



Lack of Syndecan-1 promotes the pathogenesis of experimental rheumatoid arthritis

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Abstract

Syndecan-1 (Sdc-1), a transmembrane heparan sulfate protein, is implicated in several pathophysiological processes including rheumatoid arthritis (RA). The exact role of Syndecan-1 in this autoimmune disease is still undetermined. This study explores the involvement level of Sdc-1 in the development of RA in a collagen II–induced arthritis mice model. RA was induced in two mice strains (wild-type BALB/c group and Sdc-1 knockout) by collagen II. Mice underwent regular clinical observations and scoring. After sacrifice, leg biopsies were taken from mice for histological examination, using a variety of stains. In addition, proteins were extracted, and molecular assessment of TNF- α was performed using the western blot technique. In the Sdc-1 knockout group, clinical scoring results showed a significantly more severe experimental RA; histology showed a significant increase in bone erosion, cartilage destruction, inflammation, and less granulated mast cells than the wild-type. In addition, molecular assessment of TNF- α showed more increase in expression in the Sdc-1 knockout models compared to the wild-type. Data suggest that lack of Sdc-1 enhances the inflammatory characteristics in RA. However, more molecular studies and investigations are needed to determine its exact role and possible mechanisms involved.

Keywords Syndecan-1 · Rheumatoid arthritis · Chronic inflammation · Autoimmune diseases

Introduction

Inflammation as a process can be both beneficial and detrimental depending on its duration and localization (Furman et al. 2019). It has been documented that acute inflammation after injury or infection is necessary as a defense mechanism for the body; however, a chronic inflammation that occurs in a healthy tissue can lead to the occurrence of many inflammatory diseases contributing to more than half of deaths worldwide (Furman et al. 2019). Rheumatoid arthritis (RA), the most common inflammatory arthropathy, falls in this category of inflammatory diseases, where several molecules, cells, and tissues are known to be involved in the pathophysiologic process. RA is an autoimmune systemic disorder of

unknown etiology, characterized by chronic inflammation and synovial infiltration of immune cells (Shen et al. 2021). The RA pathogenic process is quite complex involving synovial cell proliferation, fibrosis, and pannus formation, which is a large abnormal fibrovascular tissue, along with cartilage and bone erosion (Lui et al. 2022). Several inflammatory factors are involved in the pathogenesis. TNF- α , a potent cytokine involved in normal immune response, is one of the major factors present in RA, where its high levels in the synovial fluid enhance the inflammation and cause joint destruction (Vasanthi et al. 2007).

On the other hand, Syndecan-1 (Sdc-1) belongs to a family of transmembrane proteoglycans found on the surface of many cells, predominantly expressed on epithelial cells and endothelial cells, among others (Agere et al. 2018). It has been shown to play a role in numerous cell processes including inflammation, whereby it acts as an inhibitor of inflammation, in non-infectious inflammatory diseases, leading to a reduction in the expression and the activity of pro-inflammatory factors (Götte 2003). Previous studies conducted by our team have demonstrated the involvement of Sdc-1 in wound healing and regeneration, whereby

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Table 1 Induction of collagen-induced arthritis in BALB/c wild-type and Sdc-1 null mice

Wild-type	Sydecan-1 null
Group 1 Tail injection on day 0: 75 µl CII + 75 µl CFA Intraperitoneal booster injection on day 21: 75 µl CII + 75 µl PBS	Group 4 Tail injection on day 0: 75 µl CII + 75 µl CFA Intraperitoneal booster injection on day 21: 75 µl CII + 75 µl PBS
Group 2 Tail injection on day 0: 100 µl CII + 100 µl CFA Intraperitoneal booster injection on day 21: 100 µl CII + 100 µl PBS	Group 5 Tail injection on day 0: 100 µl CII + 100 µl CFA Intraperitoneal booster injection on day 21: 100 µl CII + 100 µl PBS
Group 3 Tail injection on day 0: 100 µl CFA Intraperitoneal booster injection on day 21: 100 µl PBS	Group 6 Tail injection on day 0: 100 µl CFA Intraperitoneal booster injection on day 21: 100 µl PBS
Group 7 (normal control) Non-injected	Group 8 (control) Non-injected

CII collagen II, CFA complete Freund's adjuvant, PBS phosphate-buffered saline

fibroblasts lacking Sdc-1 migrate faster than usual and fail to interact appropriately with the extracellular matrix to initiate healing (Stepp et al. 2010). Other studies showed that leukocytes lacking Sdc-1 interact more with endothelial cells, leading to an increase in the inflammatory response, thus potentially enhancing the incidence of RA (Teng et al. 2012). However, the effect of Sdc-1 on inflammatory factors is not well understood.

This study explored the involvement of Sdc-1 in the development of rheumatoid arthritis with collagen-induced arthritis in BALB/c normal mice and BALB/c Sdc-1 knockout mice. Such investigations shed light on the role of Sdc-1 in modulating and downregulating the inflammatory pathogenic process in RA and its possible mechanism of action.

Materials and methods

Animals

The BALB/c Sdc-1 knockout mice were obtained from the Department of Anatomy and Regenerative Biology, George Washington University, DC, USA. Breeding was performed at the Animal Care Facility of the American University of Beirut, Beirut, Lebanon. The entire experiment was approved by the Institutional Animal Care and Use Committee (IACUC# 19–09-552) of the American University of Beirut.

Rheumatoid arthritis: collagen II model

A total of 24 mice (12 BALB/c wild-type and 12 Sdc-1 KO), 8–10 weeks old, were divided into 8 groups according to Table 1. They were tail-injected with different concentrations of collagen II (CII) and complete Freund's adjuvant (CFA) to

induce arthritis (Williams 2004): Complete Freund's adjuvant (CFA) (7009, Lot 190,447) and bovine type II collagen (20,022, Lot 190,494) were purchased from Chondrex (Woodinville, WA 98072, USA). A two-dose regimen, 3 weeks apart, was followed according to Table 1. The first injection was intravenous and consisted for most groups of a mixture of CII and complete Freund's adjuvant (CFA), which is a solution containing inactivated mycobacteria used as an adjuvant to stimulate the immune system. An intraperitoneal booster injection was delivered on day 21, replacing CFA with phosphate-buffered saline (PBS). The controls in each category were injected only with CFA or PBS, without collagen II. Mice lived on normal diet.

Clinical signs and symptoms

Mice were monitored three times a week for 70 days, and clinical signs of redness and swelling were recorded according to the scale in Table 2 (Gaballah et al. 2022).

Histological analysis

After sacrifice, peripheral joints were taken as biopsies. Part of each joint was frozen for molecular study, while the other part was used for histological examination after 10% formalin fixation.

Table 2 Arthritis score

Score	Characteristics
0	Normal
1	Slight swelling or erythema
2	Pronounced swelling
3	Joint rigidity

Histological examination was conducted, according to standard routine procedures, to explore the joint alterations in mice. Five-micrometer-thickness tissue sections were used and four different staining methods were performed: the routine hematoxylin and eosin stain (Fischer et al. 2018), Masson Trichrome stain for the detection of connective tissue (collagen) (Rieppo et al. 2019), Safranin O stain for the identification of cartilage and bones (Zu et al. 2019), and toluidine blue stain used to stain mast cells, proteoglycans, and glycosaminoglycans in tissues such as cartilage (Raja et al. 2021). The assessment was done using Microscope Olympus CX41 for slide imaging, and selected slides were photographed.

Molecular assessment of TNF-α

Protein extraction

Mice paws were crushed into powder and put in Laemmli lysis buffer containing protease and phosphatase inhibitors.

The mix was centrifuged for 10 min at 4 °C and at 9000 rpm. The supernatant was collected and stored in – 80 °C freezers.

Western blotting The Folin-Lowry assay was used for protein quantification. Proteins were resolved by 12% polyacrylamide gel electrophoresis and were transferred to nitrocellulose membranes. The membranes were blocked in non-fat milk (5%) for 1 h at room temperature and incubated for 24 h at 4 °C with primary antibodies against β-actin (1:500; human/mouse; Santa Cruz Biotechnology, Dallas, TX, USA) and TNF-α (1:500; human/goat; R&D, Minneapolis, MN, USA). After washing three times for 10 min each with TBST, the membranes were incubated with the corresponding secondary antibodies at room temperature for 1 h and washed three times with TBST. Protein levels were determined by enhanced chemiluminescence (Bio-Rad Laboratories, Hercules, CA, USA) according to the manufacturer’s instructions. Band intensities were quantified using ImageJ software (<https://imagej.net/ij/>).

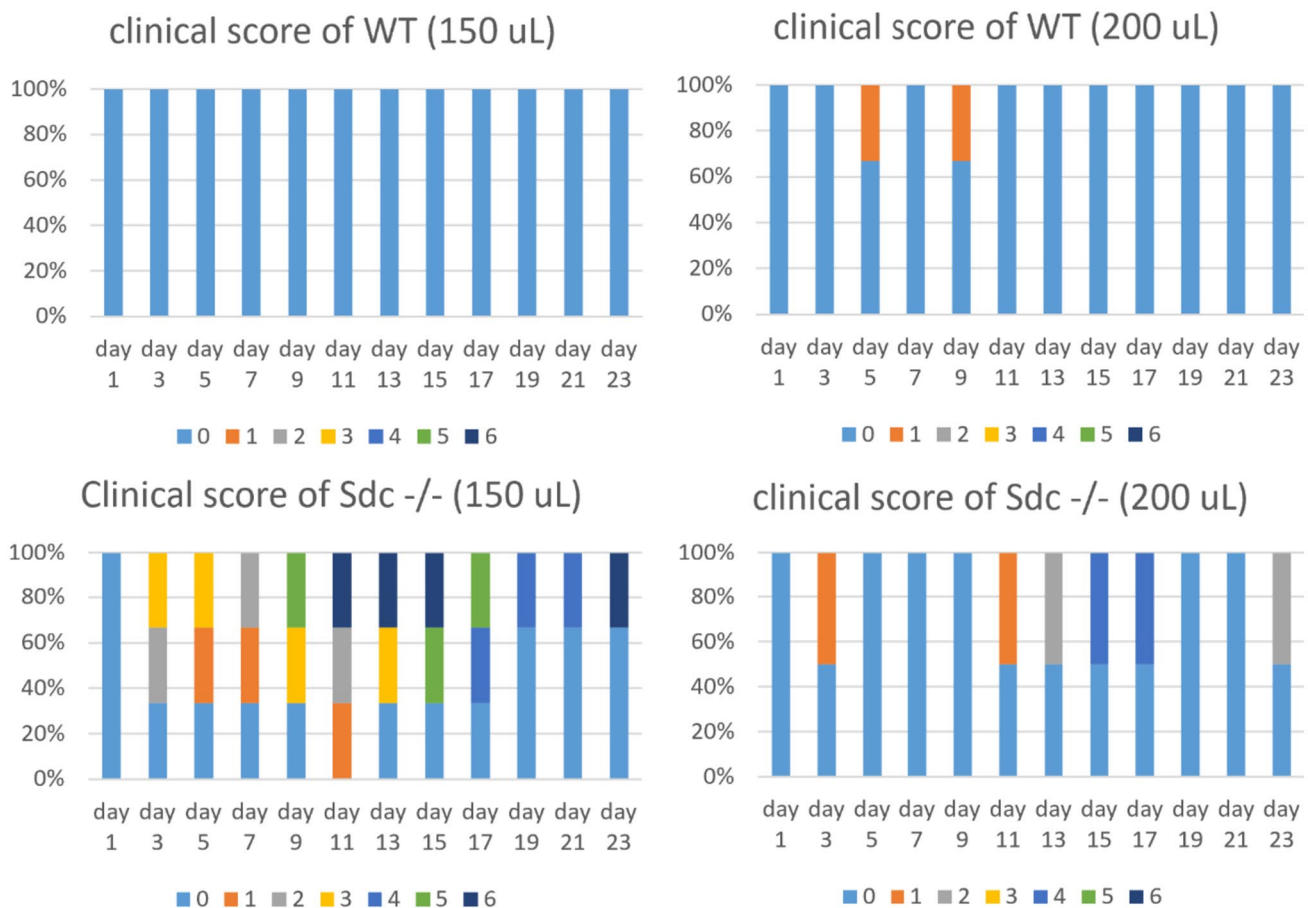


Fig. 1 Arthritis clinical score. Arthritis score of four hind paws per mouse of the groups injected with 75 μl CII+75 μl CFA and 100 μl CII+100 μl CFA was evaluated every other day for 23 days after the second injection. Each paw of each mouse is scored from 0 to

3. Then, the scoring of all paws of each mouse is added to obtain a total score varying from 0 to 12 for each mouse. The highest score obtained was 6 for the Sdc-1 knockout mice

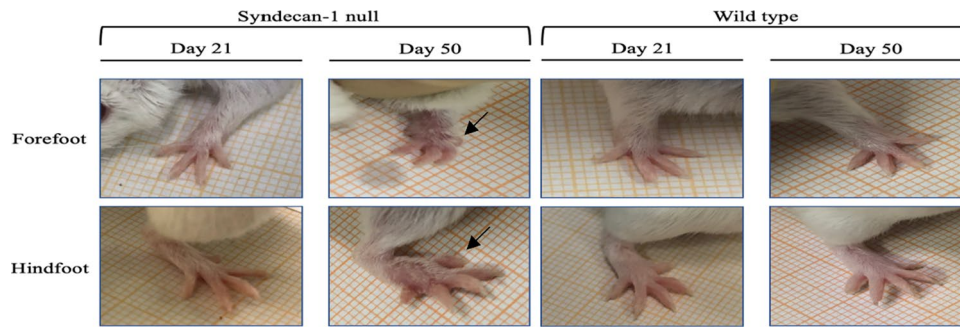


Fig. 2 Photos of forefeet and hindfeet of Sdc-1 knockout and wild-type mice on day 21 (booster injection day) and day 50 after the start of the experiment. Sdc-1 null mice developed redness and swelling on day 50 while wild-type mice did not show gross inflammatory

signs by then. Significant difference can be seen both in the forelimbs and hindlimbs: the fingers are more curved and swollen in the Sdc-1 knockout mice

Fig. 3 H&E staining for CII model. Magnification $\times 200$. Hematoxylin–eosin staining of joints for CII model mice, wild-type, and Sdc-1 knockout. Note the similarities among the control non-injected mice of both strains (a, b). On the other hand, there are significant differences between the 2 strains in the experimental groups. There exist more cartilaginous erosions in the Sdc-1 knockout mice compared to wild-type and such alteration increases with increasing concentration c–h where the joint is fully fibrotic and fused

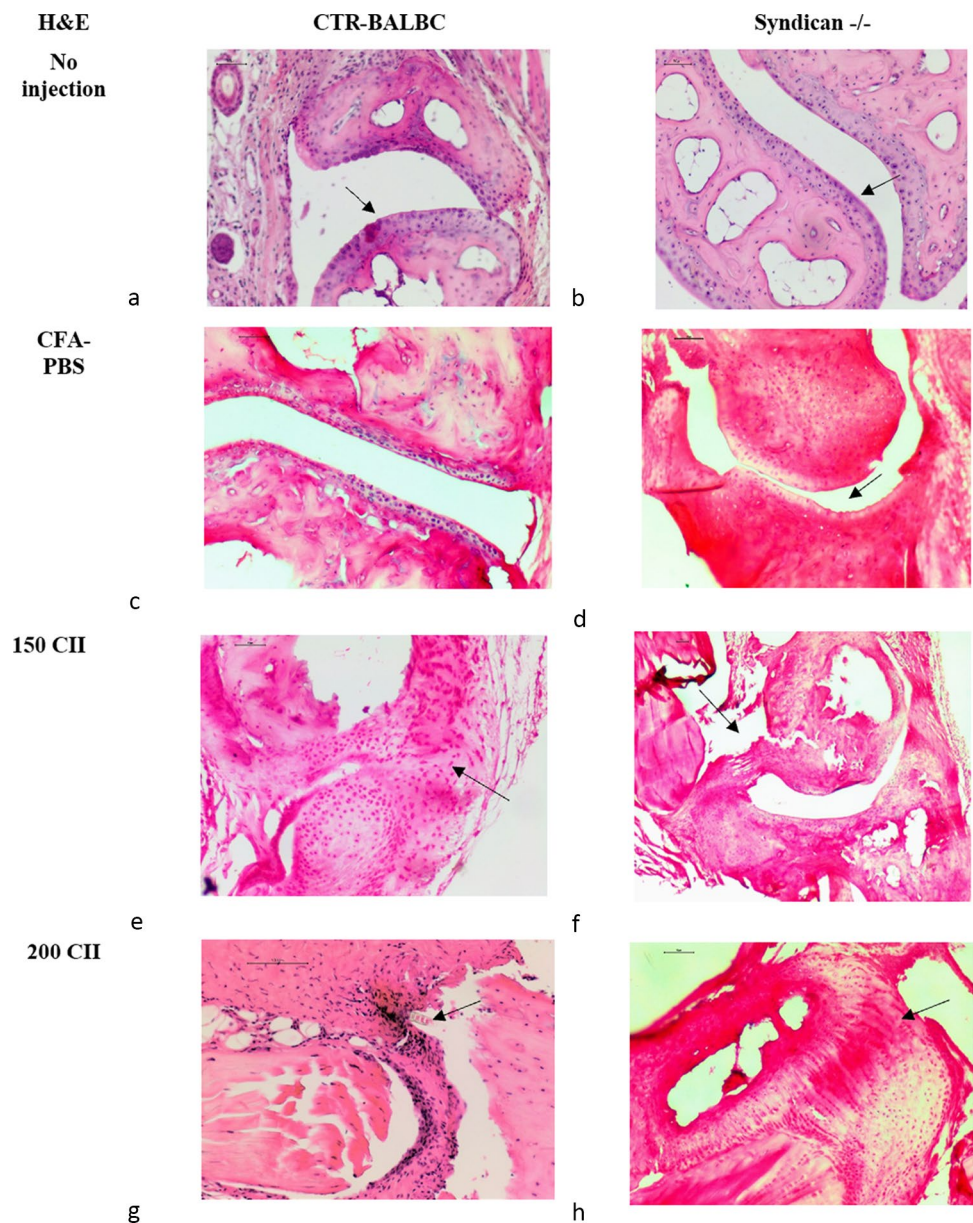
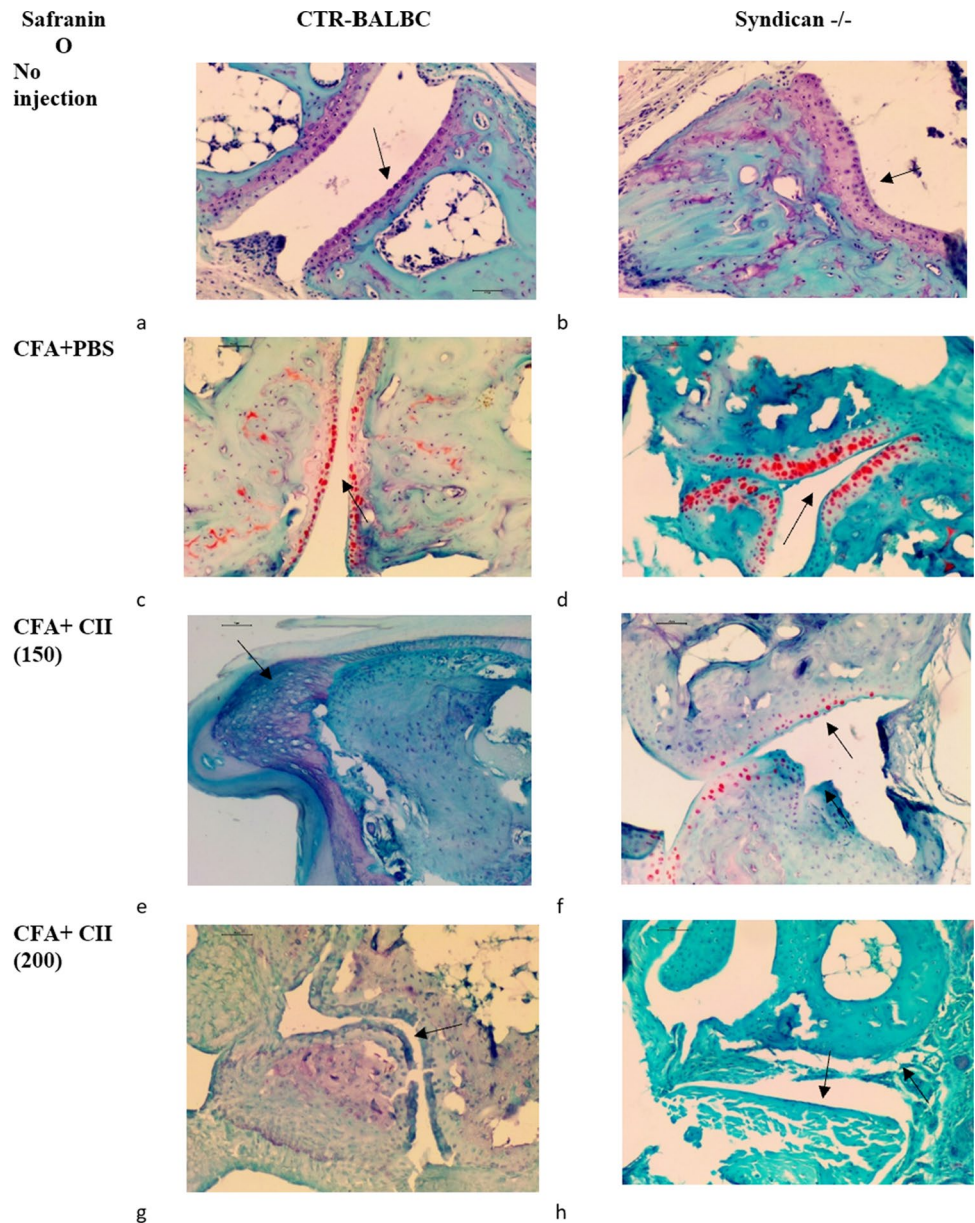


Table 3 Summary table of the bone’s characteristics in the WT and Sdc-1 knockout control and treated groups

	WT/no injection	Sdc-1/no injection	WT/ CFA + PBS	Sdc-1/ CFA + PBS	WT/150	Sdc-1/150	WT/200	Sdc-1/200
Bone tissue	Normal and healthy	Normal and healthy	Normal and healthy	Normal and healthy	Bone deterioration	Bone tissue damage, deterioration and replaced by connective tissue	Bone tissue damage and destruction	Bone fusion, abnormal bone morphology, bone deterioration
Cartilage tissue	Normal and healthy	Normal and healthy	Normal and healthy	Normal and healthy	Cartilage destruction	Cartilage destruction	Cartilage destruction	Total loss of cartilage
Inflammatory cells infiltration	Absence	Absence	Absence	Absence	Highly present	Highly present	Highly present	Highly present
Synovial hyperplasia	Absence	Absence	Absence	Absence	Presence	Highly inflamed joint	Highly inflamed joint	Highly inflamed joint

Fig. 4 Safranin O staining for CII model. Magnification $\times 200$. Safranin O stain of joints for CII model mice, wild-type, and Sdc-1 knockout. Note the significant similarities between the 2 non-injected groups (a–d). On the other hand, a thinner cartilage is seen in the Sdc-1 knockout group compared to wild-type (e, f). There is a complete cartilage degradation in the group with increased concentration of CII (g, h)



Results

Severity of the inflammation in the RA mice

Clinical signs and symptoms

To explore further the involvement of Sdc-1 in the development of rheumatoid arthritis, we established a collagen-induced arthritis mice model in 2 groups: wild-type and Sdc-1 knockout. These mice were monitored 3 times per week for 70 days and clinical scores were recorded. No significant inflammatory signs and swelling were observed in the wild-type group. However, paw thickening, pronounced swelling, and erythema were seen in the Sdc-1 null group at day 50 (Fig. 1). Moreover, a significant increase in clinical scores was found in the knockout group, compared to the wild-type (WT) group where the score was close to 0. We can deduce that redness and swelling of joints appeared stronger in the Sdc-1 null group injected with 75 μ l CII + 75 μ l CFA compared to the wild-type BALB/c group with similar treatment (Fig. 2).

Histological assessment

Histological assessment showed the structural differences between the various groups in bone, cartilage, and mast cells and inflammatory cells in general of the knee-joint and hind paw sections.

Hematoxylin–eosin staining showed more alterations in the Sdc-1 null mice group compared to the WT group, especially in the group injected with 100 μ l CII + 100 μ l CFA: bone erosion and fusion, cartilage destruction, synovial hyperplasia, inflammatory cell infiltration, subchondral bone porosity as well as replacement of bone tissue by connective tissue (Fig. 3; Table 3).

In contrast, none of these RA characteristics was observed in the control group.

Safranin O staining confirmed these results by showing an increase in articular cartilage destruction in the Sdc-1 null group compared to the WT group, especially in the group injected with 100 μ l CII + 100 μ l CFA where a total loss of cartilage is detected as well as bone deterioration, while

Fig. 5 TB staining for CII model. Magnification $\times 200$. Toluidine blue stain of joints for CII model mice, wild-type, and Sdc-1 knockout. Note the higher number of larger mast cells in the wild-type compared to smaller size and relatively lower numbers in the knockout mice (degranulating mast cells in Sdc-1 knockout mice) (a–h)

no cartilage destruction was observed in the control group (Fig. 4; Table 4).

However, more granulated mast cells can be seen in the WT group compared to the Sdc knockout group after CII and CFA injection. This can be explained by the fact that the mast cells reacted and responded to the inflammation in the RA-WT group, while in the RA-Sdc knockout group, they may had an earlier response, degranulated, and released their cytokines, and no more mast cells are produced (Fig. 5; Table 5).

TNF- α expression

TNF- α expression was also studied. We investigated its expression in normal control groups (untreated), CFA-PBS control groups, and treated groups with 150 μ l and 200 μ l CFA + CII.

Figure 6 shows a slight increase in TNF- α expression can be detected in the Sdc-1 knockout models compared to WT models among all groups (treated and untreated). In the Sdc-1 knockout group treated with 200 μ l CFA + CII, we can notice an important increase in TNF- α expression compared to the WT group treated with 200 μ l CFA + CII and compared to other groups. However, no significant differences were noted across all groups. More experimentation with a panel of cytokines is warranted before any conclusion.

Discussion

Emanating data seem to support the fact that Sdc-1 is implicated in RA pathophysiological processes when its expression is altered (Teixeira and Götte 2020). Based on several studies, the Syndecans family, a protein family, was involved in acute and chronic inflammatory diseases (Koliakou et al. 2022). However, their role differs between the types of tissues they are expressed in. The literature shows that Sdc-1 has a controversial role: it can

Table 4 Summary table of the cartilage characteristics in the WT and Sdc-1 knockout control and treated groups

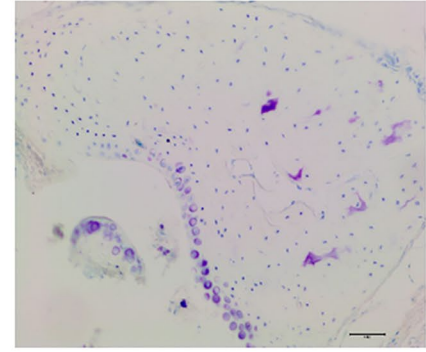
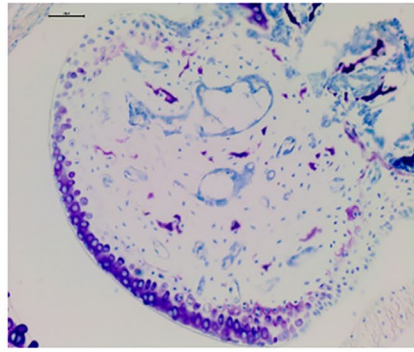
	WT/no injection	Sdc-1/no injection	WT/CFA + PBS	Sdc-1/CFA + PBS	WT/150	Sdc-1/150	WT/200	Sdc-1/200
Cartilage shape	Healthy and normal	Healthy and normal	Healthy and normal	Healthy and normal	Slight deterioration	Pronounced deterioration and destruction	Slight destruction	Total loss of cartilage
Cartilage thickness	Thick	Thick	Thick	Thick	Thick	Thin	Thin	Absent

TB

CTR-BALBC

Syndican^{-/-}

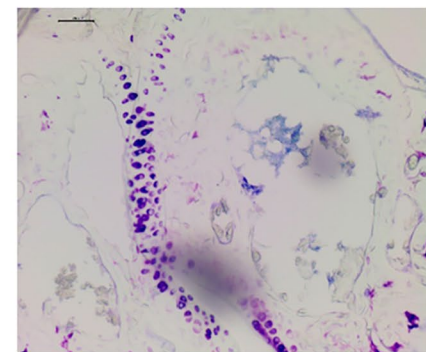
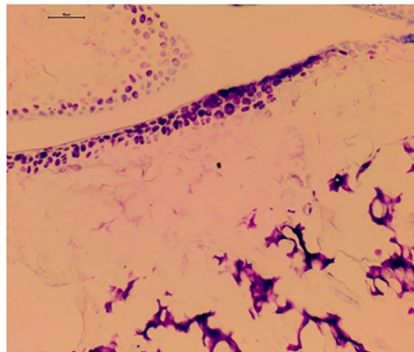
No injection



a

b

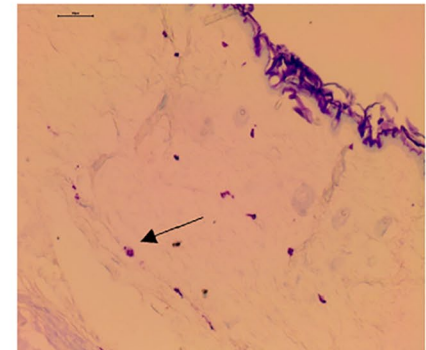
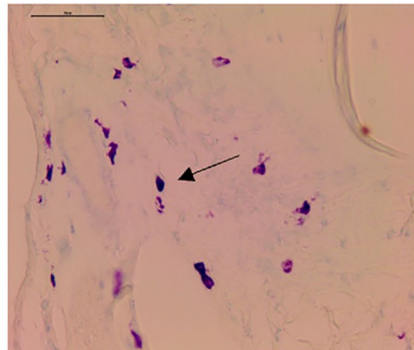
CFA+PBS



c

d

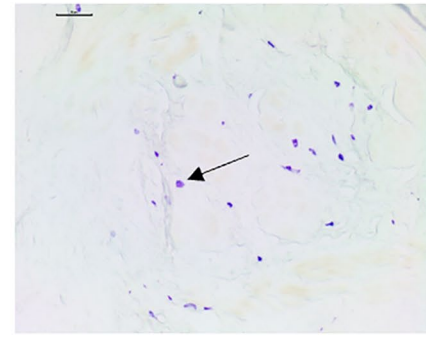
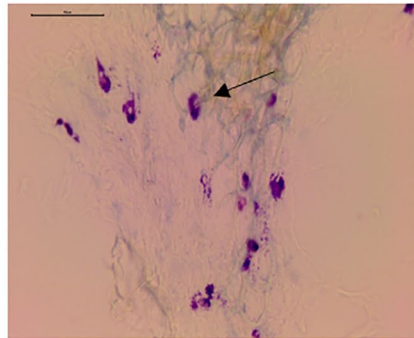
CFA+CII (150)



e

f

CFA+CII (200)



g

h

Table 5 Summary table of the mast cell characteristics in the WT and Sdc-1 knockout control and treated groups

	WT/no injection	Sdc-1/no injection	WT/CFA + PBS	Sdc-1/CFA + PBS	WT/150	Sdc-1/150	WT/200	Sdc-1/200
Absence or presence of mast cells	Absence	Absence	Absence	Absence	High presence	Light presence	High presence	Light presence
Mast cell granulation	-	-	-	-	Highly granulated	Degranulated–phantom mast cells	Highly granulated	Degranulated–phantom mast cells
Inflammatory response	-	-	-	-	Late	Early	Late	Early

exert either an anti-inflammatory or a pro-inflammatory activity depending on the type of the tissue and whether it is shed from the membrane or not (Koliakou et al. 2022). Little is known about its involvement rheumatoid arthritis. Therefore, we performed a Sdc-1 knockout mice model induced with collagen II, to assess and explore the effect of the loss of this protein on inflammation and the inflammatory process in the development of rheumatoid arthritis disease.

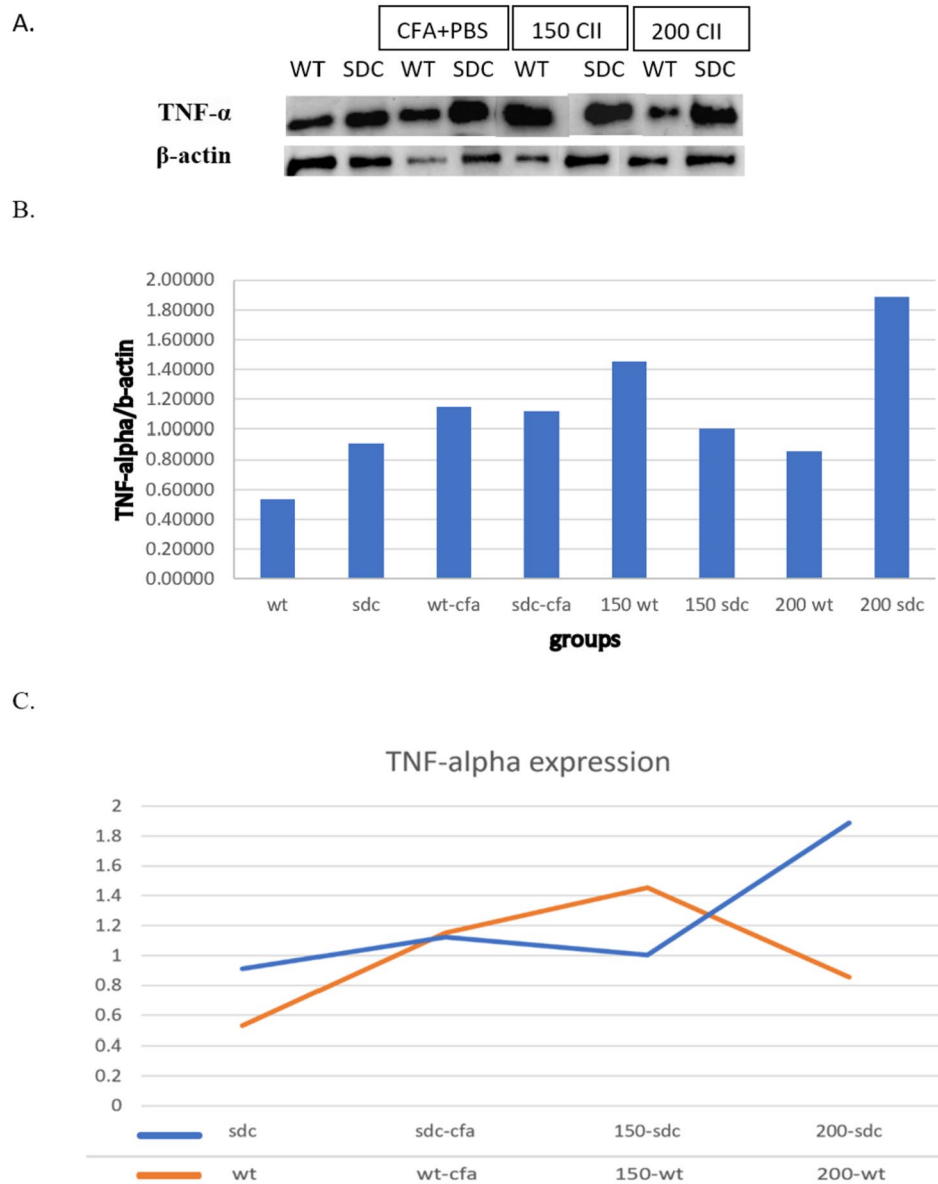
Previous studies found that Sdc-1 expression has increased in chondrocytes isolated from damaged cartilage in early stages of osteoarthritis, suggesting that this molecule is involved in the repair mechanism of the damaged joints (Salminen-Mankonen et al. 2005). However, its role in RA was still undetermined. It is only known that antirheumatic drugs decrease the serum level of Sdc-1, which might reflect a decrease in Sdc-1 shedding (Deyab et al. 2021). In our work, we noticed more histological alterations and damage with higher inflammation in joints lacking the presence of Sdc-1. Moreover, less mast cells were present in the Sdc-1 knockout group, which can be explained by 2 mechanisms: either they exerted an early response, released their components, degranulated and no more mast cells were produced, or the expression of mast cells was downregulated because of the deletion of Sdc-1 protein whereby its loss is known to have an impact on different leucocyte cell types (Gilfillan et al. 2011; Gopal et al. 2021). Mast cells strikingly increased in number in the wild-type, while it was noted that many degranulated in the Sdc-1 knockout, leaving the so-called phantom mast cells. Mast cell granules released into the tissue in response to stimuli in the microenvironment (CFA, CII) via multiple receptors can contribute to the ongoing process of inflammation via multiple mechanisms. In addition, there are numerous important mediators constitutively expressed

and stored, or newly synthesized on stimulation of mast cells including cytokines like TNF- α , chemokines, growth factors, lipids, proteases, heparin, and histamine (Mukai et al. 2018).

In the initial stages of the pathogenic process, mast cells being preferentially present around blood vessels, where injection of CFA and CII took place, could have contributed via dependence on their stimulated microenvironment, early in the process, through the multiple receptors and release of a panoply of mediators. As such, the mast cells we encountered in Sdc-1 knockout mice were mostly degranulated phantom cells.

Western blot was performed to explore the effect of Sdc-1 knockout on TNF- α expression in the CII-induced arthritis mice model. As a result, no significant increase in TNF- α expression was seen among all groups. Only a slight increase was detected in the Sdc-1 knockout mice compared to WT mice, in all groups, suggesting that the absence of Sdc-1 did not significantly affect the expression of the pro-inflammatory cytokine TNF- α . Other pro-inflammatory (IL-6 or IL-1 β) or anti-inflammatory (IL-4, IL-10, or IL-11) molecules could have been affected; they will be explored in a future study. It is also likely that this pathologic process may act through other inflammatory pathways, which were not explored. A previous study explored the regulation of Sdc-1 expression by inflammatory cytokines in IBD. Results showed that stimulation of colonic epithelial cells with TNF- α or IL-1 β downregulated the expression of Sdc-1 at both protein and mRNA levels. However, this effect was not detected when the colonic cells were stimulated with IL-6 (Day et al. 2003), proving that, even in chronic inflammatory diseases, only selective pro-inflammatory cytokines could be affected (Day et al. 2003).

Fig. 6 Tumor necrosis factor alpha (TNF- α) immunoblot analysis derived from bones. TNF- α of molecular weight 17 kDa shows an important increase in expression in the Sdc-1 knockout group treated with 200 μ l CFA+CII compared to WT. However, no significant increase in TNF- α expression can be detected among all groups



Conclusion

This study proved that the absence of Sdc-1 could lead to increase in inflammatory response. Moreover, no previous data are available showing the effect of Sdc-1 knockout on TNF- α expression in rheumatoid arthritis disease. This study is presenting preliminary data suggesting that Sdc-1 may have anti-inflammatory properties in RA arthritis disease; however, more studies at the molecular level should be conducted to confirm such a role and possible mechanistic pathways involved.

The Sdc-1 null mice developed a faster and more severe experimental RA by collagen-CFA injections; however, more characterization at the macroscopic, microscopic, and molecular levels is needed. Uncovering the role of Sdc-1 in RA paves the way for future investigations that can lead to

the emergence of new therapeutic modalities that could be adopted along with conventional treatments.

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Data availability Not applicable.

Declarations

Ethical approval The study was approved by the Institutional Animal Care and Use Committee of the American University of Beirut, Lebanon (IACUC# 19-09-552).

Consent to participate Not applicable

Competing interests The authors declare no competing interests.

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