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Dipeptidylpeptidase 4 as a Marker of Activated Fibroblasts and a Potential Target for the Treatment of Fibrosis in Systemic Sclerosis — Source link

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Abstract: BACKGROUND Dipeptidyl-peptidase-4 (DPP4) identifies a dermal fibroblast lineage involved in scaring during wound healing. The role of DDP4 in tissue fibrosis, however, is unknown. The aim of the present study was to evaluate DPP4 as a potential target for the treatment of fibrosis in systemic sclerosis (SSc). METHODS The expression of DPP4 was analyzed by real-time PCR, immunofluorescence and Western blot. The activity of DPP4 was modulated by overexpression, knockdown and pharmacological inhibition using Sitagliptin and Vildagliptin. The effects of DPP4 inhibition were analyzed in human dermal fibroblasts and in different mouse models of SSc (n=6). RESULTS The expression of DPP4 and the number of DPP4 positive fibroblasts were increased in fibrotic skin of SSc patients in a TGFdependent manner. DPP4 positive fibroblasts expressed higher levels of myofibroblast markers and collagen (p<0.001). Overexpression of DPP4 promoted fibroblast activation, whereas pharmacological or genetic inactivation of DPP4 reduced proliferation, migration, expression of contractile proteins and release of collagen by interfering with TGF--induced ERK signaling (p<0.001). DPP4-knockout mice were less sensitive to bleomycin-induced dermal and pulmonary fibrosis (p < 0.0001). Treatment with DPP4 inhibitors promoted regression of fibrosis induced by bleomycin- or chronic graft-versus-host disease and ameliorated fibrosis in TSK1 mice (p < 0.001). The antifibrotic effects were associated with reduced inflammation. CONCLUSION DPP4 characterizes a population of activated fibroblasts and regulates TGF--induced fibroblast activation. Inhibition of DPP4 exerts potent anti-fibrotic effects in well tolerated doses. These results may have direct translational implications as DPP4 inhibitors are already in clinical use for diabetes. This article is protected by copyright. All rights reserved.

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Dipeptidyl-peptidase-4 as a marker of activated fibroblasts and a potential target for the treatment of fibrosis in Systemic Sclerosis

Running Title: Dipeptidyl-peptidase-4 in SSc

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Abstract

Background: Dipeptidyl-peptidase-4 (DPP4) identifies a dermal fibroblast lineage involved in scaring during wound healing. The role of DDP4 in tissue fibrosis, however, is unknown. The aim of the present study was to evaluate DPP4 as a potential target for the treatment of fibrosis in systemic sclerosis (SSc).

Methods: The expression of DPP4 was analyzed by real-time PCR, immunofluorescence and Western blot. The activity of DPP4 was modulated by overexpression, knockdown and pharmacological inhibition using Sitagliptin and Vildagliptin. The effects of DPP4 inhibition were analyzed in human dermal fibroblasts and in different mouse models of SSc (n=6).

Results: The expression of DPP4 and the number of DPP4 positive fibroblasts were increased in fibrotic skin of SSc patients in a TGF- β dependent manner. DPP4 positive fibroblasts expressed higher levels of myofibroblast markers and collagen (p<0.001). Overexpression of DPP4 promoted fibroblast activation, whereas pharmacological or genetic inactivation of DPP4 reduced proliferation, migration, expression of contractile proteins and release of collagen by interfering with TGF- β -induced ERK signaling (p<0.001). DPP4knockout mice were less sensitive to bleomycin-induced dermal and pulmonary fibrosis (p<0.0001). Treatment with DPP4 inhibitors promoted regression of fibrosis induced by bleomycin- or chronic graft-versus-host disease and ameliorated fibrosis in TSK1 mice (p<0.001). The antifibrotic effects were associated with reduced inflammation. **Conclusion:** DPP4 characterizes a population of activated fibroblasts and regulates TGF- β induced fibroblast activation. Inhibition of DPP4 exerts potent anti-fibrotic effects in well tolerated doses. These results may have direct translational implications as DPP4 inhibitors are already in clinical use for diabetes.

Systemic sclerosis (SSc) is a chronic fibrotic disease, that is associated with the highest case specific mortality of all connective tissue diseases (1). The central histopathologic hallmark of SSc is the uncontrolled and persistent activation of fibroblasts, which release excessive amounts of extracellular matrix (2). Fibroblasts are key effector cells in fibrotic diseases. Fibroblasts are, however, not a uniform population of cells, but compose of functionally and phenotypically different subsets. Emerging evidence highlights that chronic inflammatory diseases including rheumatoid arthritis are associated with dysbalances in the ratio of different fibroblast subsets and that those shifts may drive disease progression (3-7). The complexity of different fibroblast subsets in fibrotic diseases such as SSc is less well understood. However, it is well established that fibroblasts in fibrotic diseases can acquire an activated, so-called myofibroblast phenotype (8, 9). Although transforming growth factor β (TGF- β) has emerged as a core pathway of fibroblast activation in SSc and in other fibrotic diseases, the molecular mechanisms underlying the persistent activation in fibroblasts remain incompletely understood (10).

Dipeptidyl-peptidase-4 (DPP4, also known as CD26) exists either as a type II transmembrane protein or as a soluble form (11, 12). DPP4 functions as serine protease that hydrolyses proline or alanine from the N-terminus of a broad range of polypeptides (13, 14). DPP4 inactivates incretin hormones such as glucagon-like peptides or glucose-dependent insulinotropic peptide to inhibit insulin secretion and to promote diabetes mellitus. These findings have been successfully translated from bench-to-bedside and DPP4 inhibitors are currently widely used for the treatment of diabetes mellitus (15). However, the substrates of DPP4 are not restricted to incretin hormones, but include also a broad range of other soluble mediators such as chemokines. Moreover, DPP4 can modulate intracellular signaling not only via proteolytic cleavage of substrates, but also via direct interaction with key-regulatory molecules such as CD45 or adenosine deaminase, playing immune regulatory functions (16, 17). DPP4 has been playing a costimulatory role in T cells, to promote T- and B cell activation (13, 18). Treatment with DPP4 inhibitors lowered the incidence of autoimmune disorders in patients with diabetes mellitus (19). Of particular interest, Rinkevich and coworkers recently demonstrated that DPP4 expression identifies an embryonic fibroblast lineage with unique functions in wound healing (20). They demonstrated that DPP4expressing fibroblasts accounted for the bulk of connective tissue deposition upon surgical wounding. Inhibition of DPP4 reduced scaring without negatively affecting wound closure, suggesting that DPP4 inhibition may offer an avenue for selective targeting of a fibroblast population with high capacity of matrix production, while not affecting other populations with more homeostatic functions (20). However, the concept of a DPP4-expressing subpopulation of active, matrix-producing fibroblasts has not yet been translated and validated in the context of SSc.

In the present study, we aimed to characterize the role of DPP4 in fibroblast activation and tissue fibrosis. We demonstrate that 1.) DPP4 marks a population of activated fibroblasts, increased in SSc and in experimental fibrosis in a TGF- β dependent manner; 2.) DPP4 regulates fibroblast activation and collagen release *in vitro* and *in vivo*; 3.) pharmacological inhibition of DPP4 with approved drugs induces regression of pre-established fibrosis in well tolerated doses, and 4.) despite consistent anti-inflammatory effects across different murine models, the anti-fibrotic effects are mainly mediated on DPP4 expression in tissue-resident cells such as fibroblasts.

Additional, more detailed information on Material and methods can be found in the supplement of this manuscript.

Patients and cells

Skin biopsies were obtained and dermal fibroblasts were isolated from 23 SSc patients and 21 healthy volunteers matched for sex and age. All SSc patients fulfilled the ACR/EULAR criteria for SSc (21). According to LeRoy classification of systemic sclerosis, 15 (65.2%) patients were diffuse and 8 (34.7%) patients belonged to the limited cutaneous subset of SSc. Sixteen patients were female, seven were male. The median age of patients with SSc was 52.7 ± 12.3 years (mean \pm SD), and their median disease duration was 4.3 ± 2.2 years. 5 (21.7%) patients were treated with immunosuppressive drugs at the time of biopsy (low dose steroids, methotrexate) or had received in the past cyclophosphamide or rituximab. All patients and healthy volunteers signed an informed consent approved by the local institutional review board.

Serum samples were collected from 50 SSc (53.8% diffuse cutaneous SSc) patients fulfilling the ACR/EULAR criteria and 30 healthy controls. The median age of SSc patients was 55.6 \pm 13.4 years (mean \pm SD) with a mean disease duration of 7.2 \pm 3.4 years (mean \pm SD). 11 (22,0%) patients were under treatment with immunosuppressive drugs.

In addition to human fibroblasts, murine fibroblasts were isolated from DPP4 knockout (DPP4-KO) mice and wildtype (WT) littermates (22).

Western blot analysis

Proteins were separated by SDS-PAGE and transferred to a polyvinylidene difluoride membrane. The membranes were incubated with antibodies against SMAD3 (Santa Cruz Technologies, Heidelberg, Germany), pp44/p42 MAPK (Erk1/2), p44/p42 MAPK (Erk1/2), SAPK/JNK, pp38 MAPK. p38 MAPK, pAkt, Akt (pan), pNF-kB p65, NF-kB p65, pSRC (Cell Signaling, Boston, USA) or DPP4, pJnk (Abcam, Cambridge, UK) primary antibody and HRP-conjugated secondary antibodies (Dako, Hamburg, Germany). Blots were visualized by ECL. β-actin was used as loading control (23).

Inhibition of canonical and non-canonical TGF-β pathways

To investigate the effect of TGF- β non-canonical pathway on DPP4 expression, dermal fibroblast were incubated with the ERK inhibitor FR180204 (10µM) (Tocris Bioscience, Bristol, UK), the JNK inhibitor SP600125 (80µM) (Tocris Bioscience), the p38 inhibitor SP202190 (1µM) (Tocris Bioscience), the AKT inhibitor GSK690693 (0.1µM) (Tocris Bioscience), the SRC inhibitor SU6656 (500nM) (Tocris Bioscience), the NF-kB inhibitor Bengamide B (0.1µM) (Tocris Bioscience), the ABL inhibitor imatinib (1µg/ml) (Novartis, Basel, Switzerland), or the JAK inhibitor Ruxolitinib (5µM) (LC Laboratories, Boston, USA) for 4 hours before stimulation with TGF β (10ng/ml) (24, 25). To block the canonical pathway, SIS3 (3µM, Sigma-Aldrich, Steinheim, Germany) was used. SD208 was used to inhibit the TGF- β receptor I-kinase activity (26, 27). Total RNA was isolated with the NucleoSpin RNA II extraction system (Machery-Nagel, Düren, Germany) and reverse transcribed into complementary DNA (cDNA) with random hexamers.

Immunohistochemistry and immunofluorescence staining

Formalin-fixed, paraffin-embedded human or murine skin sections or fibroblasts fixed in 4% PFA were stained with antibodies against αSMA (Life Technologies, Darmstadt, Germany), DPP4, vimentin or prolyl-4-hydroxidase, CD45, CD3 and B220 (all from Abcam), pERK (Cell Signaling). HRP-conjugated or Alexa Fluor antibodies (Life Technologies) were used as secondary antibodies. Irrelevant isotype matched antibodies served as controls (9). Nuclei were counterstained using DAPI (Santa Cruz). The staining was analyzed using a Nikon Eclipse 80i microscope (Nikon, Badhoevedorp, Netherlands).

Mouse models of SSc

Bleomycin-induced skin fibrosis - preventive and therapeutic dosing

DPP4-KO mice and wildtype (WT) littermates were injected subcutaneously with bleomycin or 0.9 % sodium chloride (NaCl), for four weeks (28, 29). Another group of mice was treated in parallel with the DPP4 inhibitor, Sitagliptin 10mg/kg (Selleckchem, Houston, USA), throughout the four weeks of bleomycin-challenge.

In the therapeutic, fibrosis was first induced by injection of bleomycin for three weeks (30). Thereafter, treatment with Sitagliptin 3mg/kg and 10mg/kg and Vildaglipin 1.5mg and 15mg/kg (Biomol, Hamburg, Germany) was initiated for another three weeks, while mice were further challenged with bleomycin. The outcome was analyzed after six weeks after the first injection of bleomycin. Mice injected with 0.9% sodium chloride served as controls.

Sclerodermatous chronic graft-versus-host disease (cGvHD)

In the B10.D2 \rightarrow Balb/c (H-2(d)) minor histocompatibility antigen-mismatched model, cGvHD was induced by allogeneic transplantation of 5×10⁶ splenocytes and 1×10⁶ bone marrow cells from B10.D2 mice into sub-lethally irradiated BALB/c (H-2d) mice. BALB/c

(H-2d) mice transplanted with splenocytes and bone marrow cells isolated from BALB/c (H-2d) mice served as controls (31, 32). Treatment started after appearance of first clinical signs of cGvHD at day 21 post transplantation, and the outcome was analyzed after six weeks.

Bleomycin-induced pulmonary fibrosis

DPP4-KO and WT mice were intratracheally injected with bleomycin or 0.9 % NaCl (27). Mice were sacrificed 4 weeks after injection. A subgroup of mice was treated with Sitagliptin in doses of 10mg/kg from days 1 - 28.

Generation of bone marrow chimeric mice

Mixed-bone marrow chimeras were generated by transplanting bone marrow from DPP4-KO mice into WT mice or vice versa. Before transplantation, recipient mice (DPP4-KO or WT-mice) were sub-lethally irradiated (33). Bone marrow cells were isolated from tibial and femoral bones of DPP4 KO donor mice and injected into WT mice (KO \rightarrow WT mice) or DPP4-KO (KO \rightarrow KO). Similarly, bone marrow cells from WT mice were injected into DPP4-KO mice (WT \rightarrow KO) and into wildtype mice (WT \rightarrow WT). Fibrosis was induced by subcutaneous or intratracheal injections of bleomycin 10 days after bone marrow transplantation.

Histological, biochemical and immunohistochemical analyses of the extent of fibrosis

The extent of fibrosis was analyzed using histological, radiologic, biochemical, and immunohistochemical readouts. Histologic readouts included quantification of the dermal thickness on Hematoxylin Eosin (HE) stained sections at eight sites at 100-fold magnification (34), evaluation of the fibrotic area as percent of total lung area in Sirius Red stained sections (35); quantification of pulmonary changes using the Ashcroft score (36) and direct

visualization of collagen by trichrome staining (27). The total collagen content was analyzed biochemically using hydroxyproline assays. In addition, myofibroblasts were identified immunohistochemically as α -smooth muscle actin (α SMA) positive.

Statistics

All data are presented as median \pm interquartile range, and differences between the groups were tested for their statistical significance by paired student t-tests for related samples and Mann-Whitney U non-parametric test for non-related samples. P-values less than 0.05 were considered significant. P-values are expressed as follows: 0.05 > p > 0.01 as *; 0.01 > p >0.001 as **; p < 0.001 as ***.

Results

DPP4 expression is increased in SSc fibroblasts

We observed increased expression of DPP4 in the skin of SSc patients compared to matched healthy individuals (Fig. 1A). Co-staining with prolyl-4-hydroxylase- β (P4H) demonstrated that fibroblasts express high levels of DDP4. DPP4 was also expressed in B and T cells of SSc patients, but fibroblasts were the dominant cell type expressing DPP4 in SSc skin (Suppl. Fig. 1). In SSc, 75.8% (\pm 7.8%) of P4H-positive fibroblasts were stained for DPP4, whereas only 29.1% (\pm 8.3%) of fibroblasts in healthy skin expressed DPP4 (Fig. 1A). Significantly more DPP4-positive fibroblasts co-expressed alpha smooth muscle actin (α SMA) as compared to DPP4-negative fibroblasts, indicating that DPP4 marks a subpopulation of activated fibroblasts (Fig. 1B). We also observed increased expression of DPP4 in murine models of SSc such as bleomycin-induced skin fibrosis (Fig. 1C) and the B10.D2 (H-2^d) \Rightarrow BALB/c (H-2^d) model of sclerodermatous chronic graft versus host disease (cGvHD) with

prominent staining of DPP4 in fibroblasts (Fig. 1C). Moreover, the protein level of DPP4 is elevated in SSc fibroblasts compared to dermal fibroblasts from healthy individuals (Fig. 1B). The levels of soluble DPP4 in the serum did not differ between patients with limited cutaneous and diffuse cutaneous SSc and healthy controls (Suppl. Fig. 2A). The enzymatic activity of DPP4 also did not differ between serum samples of SSc patients and controls.

TGF-β induces DPP4 expression in fibroblasts via ERK

Incubation of normal human dermal fibroblasts with recombinant TGF-B mimicked the DPP4 expression pattern of SSc fibroblasts with increased protein levels of DPP4 after 24 and 48 h (Fig. 2A), but normal mRNA levels (data not shown). We next analyzed, whether stimulation with TGF- β may induce the enzymatic activity of DPP4 in human dermal fibroblasts. Indeed, stimulation of SSc fibroblasts with recombinant TGF- β for 24 h upregulated DPP4 activity and this increase in activity correlated with the increase in DPP4 protein levels (Suppl. Fig. 2B). Consistent with the results in vitro, we observed increased expression of DPP4 in skin fibroblasts of mice with fibroblast-specific overexpression of a constitutively active TGF-B receptor type 1 (TBR^{act}) compared to control mice (Fig. 2A). Moreover, treatment of bleomycin-challenged mice with SD208, a specific inhibitor of the TGF-B receptor I-kinase activity, prevented the bleomycin-induced upregulation of DPP4 protein (Fig. 2A). To identify which intracellular signaling cascades mediate the stabilization of DPP4 protein by TGF-B, we knocked down SMAD3 by siRNA in fibroblasts. However, knockdown of SMAD3 did not inhibit the stimulatory effects of TGF- β on DPP4 expression (Fig. 2B). Comparable results were obtained by incubation with the SMAD inhibitor SIS3. We thus tested the role of various non-canonical TGF-β pathways using specific inhibitors against various non-canonical intracellular mediators of TGF-B. Inhibition of ERK kinases ameliorated the stimulatory effects of TGF- β on DPP4 expression, whereas inhibition of

SRC, cABL, JAK, AKT, p38, NfkB and JNK did not interfere with TGF-β-induced DPP4 expression (Fig. 2C).

DPP4 regulates fibroblast activation and collagen release

To investigate the functional role of DPP4 in fibroblast activation, we first sorted DPP4expressing and DPP4-negative fibroblasts from the skin of mice (Suppl. Fig. 3A) and analyzed the transcription of key profibrotic genes. DPP4-expressing fibroblasts expressed higher levels of *Col1a1*, *Col1a2* and *Acta2* mRNA (which encodes for αSMA) than DPP4negative fibroblasts. Similar results were obtained when DPP4-positive and DPP4-negative fibroblasts were isolated from the lungs (Suppl. Fig. 3B).

Further, we compared the fibrotic potential of fibroblasts isolated from DPP4-knockout mice (DPP4-KO fibroblasts) to that isolated from wildtype control littermates. DPP4-KO fibroblasts were less responsive to the stimulatory effects of TGF- β . Fibroblast-to-myofibroblast transition was impaired in DPP4-KO fibroblasts with reduced expression of α SMA and impaired formation of stress fibers upon stimulation with TGF- β as compared to control fibroblasts (Fig. 3A). Moreover, TGF- β failed to induce the mRNA levels of *Col1a1*, *Col1a2* and *Acta2* or to increase the release of collagen in murine DPP4-KO fibroblasts (Fig. 3A). DPP4-KO fibroblasts also demonstrated delayed closure of the gap in scratch assays as compared to control fibroblasts (Suppl. Fig. 4). Consistently, treatment with the DPP4 inhibitor sitagliptin inhibited the stimulatory effects of TGF- β on murine fibroblasts (Fig. 3A).

Overexpression of DPP4 in human fibroblasts increased the mRNA levels of *ACTA2*, *COL1A1* and *COL1A2* and the levels of collagen protein secreted into the supernatant (Suppl. Fig.5). Incubation of SSc fibroblasts with sitagliptin ameliorated TGF-β-induced fibroblast-

to-myofibroblast transition, prevented upregulation of *COL1A1* and *COL1A2* mRNA by TGF- β and reduced the release of collagen from human dermal fibroblasts (Fig. 3B).

We next aimed to characterize, how DPP4 inhibition interferes with TGF- β -induced fibroblast activation. We thus analyzed the effects of DPP4 inhibition on canonical and noncanonical TGF- β pathways that have been implicated in the pathogenesis of fibrotic diseases. Treatment of human dermal fibroblasts with sitagliptin prevented the stimulatory effects of TGF- β on ERK signaling leading to decreased levels of phosphorylated ERK (pERK) in human dermal fibroblasts (Fig. 3B). However, inhibition of DPP4 did not interfere with the TGF- β -induced SMAD3, STAT3, SRC, AKT signaling or with cJUN and FRA2 mediated AP1 signaling (Suppl. Fig. 6). Consistent results were also obtained in DPP4-KO cells (data not shown). Moreover, upon DPP4 overexpression levels of pERK increased, while pSMAD3 levels did not change, confirming the hypothesis that DPP4 exerts part of its effects via noncanonical TGF- β signaling pathway (Suppl. Fig.7).

DPP4-KO mice are protected from experimental dermal and pulmonary fibrosis

To investigate whether the inhibitory effects of DPP4 inactivation on fibroblast activation *in vitro* translate into anti-fibrotic effects *in vivo*, we evaluated the role of genetic and pharmacological inactivation of DPP4 in bleomycin-induced pulmonary and dermal fibrosis. DPP4-KO mice appear phenotypically normal and the histological architecture of the lungs and the skin were not altered under homeostatic conditions (Fig. 4A). However, DPP4-KO mice were less sensitive to bleomycin-induced fibrosis. Pulmonary fibrosis induced by intratracheal injections of bleomycin was significantly ameliorated in DPP4-KO mice with reduced fibrotic area, decreased Ashcroft scores, less pronounced fibrotic changes on CT, reduced myofibroblast counts and decreased hydroxyproline content as compared to control littermates (Fig. 4A).

DPP4-KO mice were also protected from skin fibrosis induced by subcutaneous injections of bleomycin with reduced dermal thickness, impaired myofibroblast differentiation and decreased hydroxyproline content (Fig. 4B).

In wildtype mice, preventive treatment with sitagliptin 10mg/kg, initiated together with the intratracheal or subcutaneous bleomycin-challenge, also improved all fibrotic readouts and strongly ameliorated bleomycin-induced pulmonary and dermal fibrosis. However, sitagliptin did not exert additional anti-fibrotic effects in DPP4-KO mice (Fig. 4), demonstrating that the anti-fibrotic effects of sitagliptin are indeed mediated by DPP4 inhibition and not by off-target effects.

Pharmacological inhibition of DPP4 induces regression of pre-established experimental fibrosis

Using a pharmacological approach in murine SSc models, we first employed the model of bleomycin-induced skin fibrosis. We did not use preventive dosing schedules, but started treatment with the DPP4 inhibitors sitagliptin and vildagliptin only after fibrosis has already been established (Fig. 5A). Both DPP4 inhibitors, in both doses strongly ameliorated bleomycin-induced dermal thickness, accumulation of myofibroblasts and the hydroxyproline content compared to vehicle-treated mice (Fig. 5A). DPP4 inhibition also induced regression of pre-established fibrosis when comparing to pre-treatment level (Fig. 5A). Further, consistent with the *in vitro* data, pERK expression was reduced upon treatment with DPP4 inhibitors in bleomycin challenged mice (Suppl. Fig. 8).

To further validate the anti-fibrotic effects of pharmacological DPP4-inhibition, we next analyzed the effects of sitagliptin and vildagliptin in the B10.D2 \rightarrow Balb/c (H-2(d)) model of sclerodermatous cGvHD (Fig. 5B). Treatment with sitagliptin or vildagliptin, initiated after occurrence of first clinical signs, reduced cGvHD-induced dermal thickening, collagen

deposition and myofibroblast differentiation (Fig. 5B). DDP4 inhibition also reduced the cGvHD-induced weight loss (Suppl. Fig. 9).

In addition to these two inflammation-driven models, we evaluated the antifibrotic effect of Sitagliptin in the Tsk-1 model of fibrosis as a less inflammation-dependent model of SSc. Sitagliptin significantly ameliorates fibrosis in Tsk-1 mice with reduced hypodermal thickness, myofibroblast count and hydroxyproline content as compared to vehicle-treated Tsk-1 mice. Treatment of Tsk-1 mice with Sitagliptin also significantly reduced the mRNA levels of *Colla1*, of *Colla2* and of *Acta2* (Suppl. Fig. 10).

Anti-inflammatory effects of DPP4 inhibition

DPP4 expression has been shown to be implicated in the regulation of B and T cell activity (37). Treatment with sitagliptin and vildagliptin decreased total leukocytes counts and reduced B and T cell numbers in the skin of mice with bleomycin-induced dermal fibrosis and with sclerodermatous cGvHD (Fig. 6A), demonstrating that DPP4 inhibitors also ameliorated inflammation in addition to its direct effects on fibroblasts.

We next aimed to characterize the contribution of DPP4 inactivation in leukocytes to the antifibrotic effects of DPP4 inhibition. Therefore, we generated chimeric mice by transplanting bone marrow from DPP4-KO mice in WT-littermates (DPP4-KO \rightarrow WT mice with selective knockout of DPP4 in bone marrow derived cells such as leukocytes) and vice versa (WT \rightarrow DPP4-KO mice with inactivation of resident, non-hematopoietic cells) (Fig. 6B) and induced fibrosis by intratracheal or subcutaneous injections of bleomycin, respectively. While the extent of pulmonary and dermal fibrosis of DPP4-KO \rightarrow WT was comparable to that of WT \rightarrow WT control mice, fibrosis was strongly ameliorated in WT \rightarrow DPP4-KO mice. Of note, all readouts of pulmonary and dermal fibrosis in WT \rightarrow DPP4-KO mice were comparable to that of DPP4-KO→DPP4-KO mice (Fig. 6B and Suppl. Fig. 11), characterizing resident cells such as fibroblasts as major target cells for the anti-fibrotic effects of DPP4 inhibitors.

Discussion

We demonstrate in the present study that DPP4 expression is increased in SSc patients and in different murine models of skin fibrosis. We found no difference in the serum levels of DPP4 between healthy controls and SSc patients, as previously shown (38, 39), suggesting that DDP4 expression is regulated locally in fibrotic tissues. Although DPP4 is not specifically expressed in fibroblasts, co-staining with fibroblast markers demonstrated that fibroblasts are the predominant cell type expressing DPP4 in SSc skin and that the majority of SSc fibroblasts in situ are positive for DPP4. Of particular interest, we demonstrate that DPP4 expression marks a population of activated fibroblasts. DPP4-positive fibroblasts in SSc skin showed increased expression of prototypical myofibroblast marker aSMA compared to DPP4-negative cells. Moreover, dermal and pulmonary fibroblasts expressing DPP4 demonstrated increased transcription of type I collagens and Acta2 as compared to DPP4 negative fibroblasts from the same mice. A recent landmark study by Rinkevich et al. reported a DPP4-positive fibroblast population in embryonic skin that possess high fibroproliferative potential that expands upon tissue injury to promote wound healing (20). The findings of Rinkevich et al. and our results together suggest that a subpopulation of DPP4-positive fibroblasts expand to drive persistent tissue remodeling and tissue fibrosis in SSc. However, further studies with lineage tracing experiments are required to further confirm this conclusion.

We provide evidence that TGF- β is a factor that may drive the expansion of DPP4-positive fibroblasts. We demonstrate on multiple experimental levels that non-canonical TGF- β signaling is stimulating DPP4 expression: 1.) Stimulation of cultured dermal fibroblasts with

studies.

recombinant TGF- β upregulated the expression of DPP4 protein in normal dermal fibroblasts. 2.) DPP4 levels were increased in SSc fibroblasts as compared to fibroblasts isolated from healthy individuals. 3.) Overexpression of TBRI^{act} increases DPP4 expression in the skin of mice, thus confirming that TGF- β is sufficient to increase DPP4 expression in fibroblasts *in vitro* and *in vivo*. 4.) Inhibition of the non-canonical TGF- β signaling mediator ERK, inhibits the stimulatory effects of TGF- β on DPP4 expression. 5.) Selective inhibition of TGF- β signaling prevented the upregulation of DPP4 in experimental fibrosis, highlighting that TGF- β signaling is required for the overexpression of DPP4 in experimental fibrosis.

The upregulation of DPP4 had direct functional consequences and promoted activation of certain non-canonical TGF- β pathways in fibroblasts. Inactivation of DPP4 reduced the TGF- β -induced activation of ERK signaling in cultured fibroblasts as well as experimental fibrosis. ERK is an important intracellular mediator of TGF- β , which is activated in SSc and targeted inhibition of ERK has been shown to ameliorate experimental fibrosis (40, 41). Other intracellular cascades regulated by TGF- β were not affected by DPP4 inhibition. The molecular mechanisms underlying the selective regulation of ERK by DPP4 require further studies.

Consistent with the central role of TGF- β signaling in fibrogenesis, the inhibitory effects of DPP4 on TGF- β signaling directly translated into inhibition of fibroblast activation. Inactivation of DPP4 blocked TGF- β -induced fibroblast-to-myofibroblast differentiation and reduced the release of collagen *in vitro*. Genetic or pharmacologic inhibition of DPP4 also ameliorated experimental dermal and pulmonary fibrosis induced by bleomycin or by sclerodermatous cGvHD. Moreover, inactivation of DPP4 was shown to ameliorate CCL4-induced liver fibrosis and cardiac remodeling after high-salt diet-induced heart failure (42-44). Targeted inhibition of DPP4 was also shown to reduce scar formation after cutaneous wounds. The potent anti-fibrotic effects of DPP4 inhibitors may have direct translational

implications: 1.) Pharmacologic inhibition of DPP4 did not only prevent further progression of fibrosis, but also induced regression of pre-established fibrosis to below pre-treatment levels. 2.) Potent anti-fibrotic effects were already observed with the lower doses of both DPP4 inhibitors in mice, implying that standard doses as used for the treatment of diabetes mellitus could be effective in fibrotic diseases such as SSc. 3.) Anti-fibrotic doses of DPP4 inhibitors are well tolerated and their application is not limited by adverse events in our preclinical models. 4.) DPP4 inhibitors are widely used for the treatment of diabetes, offering multiple drug candidates for further clinical studies.

We provide evidence that DPP4 inhibition does not only target fibroblast activation directly, but also reduces inflammation. Treatment with DPP4 inhibitors reduced leukocyte counts and in particular T cell and B cell infiltration in murine models of SSc; both of which are centrally involved in the pathogenesis of SSc (1, 45). Indeed, DPP4 has been shown to regulate Th2 polarization and regulate B cell activation (46-49). Despite potent effects on B-and T cell infiltration into fibrotic tissues, our bone marrow transplantation experiments actually demonstrated that the pro-fibrotic effects of DPP4 predominantly required DPP4 expression in tissue resident cells such as fibroblasts.

DPP4 inhibitors are already in clinical use for the treatment of type 2 diabetes mellitus for more than 10 years. The adverse effects include in particular arthralgia or arthritis, but also hypersensitivity, skin-related reactions and pancreatitis. The Food and Drug Administration (FDA) released a warning in 2015 that DPP4 inhibitors may cause joint pain. However, different studies showed no increased risk of arthritis in patients treated with DPP4 inhibitors compared to other second-line antidiabetics. Postmarketing events of hypersensitivity reactions like anaphylaxis and angioedema have been reported in patients treated with DPP4 inhibitors. However, more detailed studies revealed a similar incidence of angioedema in patients treated with sitaglipin compared to placebo (50). Similar results were reported also for other DDP4 inhibitors such as saxagliptin.

In out experiments, mice treated with DDP4 inhibitors did not show evidence of adverse events on clinical monitoring or on necropsy, including no evidence of arthritis or angioedema.

In summary, we provide evidence that DPP4 characterizes a population of activated fibroblasts in SSc. However, DPP4 does not only serve as an activation marker, but is also functionally required for fibroblast activation and tissue fibrosis. Targeted inactivation of DPP4 exerted potent anti-fibrotic effects in different models of experimental dermal and pulmonary fibrosis. These results may have direct translational implications as DPP4 inhibitors are already in clinical use for diabetes.

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CONTRIBUTIONS

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Figure Legend

Figure 1. DPP4 expression is increased in systemic sclerosis and murine models of SSc.

A Immunohistochemistry staining of DPP4 in skin of patients with SSc and healthy volunteers and immunofluorescence (IF) staining of DPP4 and P4H and its quantification (n=9 for each). Representative images are shown at 200- and 600-fold magnification. **B** Expression of DPP4 in α SMA positive cells and quantification (n=6 for each). Representative images are shown at 400-fold. DPP4 protein in SSc and healthy skin analyzed by Western blot and its quantification (n=6 for each). **C** DPP4 expression in bleomycin-induced (bleo) skin fibrosis analyzed by IF microscopy and Western blot. Representative images are shown at 400-fold (n=5 for each). DPP4 expression in murine fibroblasts of sclerodermatosus chronic graft versus host (cGvHD) disease model analyzed by IF microscopy and Western blot and its quantification (n=5 for each). Representative images are shown at 600-fold. Nuclei are stained with DAPI (blue). Results are shown as mean \pm SEM. *p≤0.001, ***p≤0.001 as determined by Mann-Whitney test.

Figure 2. Induction of DPP4 by TGF-β.

A Expression of DPP4 upon stimulation with TGF- β analyzed by Western blot and its quantification (n=5 for each). DPP4 protein level in murine skin overexpressing TGF- β receptor type I (TBRI^{act}) assessed by Western blot and its quantification (n=5 for each). Effects of treatment with selective TBRI inhibitor SD208 on DPP4 protein level in bleomycin induced skin fibrosis analyzed by Western blot and its quantification (n=5 for each). **B** Effect of SMAD3 knockdown on TGF- β induced expression of DPP4 and its quantification (n=5 for each). **C** TGF- β induced DPP4 protein level upon inhibition of non-canonical TGF- β pathway by SRC, ABL, JNK, JAK, NF κ B, AKT, p38, ERK and SMAD inhibitors analyzed by Western blot and its quantification of TGF- β receptor I kinase activity with SD208 served as positive control. Results are shown as mean ± SEM. *p≤0.05, **p≤0.001, ***p≤0.0001 as determined by Mann-Whitney test.

Figure 3. Inactivation of DPP4 inhibits fibroblast activation und collagen release.

A Murine DPP4-KO fibroblasts: Representative images (shown at 200-fold magnification) and quantification of α SMA and stress fibers in WT and DPP4-KO fibroblasts upon stimulation with TGF- β (n=5 for each). Nuclei are stained with DAPI (blue). mRNA levels of *Col1a1, Col1a2, Acta2* mRNA and collagen release in WT and DPP4-KO fibroblasts upon stimulation with TGF- β (n=5 for each). Murine fibroblasts with pharmaceutical inhibition of DPP4: *Col1a1, Col1a2, Acta2* mRNA levels and collagen release (n=5 for each). B Pharmacological inhibition of DPP4 in human dermal fibroblasts. Representative images (at 200-fold magnification) and quantification of α SMA and stress fibers in human dermal fibroblasts treated with sitagliptin upon TGF- β stimulation (n=5 for each). Treatment with Sitagliptin decreases the stimulatory effects of TGF- β on *COL1a1* and *COL1a2* mRNA and

Figure 4. DPP4-KO mice are partially protected from experimental bleomycin-induced pulmonary and dermal fibrosis.

A Bleomycin-induced pulmonary fibrosis: Representative images of Sirius Red, hematoxylin and eosin (HE)-stained sections (at 200-fold magnification) and high resolution computed tomography (CT) scans of the lungs. Quantification of the fibrotic area (n=6 for each), Ashcroft score (n=6 for each) and density quantification on CT in Hounsfield Units (HU) (n=6 for each). Myofibroblast counts and (n=6 for each) hydroxyproline content of the lung (n=6 for each). B Bleomycin-induced dermal fibrosis: Representative images of HE-stained sections of the bleomycin-induced skin fibrosis mouse model shown at 200-fold magnification. Dermal thickness quantification (n=8 for each). Results are shown as mean \pm SEM. p≤0.05, **p≤0.001, ***p≤0.0001 as determined by Mann-Whitney test.

Figure 5. Pharmacological inhibition of DPP4 induces the regression of bleomycin- and cGvHD-induced experimental fibrosis.

A Bleomycin-induced dermal fibrosis: Treatment scheme. Representative images of hematoxylin and eosin (HE) - stained sections of murine skin shown at 100 fold magnification (n=6 for each) and quantification of dermal thickness (n=6 for each), myofibroblast counts (n=6 for each) and hydroxyproline content of the skin (n=6 for each). B Sclerodermatous cGvHD: Experimental outline. Representative images of HE-stained sections of murine skin shown at 100 fold magnification. Dermal thickness quantification (n=6 for each), myofibroblast counts (n=6 for each) and hydroxyproline content (n=6 for each). Results are shown as mean \pm SEM. *p \leq 0.05, **p \leq 0.001, ***p \leq 0.000 as determined by Mann-Whitney test.

Figure 6. Anti-inflammatory effects of DPP4 inhibition in the lung

A Numbers of CD45, B220 and CD3 positive cells in bleomycin-challenged mice treated with DPP4 inhibitors (n=6 for each). Numbers of CD45 and B220 positive cells in cGvHD mice treated with DPP4 inhibitors. **B Chimeric mice:** Generation of mixed-bone marrow (BM) chimeras from DPP4 knockout or wildtype mice. Representative images of Sirius red staining, trichrome staining and high resolution computed tomography (CT) scans of the lungs of the chimeric mice (n=6 for each). Quantification of the fibrotic area (n=6 for each), Ashcroft score (n=6 for each) and fibrotic changes on CT-scans measured in Hounsfield Units (HU) (n=6 for each). Hydroxyproline quantification (n=6 for each) and myofibroblast counts (n=6 for each). Results are shown as mean \pm SEM. *p≤0.05, **p≤0.001, ***p≤0.000 as determined by Mann-Whitney test.



DPP4+/ P4H* cells (rel. exp.)

DPP4* / Vimentin* cells (rel. exp.)

2.5

2.0

1.5

1.0

0.0

2 0.5

synBMT

Healthy

SSc

DPP4 + / aSMA + cells (rel. exp.)

DPP4* / Vimentin* cells

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synBMT alloBMT

synBMT alloBMT

DPP4

β-actin

*

alloBMT

DPP4 (rel. exp.)

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Bleo

DPP4

β-a

Bleo







