RESEARCH ARTICLE

Strategies for blocking the fibrogenic actions of connective tissue growth factor (CCN2): From pharmacological inhibition *in vitro* to targeted siRNA therapy *in vivo*

David R. Brigstock

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Abstract Connective tissue growth factor (CCN2) is a major pro-fibrotic factor that frequently acts downstream of transforming growth factor beta (TGF-B)-mediated fibrogenic pathways. Much of our knowledge of CCN2 in fibrosis has come from studies in which its production or activity have been experimentally attenuated. These studies, performed both in vitro and in animal models, have demonstrated the utility of pharmacological inhibitors (e.g. tumor necrosis factor alpha (TNF- α), prostaglandins, peroxisome proliferator-activated receptor-gamma (PPAR- γ) agonists, statins, kinase inhibitors), neutralizing antibodies, antisense oligonucleotides, or small interfering RNA (siRNA) to probe the role of CCN2 in fibrogenic pathways. These investigations have allowed the mechanisms regulating CCN2 production to be more clearly defined, have shown that CCN2 is a rational anti-fibrotic target, and have established a framework for developing effective modalities of therapeutic intervention in vivo.

Keywords $CCN2 \cdot Connective tissue growth factor \cdot CCN2 \cdot Transforming growth factor beta \cdot TGF-\beta \cdot Gene transcription \cdot Fibrogenesis \cdot Fibrosis \cdot Therapeutic \cdot Anti-fibrotic \cdot Antisense \cdot Neutralizing \cdot SiRNA \cdot Hepatic fibrosis \cdot CTGF$

D. R. Brigstock (⊠)
The Research Institute at Nationwide Children's Hospital,
700 Children's Drive,

Columbus OH 43205, USA e-mail: David.Brigstock@NationwideChildrens.org

D. R. Brigstock Division of Pediatric Surgery, Department of Surgery, The Ohio State University, Columbus OH 43212, USA

Abbreviations

αSMA	alpha smooth muscle actin
CCl ₄	carbon tetrachloride
CCN2	connective tissue growth factor
COX	cyclo-oxygenase
CVIR	Collagen Type VI receptor
ECM	extracellular matrix
HSC	hepatic stellate cell
PG	prostaglandin
PSC	pancreatic stellate cell
RT-PCR	reverse-transcriptase polymerase chain reaction
siRNA	small interfering RNA
TGF-β	transforming growth factor beta
TIMP	tissue inhibitor of metalloprotease
TNF-α	tumor necrosis factor alpha

Introduction

Since the recognition almost 20 years ago that connective tissue growth factor (CCN2) mRNA is induced in cultured fibroblasts by transforming growth factor beta (TGF- β) (Brunner et al. 1991), there has been a concerted effort to understand the mechanistic basis of this relationship and to explore its in vivo consequences (Leask and Abraham 2003; Shi-Wen et al. 2008; Leask et al. 2009). Nowhere is this more apparent than in the field of fibrosis, which currently represents the most common pathophysiology in which CCN2 has been implicated (Rachfal and Brigstock 2005) and in which there is an extensively documented role for TGF-B as well (Verrecchia and Mauviel 2007). Fibrosis arises due to a failure of the normal wound healing response to terminate, leading to excessive scarring characterized by profound production, deposition, and contraction of extracellular matrix (ECM). This process usually occurs

over many months and years, and can lead to organ dysfunction or death. Key observations have included the following: 1) CCN2 and TGF- β are highly over-expressed and spatio-temporally correlated in numerous fibrotic lesions; 2) CCN2 induces the synthesis and secretion of ECM proteins, notably of fibrillar collagens which are a major component of fibrous deposits; and 3) TGF- β mediated collagen synthesis *in vitro* is blocked by CCN2 antagonists. These observations have been complemented by a careful molecular dissection of the TGF- β -inductive axis and important response elements in the CCN2 promoter have now been identified that are involved in the regulation of CCN2 mRNA expression, although their relative contributions vary according to cell type (Shi-Wen et al. 2008; Leask et al. 2009).

On the other hand, it is important to be mindful of several caveats. Since CCN2 is a multi-modular matricellular protein, it probably resides in the ECM or tethered to the cell surface via integrins, heparan sulfate proteoglycans, and other receptors where it likely acts as a binding partner for other cell regulatory molecules (Chen and Lau 2008). Thus the pro-fibrotic properties of CCN2 may actually be a reflection of its ability to amplify fibrogenic responses to other factors. For example, a binding interaction between the CCN2 and TGF-B proteins has been reported (Abreu et al. 2002) and this association may explain the synergistic interaction of these molecules in mouse models of dermal fibrosis whereas CCN2 alone is unable to drive sustained fibrosis (Frazier et al. 1996; Mori et al. 1999; Chujo et al. 2005). Also, there is no fibrosis in livers or kidneys of mice over-expressing tissue-specific CCN2 transgenes, yet the presence of the CCN2 transgene clearly exacerbates the population of pro-fibrogenic cell types, ECM deposition and/ or expression of fibrogenic markers after the organs undergo injury (Tong and Brigstock 2005; Yokoi et al. 2008). Thus while considerable data support a central role for CCN2 in fibrosis and its induction by TGF- β , we actually have a quite limited understanding of the various signaling events and molecular cross-talk that underlie its fibrogenic actions.

Fibrosis affects millions of people world-wide causing incalculable morbidity and mortality. In the USA it has been estimated that fibrosis of internal organs is a contributing factor in 45% of all deaths, yet there are currently no anti-fibrotics that have been approved for use by the US Food and Drug Administration. Nonetheless, prospects for effective anti-fibrotic therapies have improved rapidly in the last decade and are based on targeting key participants in fibrogenic cascades (Ghiassi-Nejad and Friedman 2008). Hence, it is not surprising that investigators have begun to address the potential therapeutic benefit of targeting CCN2 as an anti-fibrotic strategy (Blom et al. 2002; Leask et al. 2002). A particular attraction of this approach is that CCN2 acts downstream of TGF- β so targeting strategies that focus on preventing CCN2 production or action would avoid potential interference with the important tumor-suppressive and immuno-modulatory actions that are the hallmark of TGF- β function and which have complicated the targeting of TGF- β itself.

This article describes the experimental strategies that have proven effective for blocking CCN2-mediated fibrogenic pathways in cell cultures or animal models and which support further investigations of anti-CCN2 therapy for treating or preventing fibrosis in humans.

Pharmacological inhibitors

Tumor necrosis factor alpha (TNF- α)

TNF- α is mainly recognized as a proinflammatory cytokine but it also has anti-fibrotic effects in vitro and in vivo (Leask and Abraham 2004) leading several investigators to analyze its effect on CCN2 expression. Thus, TNF- α was shown to reduce basal CCN2 expression in bovine aortic endothelial cells, fibroblasts and vascular smooth muscle cells (Dammeier et al. 1998; Lin et al. 1998) as well as in TGF-\beta-stimulated fibroblasts or airway smooth muscle cells (Abraham et al. 2000; Xie et al. 2005; Beddy et al. 2006), dexamethasome-stimulated Balb/c 3 T3 cells (Dammeier et al. 1998) or histamine-stimulated lung fibroblasts (Kunzmann et al. 2007). However, in pancreatic stellate cells (PSC) or mesangial cells, the effect of TNF- α was actually to stimulate CCN2 expression (Cooker et al. 2007; Karger et al. 2008) while it had no effect on constitutive CCN2 expression in scleroderma fibroblasts (Abraham et al. 2000) or glucose-stimulated CCN2 expression in peritoneal mesothelial cells (Sakamoto et al. 2005). While the anti-fibrotic actions of TNF- α were initially attributed to interference of TGF-B pathways either by NF-kB-mediated induction of Smad7 or JNKmediated suppression of Smad 3 (Leask and Abraham 2004), the data now suggest that these pathways are overridden or inoperative under some circumstances in some cell types. Hence the use of TNF- α as a CCN2 inhibitor must therefore be carefully validated for each specific experimental system under investigation.

Prostaglandins (PG)

In fibroblasts, TGF- β or TNF- α induce expression of cyclooxygenase-1 or -2 (COX-1, COX-2) respectively, which catalyze the production of PG from arachidonic acid. A well documented effect of PG in some systems is that of being anti-fibrotic, a property that is attributed to their activation of protein kinase A and elevation of intracellular cAMP levels (Leask and Abraham 2004). Indeed, early studies showed that cAMP blocking agents such as cholera toxin, forskolin or 8-Br-cAMP were effective in preventing TGFβ-induced CCN2 production and anchorage-independent growth in NRK cells (Kothapalli et al. 1998). Forskolin also blocked CCN2 mRNA expression in TGF-\beta-stimulated human lung or renal mesangial cells (Black et al. 2007). Additionally, prostaglandin E2 (PGE2) inhibited TGF-βstimulated CCN2 production in pulmonary fibroblasts or mesangial cells, glucose-induced CCN2 levels in kidney mesangial cells, or TGF-\beta-induced CCN2 production by airway smooth muscle cells or rat-1 cells, the latter of which was mediated via EP-2 receptors (Ricupero et al. 1999; Yu et al. 2002; Makino et al. 2003; Burgess et al. 2006; Black et al. 2007). Iloprost, a synthetic analogue of prostacyclin PGI₂ that is used to help relieve Raynaud's phenomenon in scleroderma patients, elevates cAMP levels and antagonizes the ras/MEK/ERK signaling cascade necessary for induction of CCN2 (Stratton et al. 2001, 2002; Leask et al. 2003), and its inhibitory effect on CCN2 expression has been applied in an in vivo model of liver regeneration to demonstrate the CCN2-dependency of Thy-1+oval cell recruitment (Pi et al. 2005). The suppression of CCN2 or collagen production by 9-cis-retinoic acid in scleroderma fibroblasts is due to its induction of COX-2 and PGE2 expression (Xiao et al. 2008), while all-trans retinoic acid exerted anti-fibrotic effects in the liver and was associated with decreased CCN2 and TGF-B production (Wang et al. 2008). Further, the involvement of ras/ MEK/ERK in CCN2 expression has allowed these pathways to be explored with specific kinase inhibitors such as U0126 or PD98059 (Black et al. 2007; Pickles and Leask 2007; Leask et al. 2008).

It should be stressed that exceptions have been noted in how PG influences CCN2 expression in some experimental systems. For example, the action of PGF2 α in the cat iris is associated with increased CCN2 expression (Liang et al. 2003) while gingival fibroblasts are resistant to PGmediated CCN2 inhibition because the cAMP pathway is poorly coupled to the ras/MEK/ERK pathway and because PGE2 acts via the EP3 receptor to elevate JNK which opposes cAMP-mediated CCN2 inhibition (Black et al. 2007). Thus while PGs and their up- or down-stream mediators are often inhibitory for CCN2 expression, their actual effects must be experimentally validated to take account of cell- or tissuespecific variations in PG responses.

Peroxisome proliferator-activated receptor-gamma (PPAR- γ) agonists

PPAR- γ is a nuclear receptor and a ligand-activated transcription factor that acts in adipocytes as a lipid sensor for fatty acids, eicosanoids, or their metabolites. However, experimental studies of PPAR- γ using the natural ligand

15-deoxy- $\Delta^{12,14}$ prostaglandin J₂ or synthetic ligands such as rosiglitazone, pioglitazone, or GW7845 have shown that PPAR- γ acts in a variety of cell types where it can mediate suppression of basal or TGF-\beta-stimulated collagen synthesis in vitro or in vivo (Ghosh et al. 2004; Burgess et al. 2005; Milam et al. 2008). Accumulating data clearly shows that CCN2 is attenuated in many cell types upon stimulation of PPAR- γ . For example in activated hepatic stellate cells (HSC), TGF-\beta-mediated CCN2 expression was inhibited by PPAR- γ (Sun et al. 2006), the effect of which was increased by treatment of the cells with curcumin which stimulated of PPAR- γ expression (Zheng and Chen 2006). In vascular smooth muscle cells, PPAR- γ inhibited TGF- β - or angiotensin-induced CCN2 expression in smooth muscle cells via Smad-dependent mechanisms (Fu et al. 2001; Gao et al. 2007). Other in vitro experiments have shown that PPAR- γ also inhibits CCN2 expression in renal tubular epithelial cells (Wei et al. 2007a, b), renal fibroblasts (Wang et al. 2007), and hepatocytes (Gressner et al. 2008), the latter of which was promoted by caffeine, suggesting its potential use as an anti-fibrotic (Gressner et al. 2008; Leask 2008). In vivo, administration of PPAR- γ agonoists caused decreased CCN2 production in models of conjuctival scarring and lung injury (Aoki et al. 2008; Yamanaka et al. 2009).

Statins

Statins are a class of cholesterol-lowering drugs that acts by inhibiting HMG-CoA reductase, the rate-limiting enzyme of the mevalonate pathway of cholesterol biosynthesis. However they also have anti-inflammatory, anti-oxidant and immunomodulatory actions, and have also been shown to be anti-fibrotic in models of renal interstitial fibrosis in rats (Li et al. 2004; Vieira et al. 2005). Some of these anti-fibrotic effects may be due, at least in part, to the attentuation of CCN2 expression which has been documented in several different systems and in response to a variety of statins. In vitro, CCN2 expression was inhibited by treatment of mesangial cells, lung fibroblasts or renal fibroblats with simvastatin (Eberlein et al. 2001; Goppelt-Struebe et al. 2001; Heusinger-Ribeiro et al. 2004; Watts and Spiteri 2004; Watts et al. 2005), renal fibroblasts or tenon fibroblasts with lovastatin (Eberlein et al. 2001; Meyer-Ter-Vehn et al. 2008), renal tubular epithelial cells with pravastatin (Wei et al. 2007a), normal rat kidney cells with fluvastatin (Ikeuchi et al. 2004), or rat cardiac fibroblasts with atorvastatin (Martin et al. 2005). Moreover, CCN2 expression is suppressed in various animal models undergoing statin treatment including fluvastatin-treated nephropathy (Song et al. 2004), cerivastatin-treated renal fibrosis (Koepke et al. 2007), atorvastatin-treated vascular fibrosis (Ruperez et al. 2007), pravastin-treated intestinal fibrosis (Haydont et al. 2007), or simvastatin-treated pulmonary fibrosis (Ou et al.

2008). The inhibition of CCN2 expression by statins is attributable to their interference with the Rock-Rho signaling pathway (Eberlein et al. 2001; Watts and Spiteri 2004; Haydont et al. 2007; Ruperez et al. 2007) which has been recognized as a potential anti-fibrotic target (Moriyama and Nagatoya 2004).

Clearly, considerable data has been amassed regarding the pharmacological inhibition of CCN2 that has helped to understand the mechanisms up- or down-stream of CCN2 expression in normal and pathological processes, and it is possible that these findings will help to facilitate or justify the therapeutic use of these compounds for treating human fibrotic disease. On the other hand, these inhibitors generally do not act in a manner that is CCN2specific because they regulate multiple signaling pathways and gene transcription events of which CCN2 is but one component. This lack of specificity is thus a drawback when trying to definitively pin-point processes that are absolutely CCN2-dependent.

Neutralizing antibodies that block CCN2 action

Several CCN2 antibodies, generated by immunization of chickens, rabbits or mice with a variety of CCN2 proteins or peptides, have been reported to have neutralizing activity. These antibodies include the following: polyclonal chicken antibodies raised against full length human CCN2 (Kothapalli et al. 1997); polyclonal antibodies against synthetic peptides corresponding to the domain between module 3 and module 4 including anti-CCN2[244-256] (HDLEENIKKGKKC) raised in chicken (Safadi et al. 2003) and anti-CCN2[240-259] (RPCEADLEENIKKGKK CIRT) raised in rabbit (Shimo et al. 2001); two humanized monoclonal antibodies raised against recombinant human CCN2 with specificity for module 2 which have been termed "JMAb31" (human IgG2) and "FG-3019" (human IgG1k), the latter of which recognizes residues 142-157 (Aikawa et al. 2006; Shimo et al. 2006); and monoclonal antibodies generated by genetic immunization of mice with human CCN2 cDNA that are specific for module 2 or module 3 or module 4 (Ikawa et al. 2008).

In general, these antibodies are not widely available and in only a few instances have they been tested in systems that address the fibrogenic properties of CCN2. More often, they have been used to probe the role of CCN2 in other biological processes such as anchorage-independent growth, tumorigenesis, metastasis angiogenesis, osteogenesis, and chondrogenesis (Kothapalli et al. 1997; Shimo et al. 2001, 2006; Safadi et al. 2003; Aikawa et al. 2006; Dornhofer et al. 2006). Given the structural complexities of the CCN2 molecule, the presence of multiple functional sites (e.g. integrin-binding) in more than one module, its susceptibility

for proteolytic cleavage into smaller bioactive forms, and the broad repertoire of binding partners for CCN2 (the requirement for which in its full spectrum of activities has not been systematically addressed) (Brigstock 1999; Chen and Lau 2008), it is possible that a "one-size-fits-all" blocking antibody may not be technically feasible or capable of yielding definitive data for all of the many different read-outs of CCN2 activity. Indeed, the extent to which any of the aforementioned antibodies prevent CCN2 from stimulating its respective target cells by specific blocking of one or more functional sites in the CCN2 molecule versus "non-specific" steric hindrance has not been reported. While the answer to this question may be inconsequential in terms of exerting a neutralizing effect per se, it is a critical if we are to understand mechanistically the components of CCN2mediated fibrogenic cascades.

While CCN2 neutralizing antibodies have been used in in vitro experiments to demonstrate CCN2-dependency of TGF- β -induced collagen production (Duncan et al. 1999; Blalock et al. 2003), there are only a handful of published reports that support their use in fibrosis research in vivo. FG-3019 has been reported to show efficacy in models of renal fibrosis in rodents (Flyvbjerg et al. 2004; Wang et al. 2004a) and is currently under evaluation for phase I trials of kidney or lung fibrosis in humans (Mageto et al. 2004; Adler et al. 2006). Also genetic immunization of human CCN2 cDNA in mice yielded several monoclonal antibodies that reduced TGF-\beta-dependent fibrosis and collagen deposition in a mouse skin model, the most potent of which was targeted to module 2 (Ikawa et al. 2008). Thus the published data obtained with neutralizing CCN2 antibodies are very limited but do provide evidence for a role of CCN2 in fibrosis. Nonetheless, while these data suggest that module 2 is highly immunogenic and is involved somehow in fibrosis (Aikawa et al. 2006; Shimo et al. 2006; Ikawa et al. 2008), we must be mindful of the fact that forms of CCN2 comprising either module 3 alone or module 4 alone are bioactive and fibrogenic (Gao et al. 2004; Gao and Brigstock 2004, 2005, 2006; Tong and Brigstock 2006). Clearly we have much to learn about the functional domains in CCN2 and how they act independently or inter-dependently to support or drive fibrosis.

The lack of readily accessible and/or validated CCN2 neutralizing antibodies represents a huge frustration in the field. To circumvent this problem, many investigators have instead focused on antagonism of CCN2 mRNA using complementary DNA or RNA sequences - for which the reagents have been easier to develop and validate among the various laboratories involved. Thus, the role of CCN2 mRNA expression in fibrogenic pathways and its TGF- β -dependency or -independency has been investigated successfully and extensively using CCN2 antisense oligonucleotides or small interfering RNA (siRNA).

In vitro use of CCN2 antisense oligonucleotides or siRNA

The use of CCN2 antisense oligonucleotides has helped to establish the role of CCN2 in TGF- β -induced collagen production in several cell types such as kidney mesangial cells, normal rat kidney cells, corneal fibroblasts, and conjuctival fibroblasts (Duncan et al. 1999; Yokoi et al. 2001, 2002; Blalock et al. 2003; Kanemoto et al. 2003; Yamanaka et al. 2008). Similar *in vitro* studies have shown reduced collagen synthesis in scleroderma fibroblasts (Xiao et al. 2006), reduced ECM production in vascular smooth muscle cells (Liu et al. 2007), or antagonism of TGF- β - stimulated collagen production in dermal fibroblasts, osteoblasts or rat HSC (Wang et al. 2004b; Arnott et al. 2007; Li et al. 2008) after treatment of the target cells with CCN2 siRNA. Plasmids expressing short hairpin RNA were shown to disrupt CCN2 gene expression in cultured rat HSC and to cause marked attenuation in the production of collagens III and IV, laminin, and hyaluronic acid (Yuhua et al. 2008). Finally, hammerhead ribozymes designed to cleave CCN2 mRNA blocked TGF- β -mediated proliferation of dermal fibroblasts (Blalock et al. 2004) as well as basal or TGF- β stimulated collagen synthesis and entry into S phase of human HSC (D.R.B and R. Gao, unpublished observations).

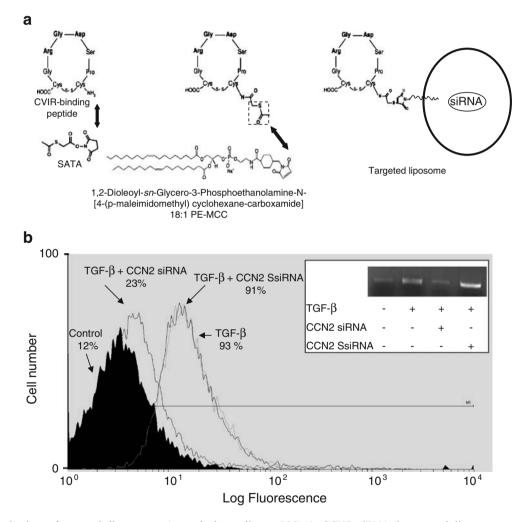


Fig. 1 a Production of targeted liposomes. A synthetic cyclic C*GRGDSPC* peptide was modified at its free N-terminus with the bifunctional linker succinimidyl-S-acetylthioacetate (SATA), the thiol group of which allowed for subsequent coupling to a maleimide linker in a modified lipid formulation (1,2-dioleoyl-*sn*-Glycero-3-Phosphoethanolamine-N-[4-(p-maleimidomethyl)cyclohexane-carboxamide]; 18:1 PE-MCC). Targeted liposomes were loaded with mouse CCN2 siRNA (sense: 5'-CCGCAAGAUUGGCGUGUGCtt; antisense: 5'-GCACACGCCAAUCUUGCGGtt; nucleotides 546-

566). **b** CCN2 siRNA in targeted liposomes attenuates TGF-βstimulated CCN2 production in mouse HSC *in vitro*. Day2 primary mouse HSC were transfected for 6 hours with either 2µM CCN2 siRNA or scrambled siRNA (SsiRNA) using targeted lipsomes. On the next day, the cells were stimulated with 20 ng/ml TGF-β for 24 hours and then analyzed for CCN2 protein or mRNA by, respectively, flow cytometry or RT-PCR (inset) as described (Leask et al. 2008; Lawrencia et al. 2009). CCN2 siRNA, but not SsiRNA, was effective in blocking TGF-β-induced CCN2 mRNA and protein

In vivo use of CCN2 antisense oligonucleotides or siRNA

Several reports have supported the efficacy of anti-CCN2 mRNA strategies in treating hepatic fibrosis *in vivo*. In carbon tetrachloride (CCl₄)-induced liver fibrosis in rats,

subcutaneous administration of anti-sense CCN2 oligonucleotides resulted in decreased hepatic mRNA levels of CCN2 and collagen type I mRNA levels, although the amount of hepatic fibrosis was not actually reduced, possibly due to high expression of tissue inhibitor of metalloprotease

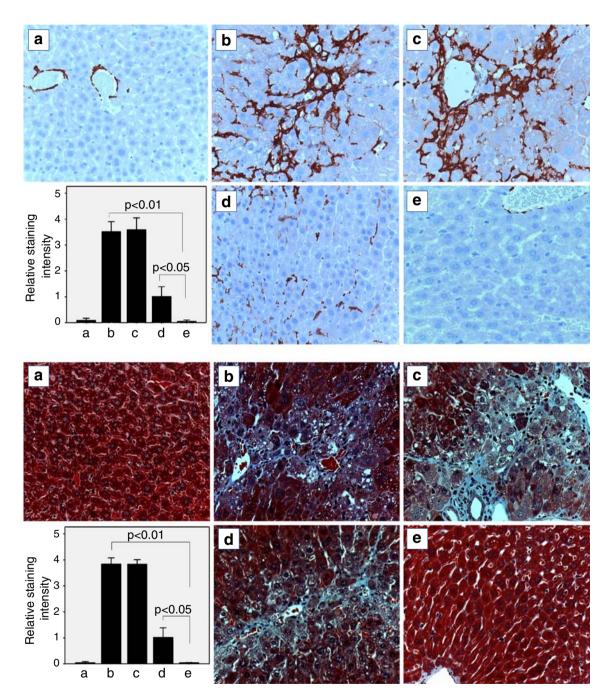


Fig. 2 Targeted CCN2 siRNA suppresses accumulation of α SMA or collagen deposition in preventative fibrosis model. Livers were sectioned and processed to detect α SMA (upper panel; brown stain) or collagen deposition (lower panel; blue stain) after administration of oil **a** or CCl₄ for 3 weeks **b**-e, the latter 2 weeks of which were given concurrent with either no therapy **b**, scrambled CCN2 siRNA in targeted liposomes **c**, CCN2 siRNA in non-targeted liposomes **d**,

or CCN2 siRNA in targeted liposomes (E). CCl₄ was given i.m. as four 30μ l doses (1:1 CCl₄ : olive oil) per week. siRNA treatments (0.1 mg/kg) were given 4 times a week by i.p. injection. The staining intensity was assessed by image analysis (50 sections; 10 sections from each of 5 mice) and is shown in the bar graph. Data shown are indicative of the data from 3 separate experiments

(TIMP)-1 (Uchio et al. 2004). On the other hand, intraportal vein injection of CCN2 siRNA in CCl_4 -treated rats reduced hepatic expression of CCN2, collagen Types 1, 2, 3, laminin, TIMP-1, and TGF- β , reduced the number of activated HSC

as assessed by α SMA staining, and further reduced serum procollagen Type 3, hepatic hydroxyproline and liver fibrosis staging (Li et al. 2006). A very similar improvement was also seen in N-nitrosodimethylamine-induced hepatic fibrosis in

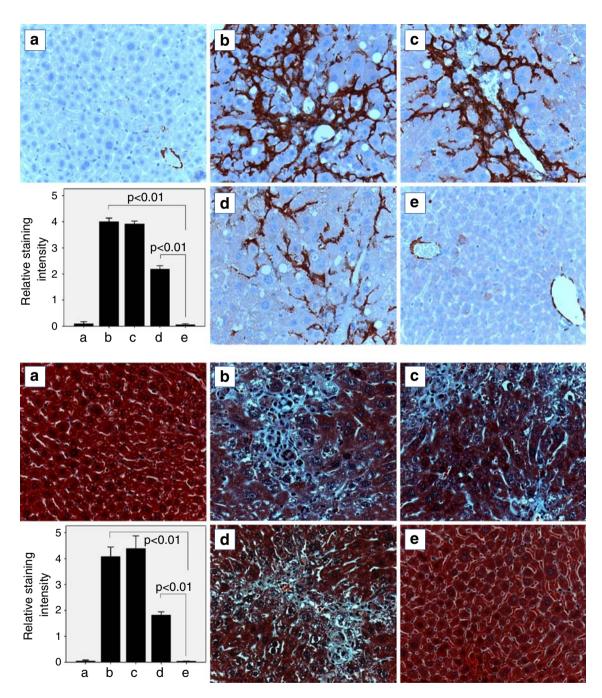


Fig. 3 Targeted CCN2 siRNA prevents accumulation of α SMA or collagen deposition in curative fibrosis model Livers were sectioned and processed to detect α SMA (upper panel; brown stain) or collagen deposition (lower panel; blue stain) after administration of oil **a** or CCl₄ for 5 weeks **b-e**, the latter 2 weeks of which were given concurrent with either no therapy **b**, scrambled CCN2 siRNA in targeted liposomes **c**, CCN2 siRNA in non-targeted liposomes **d**, or

CCN2 siRNA in targeted liposomes e. CCl₄ was given i.m. as four 30 μ l doses (1:1 CCl₄:olive oil) per week. siRNA treatments (0.1 mg/kg) were given 4 times a week by i.p. injection. The staining intensity was assessed by image analysis (50 sections; 10 sections from each of 5 mice) and is shown in the bar graph. Data shown are indicative of the data from 3 separate experiments

rats for which CCN2 siRNA was found to be effective in both preventative and curative fibrosis models (George and Tsutsumi 2007).

Similar outcomes have been reported in other model systems, most notably for kidney fibrosis. For example, in a unilateral ureteral obstruction model of renal fibrosis in the rat, blocking the activity of CCN2 by antisense oligonucleotides reduced the induction of CCN2 and, in turn, ECM genes and their corresponding proteins (Yokoi et al. 2004). CCN2 antisense oligonucleotides also significantly blocked CCN2 expression and renal interstitial fibrogenesis in a remnant kidney model of sub-totally nephrectomized mice transgenic for TGF- β , despite sustained expression of TGF- β (Okada et al. 2005). Other *in vivo* studies have shown that CCN2 siRNA delivery to rats is an effective antifibrotic therapy in renal allografts after transplantation (Luo et al. 2008). Finally, CCN2 antisense oligonucleotides have proven effective in reducing ECM expansion and scarring around breast implants (Mazaheri et al. 2003).

Clearly, we have witnessed the advent of an era in which unequivocal anti-fibrotic effects can be achieved by blocking CCN2 mRNA *in vivo*. This line of investigation is undoubtedly in its infancy but shows great promise in that some remarkable and reproducible therapeutic effects have been reported despite the use of different model systems.

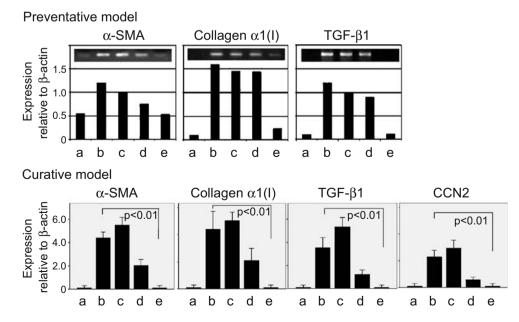
New data from the author's laboratory: Targeted CCN2 mRNA silencing *in vivo*

Previous studies have shown that the cyclic peptide C*GRGDSPC*, which selectively binds to collagen Type VI receptors (CVIR) on activated HSC (Marcelino and

McDevitt 1995), can be used as a homing device to target to HSC in fibrotic livers (Beljaars et al. 2000). This approach has attracted attention because it provides for a targeting strategy whereby anti-fibrotic agents can be delivered more efficiently to the highly pro-fibrotic HSC despite their relatively low abundance. Recently, it was reported that delivery of interferon- α 1b in liposomes coated with a slightly modified form of the CVIR-binding peptide was delivered highly efficiently to activated HSC in vivo and was anti-fibrotic in a bile duct ligation model of liver fibrosis in rats (Du et al. 2007). In concurrent experiments performed in this laboratory, a CVIR-binding cyclic peptide was covalently bound to the surface of modified liposomes (Fig. 1a) and used for delivery of CCN2 siRNA in mouse models of CCl₄-induced liver fibrosis (Lawrencia and Brigstock 2008). Prior to performing the in vivo experiments, the efficacy of the targeted liposomes in reducing CCN2 mRNA expression and protein production in primary mouse HSC was verified by reverse-transcriptase polymerase chain reaction (RT-PCR) and flow cytometry (Fig. 1b). For in vivo studies, Balb/c mice received an i.m. injection of CCl₄ on four consecutive days each week for either 1 week (preventative model) or 3 weeks (curative model). For the next 2 weeks, mice continued to receive CCl₄ under the same dosing regimen plus mouse CCN2 siRNA which was delivered 4 times a week by i.p. injection, in either targeted or non-targeted liposomes. Control mice received either oil instead of CCl₄ or a scrambled CCN2 siRNA sequence in the targeted liposomes.

In both the preventative (Fig. 2) and curative (Fig. 3) models, immunohistochemical staining for α SMA (a marker of activated HSC) and trichrome staining for collagen revealed very low levels of each molecule in control oil-

Fig. 4 Expression of fibrosisrelated genes. Livers were removed from mice after administration of oil a or CCl4 b-e for 3 weeks (preventative model; upper panel) or 5 weeks (curative model; lower panel). For the last 2 weeks in each model mice received no therapy b. scrambled CCN2 siRNA in targeted liposomes c, CCN2 siRNA in non-targeted liposomes d or CCN2 siRNA in targeted liposomes e. RNA was isolated and subjected to RT-PCR to establish the expression level of the indicated fibrotic marker genes as described (Lawrencia et al. 2009). Data are plotted relative to the expression of *β*-actin



treated animals (Figs 2, 3, panel a) and massive amounts of α SMA and collagen (bridging fibrosis) in response to CCl₄ (Figs 2, 3, panel b) as expected. This staining pattern remained unaltered in the presence of targeted scrambled CCN2 siRNA (Figs 2, 3, panel c). The CCl₄-induced levels of α SMA and collagen were diminished by non-targeted antisense CCN2 siRNA (Figs 2, 3, panel d) but the targeted antisense CCN2 siRNA approach was considerably more efficacious in that α SMA and collagen levels were reduced to background levels (Figs 2, 3, panel e) and essentially indistinguishable from those of oil treated animals.

As assessed by RT-PCR, the expression of key fibrotic markers was strongly attenuated by the targeted CCN2 therapy in both fibrosis models. The data for the preventative and curative models (Fig. 4) showed that the transcript levels for collagen $\alpha 1(I)$, α SMA, TGF- β , or CCN2 were highly induced by CCl₄ treatment as expected. These expression levels remained high in the presence of targeted scrambled CCN2 siRNA as expected and were only modestly reduced by the non-targeted CCN2 siRNA. However, the targeted CCN2 siRNA caused the mRNA levels of collagen $\alpha 1(I)$, α SMA, TGF- β or CCN2 to be reduced to background levels and were therefore indistinguishable from oil-treated controls. These data were thus consistent with the histological findings in each model (Figs 2, 3)

On the one hand, these data show that non-targeted liposomes carrying CCN2 siRNA do exhibit therapeutic efficacy as also recently demonstrated in N-nitrosodimethylamineinduced hepatic fibrosis in rats (George and Tsutsumi 2007), and this presumably reflects the fact that the liver is one of the principal organs in which liposomes tend to accumulate after injection (Ostro and Cullis 1989). On the other hand, the targeted liposomes were more effective than non-targeted liposomes in either preventing or reversing fibrosis and attenuating expression of fibrosis-related genes even in the continued presence of the fibrotic insult. While these findings reinforce a central role for CCN2 hepatic fibrosis, the fact that higher levels of CVIR are produced by HSC in fibrotic injury and were the presumptive binding site for the targeted peptide in these studies suggest that CCN2 of HSC origin may be particularly important in driving hepatic fibrosis in vivo. The data additionally highlight the potential versatility of using targeted liposomes to deliver other therapeutic siRNAs to HSC in injured liver. Indeed, a similar system was recently described in which vitamin A-coated liposomes were shown to preferentially bind to HSC and to reverse acute and chronic fibrosis and prolong survival in rats upon delivery of siRNA to rat gp46, a collagen chaperone that is homologous to human heat shock protein 47 (Sato et al. 2008). Targeted strategies have also been described for other organ systems such as the kidney

(Prakash et al. 2008) but it is uncertain whether this degree of sophistication will in fact be necessary for the successful implementation of anti-fibrotic therapy in humans (Friedman 2008).

Conclusions

A variety of CCN2 antagonistic strategies have been developed in numerous experimental systems that demonstrate a central role for CCN2 in fibrogenesis and which, in turn, have shown that targeting of CCN2 expression or action is a rational therapeutic approach in fibrotic pathologies. A diverse repertoire of techniques has been described that are effective in blocking CCN2 but it is important to appreciate their individual limitations and to be mindful of alternative approaches for studying CCN2 mechanisms. For example, CCN2-null mice are amenable to elegant exploration of CCN2-dependent pathways in vivo (Ivkovic et al. 2003; Chuva de Sousa Lopes et al. 2004; Kuiper et al. 2007, 2008; Nishida et al. 2007; Kawaki et al. 2008a, b), and the analysis of specific cell types derived from these CCN2-deficient animals hold promise for further definition of CCN2-dependent mechanisms in vitro, including those related to fibrosis (Chen et al. 2004; Shi-wen et al. 2006; Kennedy et al. 2007; Kawaki et al. 2008a; Mori et al. 2008; Pala et al. 2008; Crawford et al. 2009). Nonetheless, the ability to attenuate CCN2 production in fibrotic tissues remains a reasonable and realistic goal. With respect to antifibrotic strategies in humans, the challenges ahead will be to establish which fibrotic pathologies, if any, are best suited to CCN2 targeting (possibly in combination with targeting of other fibrogenic mediators) and to determine the optimal form(s) that this therapy should take.

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