ON) on days 8, 10, and 12. Animals were treated with rotating antibiotics consisting of intraperitoneal cefazolin, intraperitoneal ciprofloxacin, and oral sulfamethoxazole-trimethoprim.

Four groups of animals were studied, and all were killed after 5 wk. The first was a positive control group ($n = 11$) that received control virus (AdDL70). The second group ($n = 10$) received AdAngiostatin, and the third group ($n = 7$) received AdDecorin. For negative control, we used a group of animals ($n = 5$) with no catheter inserted and administered AdDL70 at initiation. At week 3 in all animals, a peritoneal fluid sample was taken 4 h after infusion of 20 ml of 2.5% Dianeal. At week 5, 20 ml of 2.5% Dianeal was infused, and 4 h later, the animal was killed, an accurate ultrafiltration volume was measured, and blood and tissue samples taken. In a subset of AdAngiostatin-treated animals, we took a peritoneal fluid sample 4 d after first administration of adenovirus to demonstrate effective gene transfer.

Whole blood was centrifuged at 5000 rpm for 10 min and the serum removed. Peritoneal fluid samples were centrifuged at 1500 rpm for 5 min. Samples were analyzed on a Hitachi 917 automated chemistry analyzer (Roche Diagnostics, Laval, Canada) for creatinine, glucose, and albumin. Mass transfer of glucose out of the peritoneum was calculated as (initial dialysate glucose \times initial volume) – (final dialysate glucose \times final volume). Creatinine and albumin clearances were calculated as mass transfer divided by the serum solute concentration. All values were corrected for animal weight at death.

To demonstrate the safety of adenovirus-mediated gene transfer of angiostatin, we studied 8 rats without peritoneal catheters. These animals were treated with 1 ($n = 4$) or 2 ($n = 4$) doses of AdAn-

giostatin (2×10^9 pfu intraperitoneally). Groups of 2 animals were sacrificed 4 and 7 d after treatment, and tissue samples were taken, fixed in neutral buffered formalin, embedded in paraffin, sectioned, and stained for hematoxylin/eosin and Masson trichrome. These sections were examined for pathologic changes in the peritoneum, liver, spleen, kidney, bowel, and ovaries.

Adenovirus

The cDNA for murine angiostatin was constructed from the four kringle regions of plasminogen ligated to the endogenous signal sequence as described previously (14). The construction of the adenovirus for human decorin (15) and control virus (AdDL70) (16) have been previously described. Adenovirus preparations were purified by CsCl gradient centrifugation and PD-10 Sephadex chromatography (Amersham Pharmacia, Baie d'Urfe, Quebec), and plaque was titered on 293 cells as described previously (17).

Histology

In daily dialysis-treated animals, tissue samples were taken at sacrifice from the lower anterior abdominal wall distant from the site of catheter placement. Sections from both sides of the midline were taken and fixed in a sufficient amount of 4% phosphate-buffered formaldehyde for 24 h. The tissue samples were then paraffin processed, embedded and 5 μ m sections cut. Cut sections were then stained for Masson trichrome, and immunohistochemistry was carried out with antibodies to von Willebrand factor-factor VIII-related antigen (Dako Corporation, Carpinteria, CA). Negative control sections were run in parallel. Sections were deparaffinized in xylene followed by 100% ethanol and then placed in a methanol H₂O₂ solution for 30 min to block endogenous peroxidase activity. After hydration to water with graded alcohols, the sections were placed in 0.05 M Tris-buffered saline (TBS), pH 7.6, digested with 0.05% Pronase (Sigma) in TBS with calcium chloride for 17 min at room temperature then blocked in 5% normal goat serum (NGS) followed by a 1 h incubation in the 1:500 rabbit anti-human factor VIII in 1% NGS. Sections were then incubated in a prediluted kit of a biotinylated goat anti-rabbit followed by a streptavidin/peroxidase conjugate (Zymed Labs, San Francisco, CA) as per manufacturer's instructions. Incubations were carried out at room temperature, and sections were washed in between incubations 3 \times 5 min with 0.05 M TBS, pH 7.6, except before the addition of the primary antibody. All sections were rinsed in 0.05 M acetate buffer, pH 5.0, before development in an aminoethylcarbazole (AEC) chromogen substrate for 15 min. All sections were counterstained in Mayer's hematoxylin for 2 min before mounting with glycerin gelatin.

We stained for apoptosis with *in situ* labeling for free 3' hydroxy-DNA using terminal deoxynucleotidyl transferase (Apotag, Intergen, Purchase, NY) according to the manufacturer's direction. Briefly, sections were deparaffinized in xylene, treated with proteinase K (20 μ g/ml) for 15 min, quenched with 3.0% hydrogen peroxide, and then exposed for 1 h to terminal deoxynucleotidyl transferase. Sections were further labeled with antidigoxigenin peroxidase conjugate, washed, and then developed with peroxidase substrate. Sections were counterstained with methyl green.

Blood Vessel Analysis

Sections of the anterior abdominal wall, immunostained for factor VIII, were studied in blinded fashion using two methods. First, we used a standardized microscope grid to count peritoneal-associated blood vessels and to measure the thickness of the submesothelial zone. Slides contained four separate transverse tissue sections taken from

the lower anterior abdominal wall, and each slide was examined at 12 random high-power views. All measurements were carried out at the same magnification. Second, we took the same sections and digitized 12 fields of view containing peritoneal tissues using a Leica DMR microscope (Leica Microsystems, Wetzlar, Germany). We then analyzed these images using Leica Qwin Image Processing Software (Leica Imaging Systems, Cambridge, England) with subroutines we created. In both methods, we limited the analysis to the submesothelial collagenous zone. Results for both analyses were reported as number of vessels/mm² of peritoneal tissue. We then multiplied these values by the thickness of the submesothelial zone to arrive at an estimate of the total vasculature measured as vessels/mm. Finally, our image-processing subroutine was able to estimate the total vessel cross-section area in each digitized image, and we could therefore calculate an average cross-sectional area per vessel for each slide analyzed.

We compared these two methods of evaluating blood vessels in histologic sections using regression analysis. This showed a very close correlation between the two methods ($r = 0.87$; $P < 0.0001$).

Hydroxyproline Assay

A portion of mesentery was taken and frozen for a hydroxyproline assay, modified from Woessner method (18). Tissues were weighed, homogenized in water, and centrifuged at 1000 rpm for 5 min, and the superficial fatty material was removed by vacuum suction. Solid material was precipitated with TCA with centrifugation at 1500 rpm for 15 min at 4°C. Samples were hydrolyzed overnight in 6 N HCl at 110°C. Hydroxyproline content is quantified by Erlich's reagent (Sigma) and assayed by measuring the optical density at 557 nm. A hydroxyproline standard sample (Sigma) was used to create a standard curve.

Western Blot Analysis

Peritoneal dialysis fluid taken at week 3 was analyzed for angiostatin and decorin. Equal volume of fluid was separated on a 15% sodium dodecyl sulfate–polyacrylamide gel under nonreducing conditions and transferred to Immobulon-P membranes (Millipore, Mississauga, ON). For angiostatin, the membrane was probed with a 1:500 dilution chicken anti-rabbit-plasminogen IgY antibody (14) (generous gift of Dr. Mark Hatton, McMaster University, Hamilton, ON) followed by a secondary rabbit anti-chicken alkaline phosphatase–conjugated antibody (Zymed). Decorin was detected by using a 1:2000 dilution of anti-decorin rabbit antibody (generous gift from Dr. Larry Fisher, National Institutes of Health, Bethesda, MD), followed by a secondary anti-rabbit alkaline phosphatase–conjugated antibody (Sigma). The probed membranes were developed by using NDT/BCIP (Promega, Madison, WI) according to the manufacturer's instructions. Standard molecular weight markers (Life Technologies, Burlington, ON) were used.

Statistical Analyses

Data is presented \pm SD unless otherwise noted. Comparison between groups was made by *t* test. Regression analysis was used to look for correlation between blood vessel number/mm, net ultrafiltration, and solute transport.

Results

Adenoviral Expression and Safety of Repeated Dose of AdAngiostatin

We have previously demonstrated that adenovirus is efficiently taken up and expressed by mesothelial cells in the

peritoneum after intraperitoneal delivery (8). We have also previously shown that the control virus, AdDL70, had little effect on the peritoneum after a mild initial inflammation lasting approximately 48 h (19). In the experiments reported here, we investigated the adenoviral gene product expression by Western blot analysis of peritoneal dialysis fluid. We showed expression of angiostatin after both the first and second administration of adenovirus (Figure 1A). Likewise, AdDecorin-treated animals showed increased presence of decorin in the peritoneal dialysate (Figure 1B), suggesting effective infection and adenoviral transgene expression in this model. We have previously shown the *in vitro* efficacy of both AdAngiostatin (14) and AdDecorin (13).

We used 2×10^9 pfu of intraperitoneally delivered adenovirus, as we have shown high levels of transgene product at this dose with little persisting adenovirus effect (8). In this model, we used two doses of adenovirus vector delivered 9 d apart. There is an immunologic response to the adenovirus that makes second administration less effective but still demonstrable (20). Others have demonstrated effective repeated intraperitoneal administration of adenovirus in a tumor model (21). In our model, we studied animals administered one or two doses of

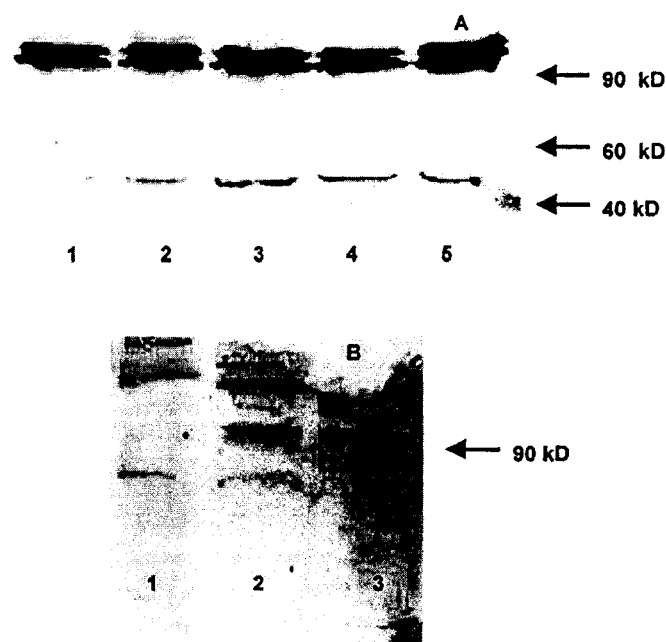


Figure 1. Western blot analysis of peritoneal dialysis fluid. (A) Angiostatin: band at 95 kD is plasminogen. Lighter band at 50 kD is angiostatin. Angiostatin is detectable in AdDL70-treated animals 7 d after 2nd virus administration (lane 1) but is greatly enhanced 4 d after 1st administration of AdAngiostatin (lanes 2 and 3) and still present 7 d after 2nd administration (lanes 4 and 5). (B) Decorin: decorin (approximately 102 kD band) was not detectable in control animals (lane 1), detectable in AdDL70, daily dialysate animals (lane 2), and enhanced after administration of AdDecorin (lane 3). Samples were separated on a 15% polyacrylamide gel, transferred to a nylon membrane, and then probed with antiplasminogen antibody (A) or anti-decorin antibody (B). Molecular markers are shown to the right of the gels.

AdAngiostatin with no daily dialysis. Histologically, we could not detect any effects of angiostatin on kidneys, spleen, liver, or ovaries. After the second administration of angiostatin, there was a mild peritoneal response with occasional eosinophils and neutrophils present in the submesothelial tissue. The mesothelial cells appeared rounded up and activated (data not shown). The effect of this peritoneal inflammatory response to adenovirus in the setting of our chronic, inflammatory model is uncertain.

Changes in Structure of the Peritoneum

We compared the histology of control animals (AdDL70, no catheter) with animals treated with 4 wk of daily dialysis, initial LPS, and either AdDL70, AdAngiostatin, or AdDecorin. Daily dialysate infusion and LPS exposure significantly altered the structure of the parietal peritoneum. As we have previously demonstrated (5), daily dialysate infusion leads to thickening of the submesothelial collagenous zone with increased total number and density of blood vessels (Figure 2; Table 1). Hydroxyproline content of mesenteric tissue is also significantly increased in animals treated with daily dialysate.

Administration of AdDecorin had significant impact on fibrogenesis in this model. There was a significant decrease in the mesenteric hydroxyproline concentration (1.8 versus 2.8 $\mu\text{g}/\text{mg}$; $P = 0.04$) (Figure 2; Table 1) compared with AdDL70, daily infusion animals. AdAngiostatin had an impact on the vascularity of the peritoneum with a significant decrease in the number ($P = 0.03$) and density ($P = 0.02$) of peritoneal vessels (Table 1). We also saw a significant decrease ($P = 0.03$) in the average cross-sectional area of blood vessels after AdAngiostatin treatment. The submesothelial thickness was not significantly reduced, but the hydroxyproline concentration of the mesentery was unchanged after AdAngiostatin treatment (Table 1).

Histologic sections of AdAngiostatin-treated animals showed areas in which blood vessels appeared to have been lost from the parietal peritoneum, and residual cells demonstrated small, condensed nuclei, suggesting apoptosis or necrosis (Figure 3). We further carried out staining for fragmented DNA using terminal deoxynucleotidyl transferase labeling and could identify an number of cells undergoing apoptosis, including endothelial cells, in sections from animals treated with AdAngiostatin. Sections from animals treated with AdDL70 did not show significant apoptosis in the submesothelial tissue (Figure 3).

Changes in Peritoneal Function

Daily dialysate also had a significant effect on the function of the peritoneum as a dialysis membrane (Table 2). Specifically, we saw a significant impairment in ultrafiltration with associated increased mass transport of glucose out of the peritoneum. There was an increase in albumin clearance along with a decrease in clearance of creatinine. When we grouped all animals studied, we identified a significant inverse correlation between the number of vessels seen in the submesothelial tissue and net ultrafiltration ($r = -0.61$; $P = 0.0002$; Figure 4). We also saw a positive correlation between the number of

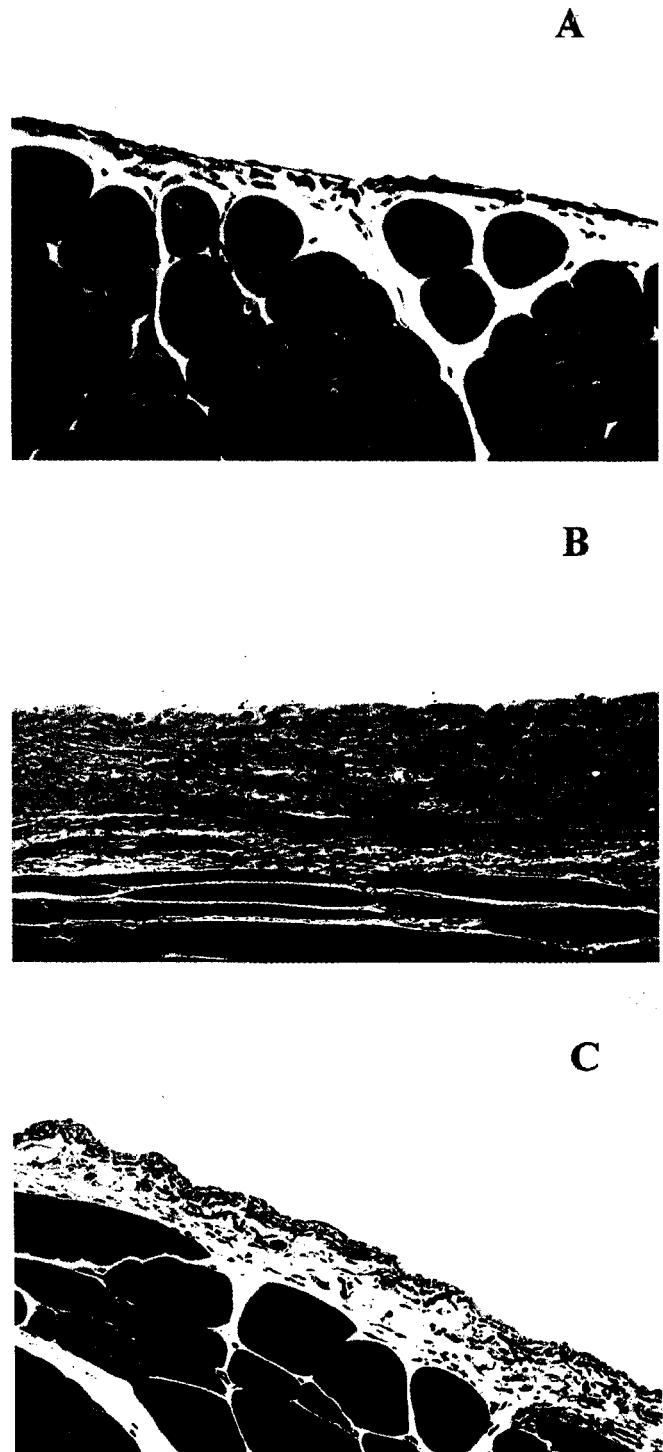


Figure 2. Histology of the anterior abdominal wall. (A) Control animal treated with AdDL70, no catheter, and no daily dialysate. Section shows normal histology with thin submesothelial zone above the abdominal wall muscles stained in red. (B) Animal treated with AdDL70, catheter, and daily dialysate. Histology is remarkably altered with thick, dense submesothelial zone and increased vascularization. (C) Animal treated with AdDecorin and daily dialysate. Section shows increased submesothelial thickening but somewhat reduced collagen deposition but similar vascularization to daily dialysate control animals. Masson trichrome stain. Magnification, $\times 50$.

Table 1. Structural changes in the peritoneum after treatment with daily dialysate and null, angiostatin, or decorin adenovirus-mediated gene transfer compared with control (no catheter) animals^a

Parameter	AdDL70, No Catheter (n = 5)	AdDL70 + Catheter (n = 11)	AdAngiostatin (n = 10)	AdDecorin (n = 7)
Thickness (μm)	30.8 \pm 13.5	193 \pm 134 ^b	122 \pm 38 ^b	200 \pm 88 ^b
Vessels/mm	0.4 \pm 0.3	12 \pm 9 ^b	5 \pm 4 ^{b,c}	11 \pm 7 ^b
Vessels/mm ²	12 \pm 10	58 \pm 21 ^b	36 \pm 19 ^c	43 \pm 16
Area/vessel (μm^2)	ND	277 \pm 96	180 \pm 86 ^c	285 \pm 128
Hydroxyproline ($\mu\text{g}/\text{mg}$)	1.4 \pm 0.3	2.8 \pm 1.0 ^b	3.2 \pm 1.9	1.8 \pm 0.9 ^c

^a ND, no data.^b $P < 0.05$ compared with AdDL70, no catheter control.^c $P < 0.05$ compared with AdDL70 + catheter control.

submesothelial vessels and the glucose transport ($r = 0.55$; $P = 0.0023$) but no association with creatinine clearance ($P = 0.3$) or albumin clearance ($P = 0.7$).

AdAngiostatin treatment appeared to have a significant positive impact on peritoneal function. There was an improvement in the ultrafiltration dysfunction ($P = 0.0004$) with a decrease in glucose transport ($P = 0.003$). The creatinine and albumin clearances were unchanged compared with the daily infusion control animals (Table 2). AdAngiostatin treatment did not completely reverse the ultrafiltration dysfunction, and the ultrafiltration in these animals was still significantly reduced compared with no catheter control animals. AdDecorin treatment had little impact on peritoneal function (Table 2). Ultrafiltration dysfunction and glucose transport were unchanged. Albumin clearance was slightly increased, but this change was not statistically significant.

Discussion

These experiments have allowed us to investigate antifibrotic and antiangiogenic intervention in an acute fibroproliferative model of peritoneal dialysis. This model employed a daily infusion of high-concentration glucose dialysate over a 4-wk period with an initial exposure to LPS and resulted in changes in the peritoneum similar to those seen in patients on long-term peritoneal dialysis. The submesothelium becomes thickened, and there is increased collagen deposition with angiogenesis. We observed associated changes in peritoneal function, including ultrafiltration dysfunction and increased glucose transport. The processes involved in peritoneal changes in patients on chronic dialysis are obviously slower and more progressive. Certain features, such as loss of mesothelium and vasculopathy seen in patients on long-term dialysis, are not seen in our model. The extent to which this daily infusion model mimics what may happen in chronic peritoneal dialysis patients is yet to be determined.

Angiogenesis is an important element of fibrosing diseases (22). We have previously shown AdAngiostatin to effectively reduce angiogenesis in a matrigel model (14) and in a direct tumor injection model (10) and to reduce lung metastases in a murine breast cancer model (14). In this work, we demonstrated a reduction in peritoneal-associated vessels with an

improvement in ultrafiltration dysfunction and decreased glucose absorption after administration of AdAngiostatin. Interestingly, we observed a reduction in submesothelial thickening that was not statistically significant, but no change in hydroxyproline concentration of the mesenteric tissue. These experiments were not designed to measure the effect of antiangiogenic therapy on fibrosis. Previous work has identified an inhibition of lung fibrosis by blocking proangiogenic CXC chemokines (23) or peritoneal adhesions by directly blocking angiogenesis (24). In both these experiments, the total amount of fibrotic tissue was assessed, whereas we measured the concentration of tissue collagen, which was not effected by angiostatin. The nonsignificant reduction of submesothelial thickness may be an indication of reduced total fibrotic tissue and therefore in agreement with previous work noted above. A larger sample size would be necessary to prove this hypothesis in our model.

The importance of angiogenesis in the alteration in peritoneal membrane function in patients on long-term dialysis has become increasingly clear (25). The interstitium of the peritoneum may have a role as a barrier to water reabsorption and larger molecule transport (26), but the capillary wall and the total vascular surface area likely have the greatest impact on transport of small solutes and ultrafiltration. In agreement with this hypothesis, our experiments demonstrated that inhibition of angiogenesis improved ultrafiltration and reduced small solute transport, but inhibition of fibrosis, through treatment with AdDecorin, did not alter peritoneal function. Also, in all animals treated with daily dialysis, we saw a strong inverse correlation between number of blood vessels and net ultrafiltration.

Decorin is a member of the small leucine-rich proteoglycan family (27). Previous work has demonstrated that decorin can alleviate organ fibrosis in several animal models (12,13). This is in agreement with the present experiments, where we show that treatment with AdDecorin in our model significantly reduced hydroxyproline concentration of mesenteric tissue, but did not significantly impact membrane-associated vessels or peritoneal function. We have previously demonstrated that the intraperitoneal delivery of AdTGF- β_1 to the peritoneum increased VEGF expression and induced peritoneal angiogenesis

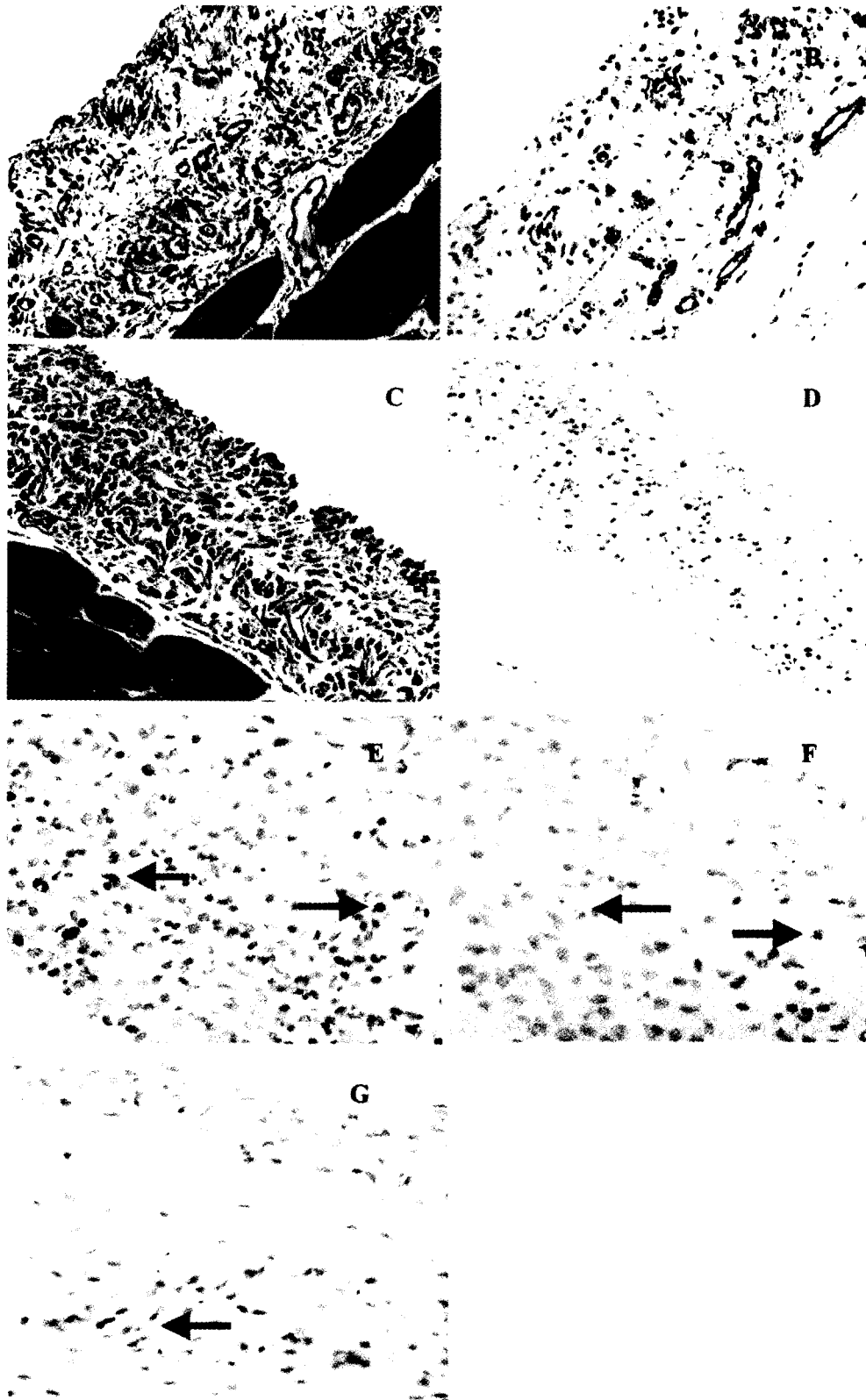


Figure 3. Histology of the anterior abdominal wall. (A) Section from animal treated with AdDL70 and daily dialysate shows submesothelial thickening and collagen deposition with increased vascularization shown by factor VIII-stained section in panel B. (C) Section from animal treated with AdAngiostatin and daily dialysate shows increased thickness and hypercellularity but a significant loss of vascularity, especially noticeable in factor VIII-stained section in panel D. (E) Staining for DNA fragmentation in animals treated with AdAngiostatin shows apoptotic cells, including endothelial cells (arrows) with corresponding factor VIII-stained section shown in panel F. (G) DNA fragmentation in animals treated with AdDL70 and daily dialysis shows virtually no apoptotic cells. Panels A and C are Masson trichrome; panels B, D, F are factor VIII; panels E and G are labeled with terminal deoxynucleotidyl transferase. Magnifications: $\times 100$ in A through D; $\times 200$ in E through G.

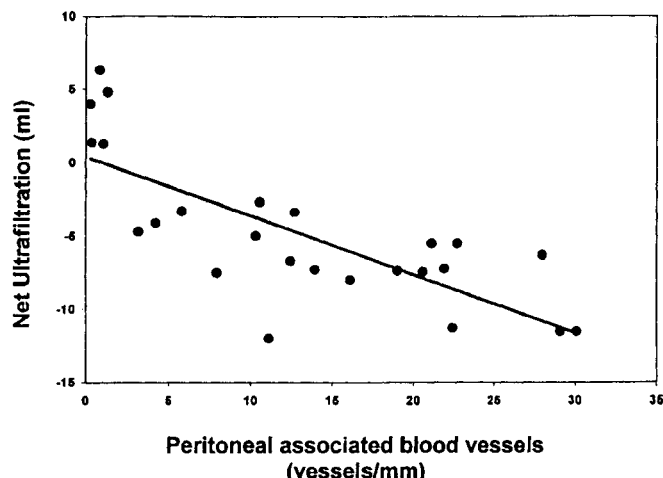


Figure 4. Inverse correlation between peritoneal-associated blood vessels counted on factor VIII-stained sections and net ultrafiltration measured after 4 wk of daily dialysate infusion. $r = -.61$; $P < 0.001$.

(8). There are likely other mechanisms in our model that lead to VEGF expression and angiogenesis, such as indirect induction through inflammatory cytokines, or direct induction through exposure to components of dialysate. TGF- β_1 may have a relatively minor role to play in the induction of angiogenesis in our model. This suggests that fibrosis does not have a significant impact on peritoneal dysfunction and again indicates the importance of the vasculature as a cause of dysfunction in peritoneal dialysis. We did see a nonsignificant increase in albumin transport in animals treated with AdDecorin. It is possible that, through alteration in the composition of the interstitium of the peritoneum that large molecule transport was affected. Further work is required to explore this possibility.

We were unable to completely reverse the effects of peritoneal dysfunction with AdAngiostatin in this model. Clearly, even high and repeated doses of angiostatin were not sufficient to block a substantial component of angiogenesis. This suggests either that the angiogenic stimuli are very strong or that other factors are present which inhibit the ability of angiostatin to induce widespread endothelial apoptosis

We believe that the results from these studies have direct therapeutic implications for patients on peritoneal dialysis. Our findings support the hypothesis that new blood vessel formation is a key component of peritoneal membrane dysfunction. Therapeutic strategies that reduce the peritoneal vasculature should be developed. We noted that creatinine clearance, although decreased in our model compared with untreated animals, was not changed after treatment with AdAngiostatin compared with AdDL70-positive control animals. This suggests that antiangiogenic therapy may not decrease the amount of dialysis provided by the peritoneum. The decreased vascular surface area and small solute transport were balanced by increased dialysate volume; therefore, creatinine clearance was preserved.

Previous work has demonstrated that anti-VEGF antibodies can prevent angiogenesis and peritoneal dysfunction in a hyperglycemic animal model (28). Other strategies targeting vascular growth factors and angiogenesis have been developed for the treatment of cancer and may be applicable to the problem of increased solute transport and ultrafiltration failure in the peritoneum (29). The most effective treatment will likely be prevention. Angiogenesis is likely driven by uremia, inflammation, and nonphysiologic dialysate. Optimizing dialysis treatment to avoid these elements may prevent peritoneal angiogenesis and best protect the peritoneum as a long-term dialysis membrane.

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Table 2. Functional properties of the peritoneal membrane in control animals (no catheter) and animals treated with daily dialysate and adenovirus

Parameter	AdDL70, No Catheter (n = 5)	AdDL70 + Catheter (n = 11)	AdAngiostatin (n = 10)	AdDecorin (n = 7)
Net ultrafiltration (ml)	4.3 ± 1.8	-8.9 ± 3.3 ^a	-3.2 ± 2.7 ^{a,b}	-9.0 ± 3.5 ^a
Glucose transport (mmol/kg)	6.8 ± 0.4	8.0 ± 0.6 ^a	7.1 ± 0.4 ^c	7.7 ± 0.2 ^a
Creatinine clearance (L/g)	49.3 ± 6.8	22 ± 10 ^a	24 ± 8 ^a	22 ± 8 ^a
Albumin clearance (L/g)	1 ± 1.3	2.9 ± 2.0 ^a	4.3 ± 2 ^a	4.9 ± 2.3 ^a

^a $P < 0.01$ compared with AdDL70, no catheter control.
^b $P = 0.0004$ compared with AdDL70 + catheter control.
^c $P = 0.0025$ compared with AdDL70 + catheter control.

- dialysis patients related to dialysis settings and peritoneal transport properties. *Kidney Int* 59(suppl 78): S42-S47, 2001
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DISCUSSION

Peritoneal dialysis (PD) is a widely used, effective therapy for patients with end stage renal disease. Ultrafiltration (UF) dysfunction is a common complication of this therapy caused by systemic factors including uremia and inflammation, and local factors including non-physiological dialysis solutions and recurrent peritonitis. The morphological changes correlating with UF dysfunction include fibrosis and angiogenesis in the peritoneal tissues.

About 14%⁷ of PD patients will have to stop using PD therapy and switch to an alternate renal replacement therapy because of UF failure. More importantly, there is increasing evidence that subtle UF dysfunction goes unrecognized clinically and patients become volume expanded^{24;182} and this leads to hypertension²⁵, hypoalbuminemia²⁷, and poor cardiovascular outcomes. Therefore, fibrosis and angiogenesis in the peritoneal tissues have important functional outcomes in PD patients that may lead to technique failure and increased morbidity and mortality. Solute removal was previously emphasized as a goal for PD therapy¹⁸³. However recent evidence suggests total solute removal is less important in PD outcomes^{184;185}, and the importance of maintaining the peritoneal membrane and maintaining UF function has been recently highlighted¹⁸⁶.

Our work has demonstrated the importance of angiogenesis in UF failure with the observation of an inverse correlation between vascularization of the peritoneal tissues and net UF⁴⁰. More importantly, we demonstrated that the use of adenovirus mediated angiostatin therapy in the chronic daily infusion model of PD led to a reduction in the number of blood vessels, and an improvement in net UF⁴⁰.

This observation not only firmly established the role of angiogenesis in the etiology of UF dysfunction, but it also points towards potential therapies for established peritoneal membrane failure. Adenovirus mediated gene therapy leads to transient gene expression, and so may not be suited for a chronic process such as peritoneal membrane alteration in PD therapy. Safety issues are also a concern. There may, however, still be other forms of gene therapy suitable to the peritoneum to modify both local processes and to deliver longer term systemic therapies ¹¹⁶.

Our demonstration of the efficacy of anti-angiogenic therapy on improving membrane function opens the door for other potential non-gene therapy approaches. Other anti-angiogenic therapies are available and some are being used clinically for applications as novel cancer therapies. There is an interest in using thalidomide in PD patients, both for its anti-inflammatory and anti-angiogenic properties. Other agents such as angiostatin ¹⁸⁷, endostatin ¹⁸⁸, TNP-470 ¹¹⁸, and novel vascular endothelial growth factor (VEGF) receptor tyrosine kinase inhibitors ¹¹⁹ are available and could be tested in an animal model for the impact on peritoneal membrane function.

Previously, the transient effect of peritonitis on peritoneal membrane function was felt to be due to vasogenic factors such as nitric oxide (NO) ⁴⁵ or prostaglandins ¹⁸⁹. Using adenovirus mediated gene transfer of the inflammatory cytokines interleukin (IL)-1 β and tumor necrosis factor (TNF)- α , we have demonstrated a direct angiogenic response in the peritoneum to these agents ⁸¹. A single episode of peritonitis has little ⁷² or no ⁵⁰ effect on long term peritoneal transport whereas severe or recurring peritonitis does ². Our study identified a differential fibrotic, angiogenic, and functional outcome with transient overexpression of TNF α or IL-1 β . IL-1 β exposure led to a prolonged

fibrogenic and angiogenic response with prolonged membrane dysfunction. $\text{TNF}\alpha$ exposure led to a very transient response⁸¹. It has yet to be shown whether severe peritonitis, or peritonitis caused by gram negative organisms, elicits a differential inflammatory response with increased expression of IL-1 and whether this explains the different functional changes observed after an episode of peritonitis. If IL-1 is found to be associated with persistent peritoneal membrane dysfunction in patients as it was in our animal study, then this suggests that combining prompt antibiotic therapy with IL-1 inhibition, perhaps through IL-1 receptor antagonist⁸⁹, may preserve the peritoneal membrane function better than antibiotics alone.

Several previous studies have demonstrated that the profibrotic cytokine transforming growth factor (TGF)- β is likely involved in angiogenesis. TGF β appears to have direct effects through regulation of VEGF, platelet derived growth factor, and fibroblast growth factors, along with secondary vascular modification through angiopoietin (ANG) -1^{160;170}. We have clearly demonstrated the role of TGF β 1 in the angiogenic response. Transient overexpression of TGF β 1 led to significant peritoneal fibrosis associated with increased vascularization¹¹³. This increased angiogenesis was associated with persistent UF dysfunction. TGF β 1 appeared to upregulate VEGF in vivo and in vitro in mesothelial cell culture. These results have recently been replicated in other experimental systems^{169;190}.

Studies of the peritoneum have allowed us to gain some insight into the fibrogenic process. It is clear that TGF β 1 is a key profibrotic cytokine and alone can initiate and maintain peritoneal fibrosis¹¹³. What was also of interest was our observation that, 7 days after peritoneal infection with adenovirus, $\text{TNF}\alpha$ caused a similar upregulation of TGF β 1

and deposition of collagen in peritoneal tissues as IL-1 β ⁸¹. However, at later time points, the fibrotic process was maintained in animals exposed to IL-1 β . We found that tissue inhibitor of metalloproteinase (TIMP)-1 was upregulated by IL-1 β but not by TNF α . TIMP-1 inhibits the metalloproteinases (MMP) which in turn are responsible for collagenolysis. These results suggest that initiation of fibrosis is due to inflammation and TGF β 1, whereas persisting fibrosis is due to an imbalance between collagen deposition and degradation. We have similar findings in an analysis of fibrosis-resistant and fibrosis-prone mice strains¹⁹¹. Fibrosis-resistant mice demonstrate a similar expression of collagen after exposure to TGF β as fibrosis-prone mice, but the expression of TIMP-1 is much lower.

These observations have therapeutic implications. Anti-fibrotic therapy in general has been targeted toward inhibiting the initial inflammatory response to injury or activity of TGF β ¹⁹². Often, this initial phase of the fibrogenic process is not recognized and patients present with established fibrosis. Our findings suggest that, in this setting, altering the collagen metabolic balance towards net collagenolysis may have therapeutic benefits^{81;191}.

In the peritoneum, the formation and modification of collagen in response to cytokine stimulation or injury is not well studied. It is known that the composition of the interstitium has an effect on the hydrodynamic properties of the peritoneum and will affect fluid absorption and macromolecular transport by tissues and lymphatics⁴⁸. Albumin transport appears to correlate inversely with hydroxyproline concentration in the peritoneal tissues. This suggests that fibrosis may impede macromolecular transport. Our model of adenovirus mediated TGF β 1 induced peritoneal fibrosis will allow the further

study of alteration in macromolecular transport, fluid absorption, tissue collagens and hyaluronan concentrations. These studies are important as pharmacologic intervention, such as hyaluronan¹⁹³, may have an effect on the peritoneal interstitium.

The different angiogenic responses we have observed after transient adenovirus mediated cytokine overexpression are intriguing and are a focus of ongoing and future research. While, angiogenesis has been extensively studied in the setting of malignancy, not a great deal is known about angiogenesis in the setting of fibrosis¹⁴³. Various cytokines such as TNF α induce a transient angiogenesis. The blood vessels induced after over expression of TNF α demonstrate increased permeability to macromolecules such as albumin. TGF β 1 and IL-1 β both induce a persisting peritoneal vasculature with comparatively less albumin transport. This suggests that the fibrogenic environment is associated with a more mature vascular phenotype. I therefore suggest that the fibrogenic response is associated with upregulation of ANG-1 which interacts with endothelial cells to reduce susceptibility to apoptosis¹⁹⁴ and decrease macromolecular transport¹⁴⁵. Preliminary data suggests that in vivo, TGF β 1 is associated with increased ANG-1 expression compared with VEGF, which appears to upregulate ANG-2¹⁵⁹. TGF β may also act through the receptor activin receptor like kinase (ALK) -1 which has been shown in knock out mice to be important in blood vessel maturation¹⁷².

This finding has implications for PD research and therapy. We were unable to fully reverse angiogenesis and functional changes in our chronic model even with high doses of adenovirus mediated gene transfer of angiostatin⁴⁰. This may partially be a result of the fibrogenic response generated in this model and the possibility that this could upregulate ANG-1 and protect the vasculature. Decorin, a proteoglycan that binds to and

inhibits TGF β 1, did not have an impact alone on the vasculature but did reduce the collagen content of the peritoneal tissue ⁴⁰. It is possible that the combination of therapy, anti-angiogenic and anti-fibrotic, is necessary to further impact on peritoneal angiogenesis. Also, altering the angiopoietins balance in favor of ANG-2 may make the endothelium more sensitive to the inhibiting effect of angiostatin.

TGF β is associated with poor prognosis in malignancy and this has been thought to be due to its immunomodulating effects or an effect on cell cycle regulation ¹³⁴. More recently, it has been observed that TGF β may have a role in the progression of malignancy through its synergistic effect with hypoxia on VEGF expression and angiogenesis ¹³¹. TGF β has been associated with worse prognosis and increased vascular density in non-small cell lung cancer ¹⁹⁵. Angiopoietins also are involved in the angiogenesis related to malignancy ¹⁹⁶. The effect of TGF β on angiopoietins expression in malignancy has not yet been fully explored.

Adenovirus mediated gene transfer of VEGF has been used to treat ischemic heart disease in humans in several clinical trials ¹⁹⁷. One of the potential side effects has been a vascular leakage phenomenon and the transient nature of the angiogenic response. Future improvements may include combined angiogenic therapy using VEGF and ANG-1 to mature and prolong the vascularization.

Our work has introduced a new animal model for research in PD. Adenovirus mediated gene transfer into the peritoneum is very effective with a transient, high level expression of adenovirus ¹¹³. This technique has been used in combination with a chronic infusion model to demonstrate potential therapeutic options ⁴⁰. Our results have led to insights into the structure and function of the peritoneum as a dialysis membrane and may

eventually lead to improvements in PD therapy that can benefit patients with prolongation of therapy and reduced morbidity and mortality. The importance of the angiogenic response in alteration of the function of the peritoneal membrane has provided a unique opportunity to develop techniques to study blood vessel growth in the setting of fibrogenesis. We have provided some early evidence of the role of TGF β 1 and other inflammatory cytokines in the induction and modification of angiogenesis.

Future work will include further evaluation of the role of TGF β 1 in angiogenesis through its regulation of ANG-1. The peritoneum also provides a simple environment to study the relative contribution of TIMPs and MMPs to the maintenance of fibrosis and the interaction between TIMPs, MMPs and angiogenesis. This model, combined with more detailed functional studies of the peritoneum, will allow the assessment of the relative contribution of vascularization, aquaporin dysfunction, and interstitial / lymphatic reabsorption to UF dysfunction after a fibrogenic stimulus. These results will continue to provide insights and new possible therapies for patients on PD and may provide further insight into angiogenesis in other disease processes.

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