


# Microparticles in systemic sclerosis, targets or tools to control fibrosis: This is the question!

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## Abstract

Systemic sclerosis is the main systemic fibrotic disease with unknown etiology characterized by peripheral microvascular injury, activation of immune system, and wide-spread progressive fibrosis. Microparticles can be derived from any cell type during normal cellular differentiation, senescence, and apoptosis, and also upon cellular activation. Carrying along a broad range of surface cytoplasmic and nuclear molecules of originating cells, microparticles are closely implicated in inflammation, thrombosis, angiogenesis, and immunopathogenesis. Recently, microparticles have been proposed as biomarkers of endothelial injury, which is the primary event in the genesis of tissue fibrosis. Microparticles may have a role in fostering endothelial to mesenchymal transition, thus giving a significant contribution to the development of myofibroblasts, the most important final effectors responsible for tissue fibrosis and fibroproliferative vasculopathy. Thanks to potent profibrotic mediators, such as transforming growth factor beta, platelet-derived growth factor, high mobility group box I protein, nicotinamide adenine dinucleotide phosphate oxidase 4, and antifibrotic agents, such as matrix metalloproteinases, microparticles may play an opposite role in fibrosis.

## Keywords

Systemic sclerosis, microparticles, biomarkers, endothelial damage, fibrosis

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## Introduction

Fibrotic diseases are one of the leading causes of mortality due to the scarce therapeutic options available today.<sup>1,2</sup> Despite numerous recent advances in the understanding of various complex mechanisms responsible for the pathogenesis and progression of fibrotic diseases, this topic still remains elusive.<sup>3</sup> The pathogenetic pathways leading to fibrosis are multiple, but a classic model of fibrotic diseases is represented by systemic sclerosis (SSc). This disease is characterized by a complex pathophysiology<sup>4</sup> which is not only represented by an immune dysfunction but also by the peculiar involvement of the microvascular system.<sup>5</sup> The sufferance of the endothelial cells (ECs) covering the vessel wall is a pivotal event in the disease pathogenesis also for the endothelial transition into myofibroblasts (EndoMT).<sup>6</sup> Therefore, the endothelium injury may be considered as a primary event which leads to tissue fibrosis. For this reason, the markers of endothelial

injury have been studied at large, but none of them has been identified as useful and reliable marker of the endothelial sufferance so far.

Microparticles (MPs) can regulate vascular thrombosis, angiogenesis, vascular reactivity, and inflammation.<sup>7</sup> Epigenetic studies contributed to better understanding of MPs and its potential role in fibrosis.<sup>8–10</sup> It is well known that MPs have been also proposed as markers of endothe-

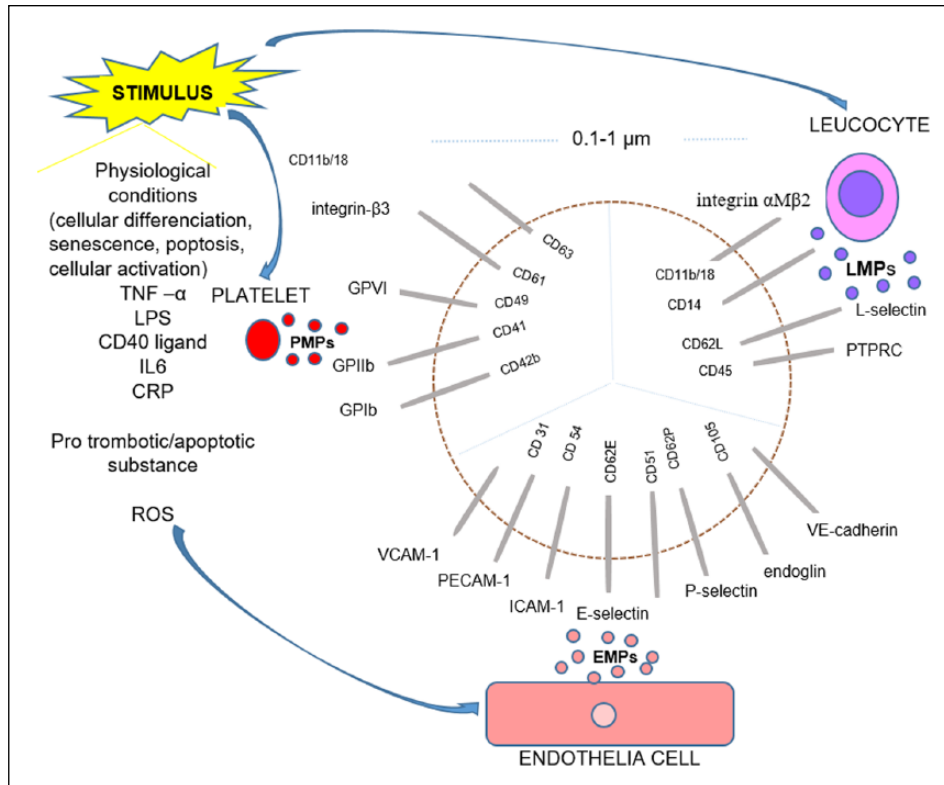
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**Figure 1.** Schematic presentation of MPs surface main components.

EMPs: endothelial cells-derived microparticles; PMPs: platelet-derived microparticles; LMPs: leukocyte-derived microparticles; GPVI: glycoprotein VI; GPIIb: glycoprotein IIb; GPIbα: glycoprotein Ibα; VCAM-1: vascular cell adhesion molecule-1; PECAM-1: platelet endothelial cell adhesion molecule-1; ICAM-1: intercellular cell adhesion molecule-1; E-selectin: endothelial selectin; CD51: vitronectin receptor; P-selectin: platelet selectin; L-selectin: leukocyte selectin.

lial involvement, and their role in the genesis and maintenance of fibrosis is recently hypothesized.<sup>11,12</sup>

The aim of this review is to assess the importance of MPs in the pathogenetic cascade involved in tissue fibrosis.

### MPs: state of the art

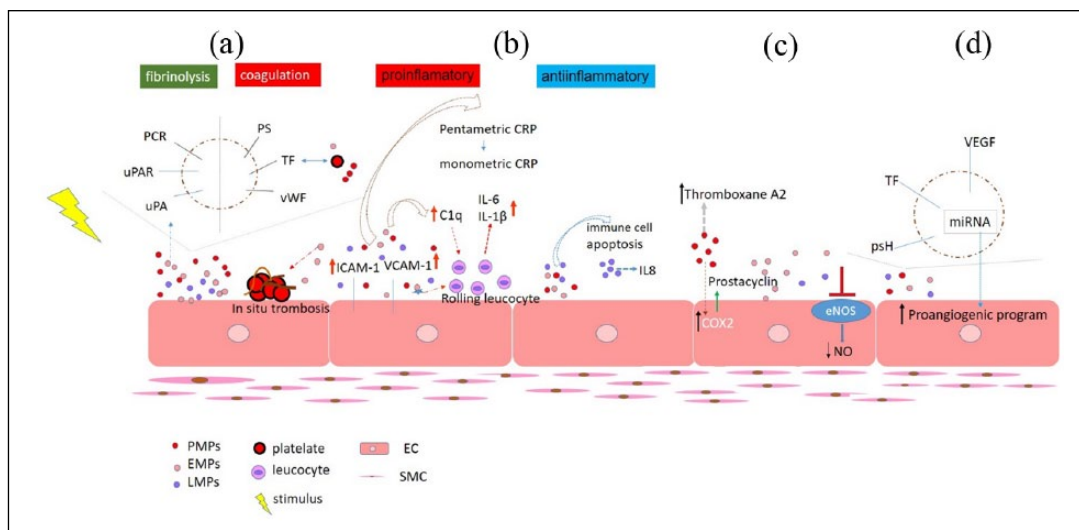
In the literature, MPs, also called microvesicles and ectosomes, are heterogeneous population of membrane-coated vesicles generated from the cells via outward blebbing of the plasma membrane under both physiological and pathological conditions.<sup>13–15</sup> Although MPs may be distinguished from other extracellular vesicles (such as exomes and apoptotic bodies) according to the mechanism of formation and their content, the most commonly used differentiating parameter is size. MPs are typically defined as 0.1–1 μm in diameter, while exomes are smaller (approximately 40–100 nm). Apoptotic bodies are much larger compared to both exomes and MPs, with size of 1–5 μm.<sup>14</sup>

MPs are generated and released during different biological processes, including not only normal cellular differentiation, senescence, or apoptosis but also upon cellular activation following stimulation with proinflammatory cytokines (i.e.

tumor necrosis factor α (TNF-α), lipopolysaccharides (LPS), soluble CD40 ligand, interleukin-6 (IL-6), IL-1α, and C-reactive protein (CRP)), prothrombotic (i.e. thrombin, collagen, proteinase-activated receptor agonists, and plasminogen activator inhibitor-1β (PAI-1)), or proapoptotic substances and exposure to high shear stress (Figure 1).<sup>13,16–19</sup>

These small particles (0.1 and 1 μm in diameter) may be distinguished from other groups of cell-derived vesicles because the MPs membrane and its proteins originate from their parental cells, reflecting both the type and state of their cellular origin.<sup>13,14</sup> Thus, MPs express a broad range of surface cytoplasmic and nuclear molecules that are incorporated into membrane-bound structures including bioactive lipids, integrins, cytokines, and enzymes.<sup>13,14,16,20</sup> MPs contain DNA, RNA, including micro RNA (miRNA), histones, and damage-associated molecular patterns (DAMPs).<sup>21,22</sup> Once MPs are released into circulation, they bind and fuse with their target cells through receptor/ligand interaction and act as biological vectors.<sup>13</sup>

MPs can deliver miRNA into recipient cells, where the exogenous miRNA may regulate target gene expression and modulate the function of recipient cells. Furthermore, immune complexes with autoantigens presented by MPs may induce immune response; in addition, the particles



**Figure 2.** Schematic presentation of possible microparticles activity: (a) Hemostatic properties of MPs. Platelet aggregation and spreading of procoagulant potential; (b) Induction and amplification of inflammation. Immune cell apoptosis and production of anti-inflammatory mediators; (c) Vascular reactivity and endothelial dysfunction; and (d) Angiogenesis.

MPs: microparticles; PS: phosphatidylserine; TF: tissue factor; vWF: large von Willebrand factor; PCR: protein C receptor; uPAR: urokinase-type plasminogen activator receptor; ICAM-1: intercellular adhesion molecule 1; VCAM-1: vascular cell adhesion molecule-1; C1q: complement component 1q; COX 2: cyclooxygenase 2; NO: nitric oxide; eNOS: endothelial nitric oxide synthase; psH: protein sonic Hedgehog; PMPs: platelet-derived MPs; EMPs: endothelial cells-derived MPs; LMPs: leukocyte-derived MPs; EC: endothelial cell; SMC: smooth muscle cell.

themselves may coordinate functions of different cells via both autocrine and paracrine ways. Thus, MPs are found as central mediators of a communication network for the local and systemic intercellular exchange of biological information and cell–cell interaction.<sup>17,23</sup> Some components of MPs are selectively enriched compared to their cell of the origin, and even more, the composition and the function of MPs not only depend on the cellular origin but also on the inducing triggers and the microenvironment of the parental cell.<sup>23,24</sup>

Although circulating MPs can be derived from virtually any cell type, including immune cells, they are most commonly originated from the vasculature and circulating blood cells: endothelial cells (endothelial cells–derived microparticles (EMPs)), platelets (platelet-derived microparticles (PMPs)), leukocytes (leukocyte-derived microparticles (LMPs)), and vascular smooth muscle cells.<sup>13</sup> The components on the MPs surface are the most notable, since they allow detection by flow cytometry (FC). Thus, EMPs may display platelet endothelial cell adhesion molecule-1 (PECAM-1; CD31), vascular endothelial cadherin (VE-cadherin; CD144), vitronectin receptor (CD51), ICAM-1 (CD54), vascular cell adhesion molecule-1 (VCAM-1; CD106), endothelial selectin (E-selectin; CD62E), platelet selectin (P-selectin; CD62P), or endoglin (CD105).<sup>25</sup> PMPs can exhibit glycoprotein Iba polypeptide (GPIba; also known as CD42b), glycoprotein IIb (GPIIb; CD41), glycoprotein VI (GPVI; CD49), integrin- $\beta 3$  (CD61), and the lysosomal markers such as CD68 or CD63, while LMPs may express protein tyrosine phosphatase receptor type C (PTPRC; CD45), CD14, L-selectine (CD62L), and integrin  $\alpha M\beta 2$  (CD11b/18)<sup>25–27</sup>.

The expression of different antigens on the MPs surface depends on the state of the origin cell. Thus, the activated ECs increase expression of inducible antigens on EMPs (e.g. CD62E, CD106, and CD54), while ECs undergoing apoptosis enhance expression of constitutive antigens (e.g. CD31 and CD105) and increase binding of annexinV.<sup>13,28</sup>

Presence of different components/antigens suggests a wide range of MPs activities, mainly in hemostasis, inflammation, vascular reactivity, and angiogenesis.<sup>4,9,29–33</sup>

Increased number of EMPs and PMPs are found in patients with antiphospholipid syndrome, suggesting a role of MPs in thrombotic events and pregnancy complications.<sup>7</sup> PMPs could be considered as a biomarker for the risk of thrombosis or miscarriage in individuals with antiphospholipid antibodies.<sup>34</sup> Enhanced TF expression on both EMPs and LMPs has been shown in systemic lupus erythematosus (SLE) patients, indicating an active role of TF-positive MPs in thrombosis.<sup>35</sup>

Furthermore, EMPs vesiculation correlate with IL-6 release, showing that close relationship between endothelial vesiculation and classical inflammatory pathway exist and that MPs are implicated in inflammation.<sup>9</sup>

Generally, MPs can affect many different processes of the vasculature. PMPs may induce the expression of cyclooxygenase 2, leading to the release of the vasodilative mediator prostacyclin. On the contrary, PMPs may contain the arachidonic acid metabolite thromboxane A2, which increases vascular contraction. Furthermore, EMPs have an endothelial-dependent vasodilation effect influencing directly vascular tone (Figure 2).<sup>36,37</sup>

MPs may promote endothelial dysfunction by impairing the endothelial nitric oxide (NO) pathway and inducing proinflammatory response (Figure 2).<sup>38</sup> EMPs are considered as a new useful and reliable marker of endothelial dysfunction. Furthermore, it has been shown that PMPs can stimulate angiogenesis and revascularization in ischemic heart disease.<sup>10</sup>

Despite increasing scientific and clinical interest, methodology for MPs assessment is still an area of great debate, which is impeded by technological issues.<sup>39,40</sup> Nevertheless, many different methods of MPs detection in biological samples have been described in the literature so far.<sup>40,41</sup> No standardized protocols are available for the isolation, detection, and characterization of MPs.

All preanalytical steps, from blood sampling to sample freezing, should be considered as a source of variation in MPs analysis. For instance, isolation of MPs from blood is affected by the following: venepuncture and the diameter of the needle; time between blood collection and first centrifugation, which should be within 30 min to 1 h from sampling, no more than 2 h; type of anticoagulant used; and freezing and storage of the samples (storage no more than 1 year after freezing until analysis at a temperature below 80°C), thawing, centrifugation, and washing procedures.<sup>40,42–44</sup> After blood collection, platelets need to be removed from the plasma in order to avoid cellular activation, leading to involuntary production of MPs. Centrifugation protocols for preparing platelet-free plasma and isolated MPs have major influences on MP analysis. The common centrifugation parameters used for MPs isolation vary between 1500g and 10,000g for 5–20 min in the first centrifugation step intended to remove cells and large particles (including platelets), followed by 13,000g–100,000g for 30–60 min to exclude residual platelets obtaining MPs pellet. These differences in centrifugation speed and time greatly affect the final MP counts. It has been shown that initial low-speed centrifugation between 1200g and 2000g for 15–20 min could effectively remove erythrocytes, platelets, and large membranous fragments, whereas speeds >2000g lead to a substantial loss of MPs.<sup>40,45</sup> Sometimes, in order to remove residual platelets, filtration with 0.8- $\mu$ m porous membranes is used;<sup>46</sup> however, filtration may activate platelets and induce MPs fragmentation, leading to serious loss of MPs.<sup>45</sup>

Isolation of MPs in SSc studies have included either one step (vary from 1500g to 2000g, 10–15 min)<sup>47–49</sup> or two step (first: 200g–1800g, 10 min and second: 800g–20,000g, 6–10 min) centrifugation.<sup>50–53</sup> After centrifugation and freeze-thaw steps had been done, one more centrifugation with two steps (1500g for 5 min and 100,000g for 20 min) was performed in one study.<sup>49</sup>

Arising concern is that isolation could result in getting MPs contaminated with exosomes; apoptotic blebs; protein aggregates, including immune complexes; the presence of lipoprotein particles; and small platelets within the size

range of MPs.<sup>46,54,55</sup> In spite of these limitations, the combination of differential centrifugation and FC has proven to be invaluable for the detection of MPs.<sup>49–51,53,56–59</sup>

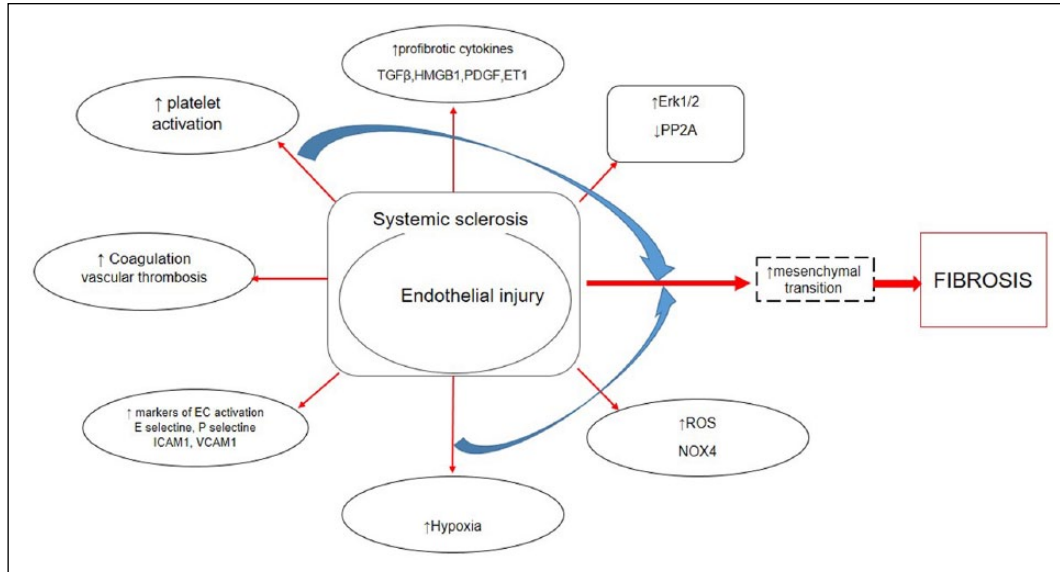
Different optical and non-optical detection methods have been utilized for the assessment/quantification of MPs, including immunoassays, FC, electron microscopy, atomic force microscopy, and dynamic light scattering.<sup>40,41</sup>

FC analysis of blood MPs appears to be the most favored analytical method of identification, quantification, and size assessment of the microvesicles.<sup>40,60</sup> MPs are typically detected in terms of size by FC based on the intensity of light scattering and exposure of phosphatidylserine (PS), an “eat me signal” for the immune system, identified by staining with annexin V and further characterized with fluorescent-labeled antibodies against specific surface antigens.<sup>13,39–41,60,61</sup> One of the biggest FC limitation is MPs size detection, regarding the fact that MPs are too small and heterogeneous in size to be detected and clusters of small MPs might be counted as one event.<sup>39,40</sup> Furthermore, immune complexes can overlap with size of MPs appearing as MPs by FC.<sup>46</sup> Quantification of MPs could be done with detection of PS-rich surface; annexin V (AnxV) binding is often used as identifier.<sup>39,40,46,60–62</sup> The mechanisms of generating AnxV non-binding MPs (AnxV-MPs) are not fully known, but because PS exposure is a typical feature of apoptosis, it may be hypothesized that AnxV-MPs are generated by cellular activation. However, a significant proportion of the MPs are annexin V negative, suggesting either heterogeneity in the mechanism of production or the presence of PS at concentrations below the limits of detection. As an alternative, lactadherin could be used with higher affinity for PS and the potential advantage of non-calcium dependent binding to PS compared to AnxV. BODIPY maleimide, calcein AM, and SYTO 13 are other alternative probes that could detect all circulating MPs also.<sup>13,39,63,64</sup> Identification the MPs origin with fluorescent-labeled antibodies also has limitation in that antibodies are dissimilar according to specificity and sensitivity for specific antigens.<sup>39</sup> Finally, the standardization of both preanalytic and analytic methods still remains a significant challenge.<sup>40</sup>

The link between inflammation, coagulation, and fibrosis is well documented. Even low-grade persistent inflammation is enough to promote fibrosis. Alike, plasma coagulation cascade proteases are also involved in fibrosis via induction of profibrotic molecules.<sup>65</sup> However, our knowledge of MPs in the development of fibrosis is unclear.

## Fibrosis: state of the art

Fibrosis is characterized by excessive deposition of extracellular matrix (ECM) proteins, mainly collagen and fibronectin, in response to injury and is quite important in wound healing. However, when fibrotic process is chronically active, it could lead to permanent tissue remodeling and significant organ impairment.<sup>5</sup> It has been already known that



**Figure 3.** Illustration of the link between endothelial injury and fibrosis.

fibrosis is a feature of different connective tissue diseases (CTDs) and is the hallmark of the main systemic fibrotic disease—SSc. SSc is a complex, multifaceted CTD of unknown etiology, characterized by a peripheral microvascular injury spreading into progressive fibrosis of the skin and multiple internal organs.<sup>5</sup> We have already pointed out that endothelial injury is a key pathological event in SSc, contributing to enhanced leukocyte, platelet activation, and coagulation pathways; production of proinflammatory and profibrotic cytokines; generation of reactive oxygen species (ROS); defective angiogenesis; and vasculogenesis, resulting in development of severe vasculopathy and further fibrosis (Figure 3).<sup>5,66,67</sup>

SSc-associated interstitial lung disease (ILD) and pulmonary arterial hypertension (PAH) are the leading causes of impaired quality of life and mortality in SSc.<sup>1–3,68</sup> It has been reported that endothelium and pericyte activation, telocytes loss, aberrant immune responses, endoplasmic reticulum stress, and chronic tissue injury are involved in the initiation of fibrosis in SSc.<sup>69–71</sup> Although the pathogenesis of SSc fibrosis is still elusive, strong evidence suggests that myofibroblasts (MFs) are the main final effectors responsible for tissue fibrosis and fibroproliferative vasculopathy (Figure 3).<sup>72</sup> These cells contribute to the progressive increase in tissue stiffness, further enhancing the profibrotic process.<sup>73</sup>

Extensive investigations have revealed that MFs originate from multiple cellular sources, including resident tissue fibroblasts, bone marrow–derived circulating fibroblast precursors (also known as fibrocytes), and epithelial cells via a phenotypic transition into mesenchymal cells (EMT), by which epithelial cells modify adhesive properties and polarity and acquire ECM-producing MFs features. Other cell types such as pericytes, adipocytes, pleural mesothelial cells, or macrophages are also potential sources of MFs.<sup>72,74</sup>

More recent studies have shown that another source of activated MFs are ECs that have acquired a mesenchymal phenotype through an endothelial to mesenchymal transition (EndoMT) process. Today, we know that EndoMT plays an important role during several pathological conditions, including cardiac, pulmonary, and renal fibrosis; carcinoma-associated interstitial fibrosis; idiopathic portal hypertension; intestinal fibrosis; and diabetic nephropathy.<sup>72,74–77</sup> Moreover, EndoMT may play a role in the development of tissue fibrosis and fibroproliferative vasculopathy in SSc.<sup>72</sup> One more evidence which supports this hypothesis is that EndoMT may occur in SSc dermal endothelium, contributing to the development of dermal fibrosis.<sup>6</sup> During EndoMT, ECs become detached from the endothelial layer; change their morphologic characteristics; lose their specific EC markers such as CD31/PECAM-1, large von Willebrand factor (vWF), occluding, and VE-cadherin; and initiate the expression of mesenchymal/myofibroblast phenotype characterized by the expression of alpha smooth muscle actin ( $\alpha$ -SMA), vimentin, S100A4/fibroblast-specific protein-1, and type I collagen (CI). It has been shown that EndoMT may support loss of microvascular EC and thus contribute to capillary rarefaction, leading to chronic tissue ischemia and amplifying further fibrotic process.<sup>72,78,79</sup>

Multiple pathways are implicated in the fibrotic process. The potent profibrotic transforming growth factor beta (TGF- $\beta$ ) has been highlighted as a key player in fibrosis as well as in EndoMT and EMT, confirmed both in vitro and in vivo studies.<sup>78,80,81</sup> TGF- $\beta$ -regulated genes are expressed in the skin and the lung of patients with SSc, and the extent of the cytokine expression correlates with the disease activity.<sup>82</sup> Thrombospondin-1 (TSP-1) is important in controlling TGF- $\beta$  activation in fibrotic diseases.<sup>83</sup> Smad-dependent and Smad-independent pathways and numerous transcriptional regulators such as Snail, Snail2 (or Slug), Twist, and

some members of Zeb family of proteins are implicated in EndoMT.<sup>84,85</sup> According to Smad-independent pathway, extracellular signal-regulated protein kinase (Erk) 1/2 has been suggested to have an important role in fibrosis by regulating MF transdifferentiation, cell proliferation, and survival, as well as matrix synthesis. Erk1/2 pathway is induced by TGF- $\beta$  in dermal fibroblasts and ECs.<sup>86</sup> The protein phosphatase 2A (PP2A) dephosphorylates and blocks activation of ERK1/2.<sup>87</sup> The PP2A mRNA and protein expression are significantly reduced in SSc fibroblasts and correlate with an increase in ERK1/2 phosphorylation and collagen expression.<sup>88</sup> Furthermore, aberrant activation of Notch, Sonic Hedgehog, and Wnt/13-catenin pathways may lead to various pathological consequences, including the development of fibrotic diseases via, in part, EndoMT (Figure 3).<sup>89–91</sup>

EndoMT may be mediated by other potential factors including platelet-derived growth factor (PDGF), vascular endothelial growth factor (VEGF), insulin-derived growth factor, connective tissue growth factor (CTGF), and endothelin-1 (ET-1).<sup>92</sup> It has been proven that ET-1 is capable of generating EndoMT either by itself or in combination with TGF- $\beta$ , and despite this, the synergistic interaction with TGF- $\beta$  is still widely supported. CTGF is a common target for both ET-1 and TGF- $\beta$ . In different cells type, CTGF regulates cell proliferation, apoptosis, migration, mesenchymal cell activation, and ECM accumulation.<sup>93</sup>

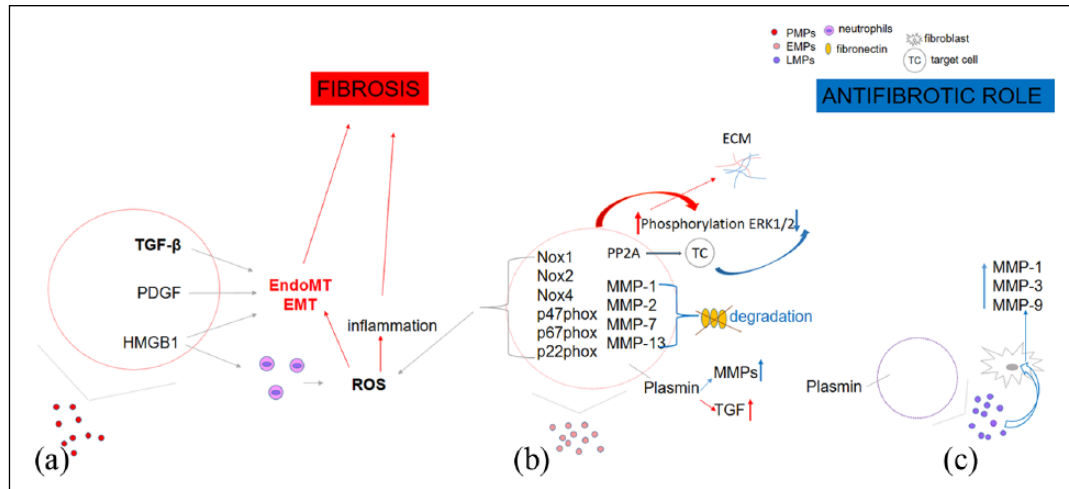
Hypoxia represents a potent stimulus for the generation of various growth factors and ROS, which influence the fate of ECs, promoting mesenchymal transition and fibrosis.<sup>94</sup> Skin hypoxia has been documented in SSc patients.<sup>95</sup> Oxidative stress is implicated in various features of idiopathic pulmonary fibrosis (IPF), liver fibrosis, and SSc, mediated by ROS. Mainly, ROS production derives from the activation of the nicotinamide adenine dinucleotide phosphate oxidase (NOX) family. NOX4 has a central role in the initiation and maintenance of fibrosis. Increased expression of NOX4 transcripts and also level of NOX4 in affected SSc skin have been demonstrated.<sup>96</sup> Moreover, TGF- $\beta$ 1-mediated expression of NOX4 is closely related to myofibroblastic differentiation; on the contrary, MF differentiation is dependent on the generation of ROS by NOX4, suggesting that ROS and TGF- $\beta$ 1 are essential for manifestation of the MF phenotype (Figure 3).<sup>97,98</sup> Activation of HIF-1 signaling in renal epithelial cells may promote fibrogenesis by increasing expression of ECM modifying and by facilitating EMT.<sup>99,100</sup> Furthermore, activation of HIF-1 $\alpha$  in dermal fibroblasts of SSc may upregulate CTGF expression, contributing to the progression of skin fibrosis.<sup>101</sup>

Several studies have shown a significant deregulation of miRNAs involved in angiogenesis, vascular repair, and endothelial homeostasis.<sup>102</sup> In SSc, several miRNAs are associated with kTGF- $\beta$  and CI expression. Thus, microRNA-29a and miRNA-196a can suppress CI gene expression, but they are downregulated in SSc, suggesting that

their low-level expression may promote upregulation of CI by TGF- $\beta$  in SSc fibrogenesis. Moreover, levels of miRNA-196 inversely correlate with prevalence of pitting scars and modified Rodnan skin thickness score (mRSS) score.<sup>103,104</sup> The downregulation of miRNA let7a leads to the excessive CI expression and, recently, has been shown that treatment with let7a improves the skin fibrosis in SSc.<sup>105</sup> Several studies have proven that using strategy with combination of miRNAs can transdifferentiate fibroblasts into cardiomyocytes or neuronal tissue. However, the rate of reprogramming fibroblasts is low and insufficient to translate into a clinical setting.<sup>106,107</sup> Recently, it has been demonstrated that MPs derived from endothelial progenitor cells could protect the kidney from ischemic acute injury by miRNA-dependent reprogramming of hypoxic resident renal cells to a regenerative program.<sup>108</sup>

Circulating blood cells can mediate various features in the fibrotic diseases, through pleiotropic functions. Platelets are critical players in SSc pathogenesis. They represent source of different profibrotic signals such as VEGF, TGF- $\beta$ , PDGF, and serotonin.<sup>109</sup> Platelets also contain high mobility group box 1 (HMGB1) protein (Figure 3). This protein plays multiple roles in the pathogenesis of inflammatory and autoimmune diseases and mediates processes that range from inflammation to repair. It targets various immunologically relevant systems, including p53, nuclear factor (NF)- $\kappa$ B, the glucocorticoid receptor, and the receptor for advanced glycation end products (RAGE). The signaling of HMGB1 or with its receptors plays a crucial role in mediating fibrotic diseases in liver, renal, lung, and myocardial.<sup>11,109</sup> Further studies have revealed that serum HMGB1 level in SSc is higher compared with healthy controls and control mice, while SSc patients with elevated HMGB1 level have more frequent involvement of several organs and immunological abnormalities than those with normal level.<sup>110</sup> The bioactive HMGB1 can stimulate neutrophils to generate ROS via P-selectin, thus contributing to increased vessel inflammation in SSc and oxidative stress development. The oxidation of HMGB1 further amplified its ability to activate neutrophils. It has been demonstrated that HMGB1 + MPs purified from SSc patients activate *in vitro* healthy neutrophils maintaining of sterile inflammation in SSc patients.<sup>111</sup> Moreover, the activation of HMGB1 is associated with the loss of telocytes.<sup>70,71</sup> Finally, HMGB1 may contribute to EMT and EndoMT in various fibrotic diseases (Figure 3).<sup>11</sup>

Data from diverse fibrosis models indicate that matrix metalloproteinases (MMP) may modulate a range of biological processes, especially those related to immunity and tissue repair and/or remodeling, having both inhibitory and stimulatory roles in fibrosis.<sup>112</sup> Since MMP3, MMP2, and MMP9 might stimulate EMT, MMP9 can activate TGF- $\beta$ , contributing to enhance the pool of active TGF- $\beta$ . MMP7-deficient mice are protected from bleomycin-induced lung fibrosis, suggesting profibrotic role



**Figure 4.** Illustration of the link between microparticles and fibrosis: (a) Platelet-derived microparticles (PMPs) secretome may contribute to fibrosis via EndoMT, EMT process, and ROS-mediated pathway; (b) Fibrotic and antifibrotic role of endothelial cells-derived microparticles (EMPs); and (c) Profibrotic and antifibrotic properties of leukocyte-derived microparticles (LMPs). ECM: extracellular matrix; EMT: epithelial to mesenchymal transition; EndoMT: endothelial to mesenchymal transition; ERK1/2: extracellular signal-regulated protein kinase 1/2; HMGB1: High mobility group box 1; NOX: NADP(H) oxidase; p47phox, p67phox, p22phox: NADP(H) oxidase subunits; PP2A: protein phosphatase 2A; ROS: reactive oxygen species.

of these MMPs. Even though, in lung fibrosis model, at the beginning MMP7s facilitate neutrophil influx and activation, leading to epithelial damage and an enhanced fibrotic environment, later epithelial-derived MMP-7s promote resolution by attracting an influx of immunosuppressive leukocytes reflecting also antifibrotic role. In a model of liver fibrosis, MMP-1 and MMP-13 lead to resolution of fibrosis.<sup>112,113</sup> In a SSc patient, MMP-7 serum level is higher than those in control group, and patients with lung fibrosis have higher levels than those without. Interestingly, fibroblasts in early stage of SSc exhibit higher levels of MMP-1, MMP-3, and TIMP-1, unlike the gene expression of MMP-1, MMP-2, and MMP-3, which is decreased in fibroblast from SSc patients with mild stage of disease. MMP-9 concentration positively correlates with the mRSS, and one of the sources of MMP-9 is dermal fibroblast.<sup>114,115</sup>

### Is there a link between MPs and fibrosis?

A few important findings suggest direct implication of MPs in fibrosis and possibly a role in EMT and EndoMT. PMPs secretome contains a range of vasoactive mediators favoring vasoconstriction (e.g. thromboxane), growth factors (TGF-β and PDGF), and HMGB1 protein that may contribute to fibrosis.<sup>9,11</sup> Very new precious data have shown that PMPs from SSc patients interact with neutrophils in vitro and in mice, promoting neutrophil autophagy and leading to generation of neutrophil extracellular traps (NETs).<sup>116</sup> Taken together, all of these data may support the role of PMPs in EndoMT due to fact that NETosis itself is contributing to EndoMT.<sup>117</sup> Furthermore, EMPs may contain Nox1, Nox2, Nox4,

p47phox, p67phox, and p22phox and have the ability to produce ROS through Nox-dependent processes. Moreover, EMPs can increase phosphorylation of ERK1/2 and Src via ROS-independent way, contributing to increased ECM accumulation. A recent proteomic analysis found that PP2A, which is present in human EMPs, may be transferred to target cells, giving insight to decreased ERK1/2 phosphorylation, but this has not been shown experimentally yet.<sup>8</sup> In contrast, MPs from LPS-treated THP-1 monocytes cell may induce phosphorylation of ERK1/2, activation of the nuclear factor-κB pathway, and expression of cell adhesion molecules ICAM-1, VCAM-1, and E-selectin, promoting proinflammatory and profibrotic role (Figure 4).<sup>118</sup>

Podocyte-derived MPs might increase p38 and Smad3 phosphorylation and expression of the ECM proteins in proximal tubule epithelial cells, suggesting their role in EMT and tubular fibrosis.<sup>119</sup> EMT could be fostered by EMPs coming from activated ECs via increasing the expression of HIF-α/VEGF-A in a COX-2/EP2 receptor dependent manner.<sup>120</sup>

As MPs contain proteolytic enzymes, it is possible that MPs may contribute to alteration of the ECM and cleavage of signaling molecules. For example, MPs derived from microvascular ECs contain MMP-1, MMP-2, MMP-7, and MMP-13 and may degrade fibronectin in vitro.<sup>121</sup> Furthermore, EMPs have ability to bind and activate both endogenous and exogenous proMMP-2, leading to vascular matrix remodeling.<sup>122</sup> Very new data have proven that LMPs derived from T cells and monocytes potentially induce the synthesis of MMP-1, MMP-3, MMP-9, and MMP-13 in fibroblasts in a time-dependent manner. On the contrary, no contraregulatory induction of the

expression of tissue inhibitors of MMPs was observed.<sup>14</sup> In SSc, the inverse correlation between the mRSS and values of total EMPs and PMPs has been demonstrated (Figure 4).<sup>49,50</sup>

EMPs and LMPs may generate on the surface plasmin, which has both profibrotic and antifibrotic properties by activating TGF beta on one hand and on the other hepatocyte growth factors and MMPs (Figure 4).<sup>123</sup>

MMPs and plasmin influence migration capacity of cells which could be implicated in fibrosis. MPs promote fibroblast activation and migration. ROS, namely H<sub>2</sub>O<sub>2</sub>, may increase the generation of procoagulant MPs, TF-bearing MPs, by alveolar epithelial cells that could activate local synthesis of factor Xa leading to PAR-1-mediated activation of fibroblasts and a profibrotic response.<sup>124</sup> Furthermore, EMPs isolated from idiopathic pulmonary patients might induce migration capacity of the lung fibroblast, increasing formation of F-actin fibers, by their fibrinolytic activity contributing to fibrogenesis.<sup>125</sup> EMPs are elevated among IPF patients with severe reduced diffusing capacity of the lung for carbon monoxide (DLCO), while TF-bearing MPs negatively correlate with both forced vital capacity (FVC) and DLCO,<sup>125</sup> suggesting their implication in lung fibrosis disease. Evidence that MPs are involved in pathogenesis of lung fibrosis also comes from the finding that LMPs might lead to extensive entry of neutrophils into airways and aggregation at the epithelial surface of the respiratory tract in cystic fibrosis patients.<sup>126</sup> Regarding the well-known profibrotic effect of IL17,<sup>127,128</sup> PMPs could be considered as antifibrotic mediators through their ability to prevent the differentiation of Tregs into IL-17 and IFN- $\gamma$ —producing cells in P-selectin-dependent manner.<sup>129</sup> Moreover, PMPs may deliver miRNA let7a to the ECs and reduce production of TSP-1, possibly influencing TGF- $\beta$  pathway.<sup>130</sup>

## The role of MPs in SSc

Recently, a few research groups reported plasma levels of MPs and their clinical association in SSc patients giving divergent results.<sup>47–53,59,111,116,131</sup> EMPs, PMPs, and LMPs are predominantly investigated, and labeling antigens for detection-specific MPs population has been dissimilar across studies. Furthermore, some studies have shown that patients with SSc have increased concentration of MPs compared to healthy controls,<sup>48–52,111,116,131</sup> while others have demonstrated opposite results (Table 1).<sup>47,59</sup>

Even though some of them have investigated same population of MPs, heterogeneity of the studies with respect to eligibility criteria, study population, methods, and choice of outcome statistics make the comparison difficult.

In spite of the study differences, the association of MPs with hemostasis disturbance, microangiopathy, disease activity, inflammation, and organ involvement in SSc might be speculated.

## Hemostasis

As a common feature, particles expose PS, as a consequence of membrane flipping during apoptosis, which can bind and activate different coagulation factors promoting conversion of prothrombin into thrombin. Furthermore, MPs may express tissue factor (TF) and vWF multimers, which may initiate the extrinsic coagulation pathway and promote platelet aggregation.<sup>29,30,132</sup> EMPs from activated ECs may trigger TF-dependent thrombin formation in vitro and thrombus formation in vivo.<sup>133</sup> In addition, TF can be transferred between MPs and different cell types, spreading procoagulant potential.<sup>134</sup>

Recent studies have demonstrated that EMPs expose endothelial protein C receptor, urokinase-type plasminogen activator, and its receptor, suggesting that these particles also have anticoagulant properties (Figure 2).<sup>12</sup>

Concomitant to the changes in the SSc endothelial lining, platelets undergo activation. Enhanced activation of platelets, increased tendency to aggregation, and activation of coagulative cascade have long been observed in SSc patients.<sup>109,135,136</sup> Different stimulus may influence the emergence of procoagulant MPs. An oxidative stimulus, namely H<sub>2</sub>O<sub>2</sub>, increases the production of procoagulant MPs by alveolar epithelial cells in culture. Increased number of TF-bearing MPs has been found in SSc interstitial lung disease.<sup>124</sup>

## Vascular health and microangiopathy

Few lines of evidence support the hypothesis that certain MPs subpopulation can induce angiogenesis and vascular remodeling. It has been postulated that expressing the VEGF, TF, and the protein sonic Hedgehog can define MPs as proangiogenic structures. Since MPs contain miRNA, they are able to activate a proangiogenic program in ECs.<sup>4</sup> Incubation of human microvascular endothelial cell line (HMEC-1) with THP-1 MPs leads to transfer of miRNA 150 from MP to recipient EC promoting angiogenesis, leading finally to the developed capillary-like structures out of existing blood vessels.<sup>31</sup> EMPs may transfer miRNA to ECs promoting angiogenesis.<sup>32</sup> In addition, activated subtype of EMPs positively correlate with number of ramified capillaries, indicating their role in angiogenesis and vessel regeneration.<sup>33</sup> In contrast, high levels of EMPs isolated from human umbilical vein ECs may reduce angiogenesis, while low concentration of EMPs stimulate formation of capillary-like structure (Figure 2).<sup>9</sup>

Recently, it has been shown within SSc patients that the active nailfold videocapillaroscopy (NVC) pattern is associated with higher concentration of activated E-selectin-positive EMPs (CD62+AnxV-) compared to early microvascular involvement. This subpopulation of EMPs is also increased in patients with specific microvascular alterations: pericapillary edema and giant capillaries or frequent



**Table 1.** Labeling of MPs and differences across studies.

|           | Labeling                                     | SSc vs HC | ISSc vs HC | dSSc vs HC | ISSc vs dSSc | Reference                              |
|-----------|--|-----------|------------|------------|--------------|--|
| tMP       |  | ↑**       | ↑*         | ↑*         |              | Guiducci et al. <sup>49</sup>          |
| AnxV- MPs | Total  | ↓**       | ns         | ns         |              | Iversen et al. <sup>59</sup>           |
|           | Total fraction                               | ↑*        |            |            |              | Iversen et al. <sup>59</sup>           |
| AnxV+ MPs | CD62E+                                       | ↑**       | ↑*         | ↑*         | ↑ns          | Michalska-Jakubus et al. <sup>50</sup> |
|           | Total  | ↓*        |            |            |              | Iversen et al. <sup>59</sup>           |
|           | CD62E+                                       | ↑**       | ↑*         | ↑*         | ↑ns          | Michalska-Jakubus et al. <sup>50</sup> |
|           | CD31+/CD42b-                                 | ↓*        |            |            |              | Jung et al. <sup>47</sup>              |
|           | CD31+/CD42b- or CD31+/CD42b+ or CD31-/CD42b+ | ↑ns       |            |            |              | McCarthy et al. <sup>51</sup>          |
| EMPs      | CD144+                                       | ↑**       | ↑*         | ↑**        | ↑ns          | Guiducci et al. <sup>49</sup>          |
|           | CD146+                                       | ↓*        |            |            |              | Iversen et al. <sup>59</sup>           |
|           | <sup>a</sup> CD146+                          | ↑ns       | ↑ns        | ↑ns        |              |  |
|           | CD31+/CD42b-                                 | ↑**       | ↑*         | ↑*         | ↑ns          | Michalska-Jakubus et al. <sup>50</sup> |
|           | CD51+  | ↑*        | ↑*         | ↑*         |              |  |
| PMPs      | CD42+  | ↑**       | ↑*         | ↑*         | ↑ns          | Guiducci et al. <sup>49</sup>          |
|           | <sup>b</sup> CD42a+                          | ↑ns       | ↑ns        | ↑ns        |              | Iversen et al. <sup>59</sup>           |
|           | CD42a+                                       | ↓*        | ↑**        |            |              | Nomura et al. <sup>52</sup>            |
|           | CD61+HMGB1+                                  | ↑**       |            |            |              | 111,116,131                            |
| LMPs      | <sup>c</sup> CD14+                           | ↑**       | ↑*         | ↑*         | ↑ns          | Guiducci et al. <sup>49</sup>          |
|           | <sup>d</sup> CD3+                            | ↑*        | ↑*         | ↑*         | ↑ns          |  |
|           | CD45+  | ↓*        | ↑ns        | ↑ns        |              | Iversen et al. <sup>59</sup>           |
|           | <sup>e</sup> CD45+                           | ↑ns       |            |            |              |  |

HC: healthy controls; tMP: total number of MPs; AnxV: annexin V; MP: microparticles; EMPs: endothelial cells-derived microparticles; PMPs: platelet-derived microparticles; LMPs: leukocyte-derived microparticles; SSc: systemic sclerosis; ISSc: limited systemic sclerosis; dSSc: diffuse systemic sclerosis.

<sup>a</sup>Fraction of AnxV-CD146+.

<sup>b</sup>Fraction of AnxV-CD42a+.

<sup>c</sup>Monocytes.

<sup>d</sup>T cells.

<sup>e</sup>Fraction of AnxV-CD45+.

\*p < .05; \*\*p < .001; ns p ≥ .05.

microhemorrhages, confirming that endothelial activation is enhanced in the active phase of SSc-related microangiopathy and also suggesting that their increased concentration might be a sensitive marker for early EC dysfunction. Furthermore, activated EMPs may reflect early step of angiogenesis since they positively correlate with number of ramified capillaries. Total number of EMPs is associated with the overall number of microvessels reflecting the severity of avascularizations. The confirmation proof of this is that the total number of EMPs is decreased in late NVC compared to early pattern and inversely correlates with number of ramified loops in SSc patients.<sup>50</sup> Apoptotic MPs phenotype (AnxV+ MP) has shown positive association with the avascular and microvasculopathy scores objected by NVC in the autoimmune disease patients with Raynaud's phenomenon, reflecting the existence of critical tissue hypoxia.<sup>48</sup> Furthermore, it has been shown that both higher annexin-positive EMPs and PMPs levels are associated with better digital perfusion assessed using laser speckle contrast imaging (LSCI) in patients with primary RP and SSc, reflecting vascular perfusion across diseases.<sup>51</sup>

It is well known that calcinosis and digital ulcers are associated with the late NVC pattern.<sup>137,138</sup> Regarding this, we could expect that in patients with this features of disease, total number of EMPs is decreased or apoptotic phenotype increased. Indeed, the significantly decreased numbers of both total MPs shading from various cells and PMPs have been demonstrated in patients with present cutaneous ulcers.<sup>49</sup> Furthermore, total EMPs levels also tended to be lower in SSc patients with active digital ulcers,<sup>50</sup> and higher levels of apoptotic EMPs (CD31+/CD42b-AnxV+) are associated with a history of digital ulceration/pitting.<sup>51</sup> In contrast, patients with calcinosis have increased level of activated (CD146+ AnxV-) EMPs subpopulation.<sup>53</sup>

### Inflammation and disease activity

MPs can activate complement cascade (C1q), enhance leukocyte rolling, and stimulate the release of broad proinflammatory mediators (e.g. IL-6 and IL-1 β; Figure 2). IL-6 has been implicated in the pathogenesis of SSc via stimulation of fibroblasts to produce excess collagen

and glycosaminoglycan, but to the best of our knowledge, no study so far has tested the role of MPs bearing IL6 in SSc.<sup>139</sup> Interaction between EMPs and naïve EC triggers proinflammatory response by upregulation of ICAM-1, messenger RNA (mRNA) expression, and soluble ICAM-1 shedding from target cells. Furthermore, EMPs which are triggered by transforming growth factor- $\alpha$  increase the release of soluble ICAM-1 secretion from ECs, enhancing the endothelial response to inflammation. This paracrine effect of MPs could not be observed using EMPs from unstimulated ECs, suggesting that these MPs may be both a consequence and a cause of the inflammatory response.<sup>9</sup>

In inflammation state, MPs have ability to transfer chemokine receptors and arachidonic acid between cells, leading to induction of adhesion molecules such as intracellular adhesion molecule-1 (ICAM-1) and VCAM-1. Adhesion molecules on EMPs can mediate adhesion of monocytes to ECs in vitro, leading to the maintenance of inflammation (Figure 2). Several studies have demonstrated increased sVCAM-1, sICAM1, sE-selectin, and sP-selectin levels in serum of SSc patients compared with the healthy controls,<sup>66,140,141</sup> raising the question: do MPs have together with adhesion molecules role in SSc pathogenesis. It has been found that sE- and sP-selectins<sup>142</sup> strongly correlate with either fraction (F) of AnxV-negative EMPs or PMPs in SSc with difference regarding to subtypes. In patients with diffuse systemic sclerosis (dSSc), association between F EMPs and both sP and sE selectins was observed, while in limited systemic sclerosis (lSSc), the only association was between PMPs and sP.<sup>59</sup>

MPs are capable of converting pentameric CRP into proinflammatory monomeric CRP. Furthermore, MPs containing CRP monomers can bind to the surface of ECs and generate proinflammatory signals in vitro.<sup>143</sup> In early inflammation, MPs may also induce immune cell apoptosis and the production of anti-inflammatory mediators such as IL8 predominately from LMPs (Figure 2).<sup>144</sup> CRP is one of the revised European Scleroderma Trials and Research group (EUSTAR) index component.<sup>145</sup> The highest number of total MPs has been found in a SSc patient with elevated CRP and an increased disease activity score of 3.5.<sup>49</sup> In spite of no study showing significant correlation between MPs and EScSG disease activity index score<sup>49,50</sup> so far, some data suggest that this association might exist. Thus, C3 complement, one of the EScSG component and possible marker of vascular injury,<sup>146</sup> inversely correlates with values of activated EMPs (CD62+ AnxV-).<sup>50</sup>

Recently, strong association has been demonstrated between EMPs (CD62+) levels and perivascular soft tissue inflammation, visualized by fluorescence optical imaging (FOI) in SSc patients.<sup>47</sup> An enhancement of fluorescence optical contrast media has been observed in vivo in the inflammatory tissue as visualized by FOI with an excellent correlation to histopathology.<sup>147</sup>

## Organ involvement

The levels of EMPs, MPs, and PMPs total number inversely correlate with the severity of skin involvement assessed by mRSS,<sup>49,50</sup> the best validated outcome measure for skin fibrosis in SSc.<sup>146</sup> Lower levels of PMPs and total MPs have been reported in patients with mRSS  $\geq 10$ ,<sup>49</sup> indicating that numbers of MPs could be associated with milder dermal fibrosis in SSc.

Recently, it has been shown that worse lung function measured by DLCO and FVC correlates with higher levels of both AnxV non-binding EMPs and LMPs, and these findings have been dissimilar in patients with limited and diffuse disease. In dSSc, increased concentration of both AnxV- EMPs and LMPs is related to a reduction of FVC, whereas in lSSc, the same MPs are associated with a reduction of DLCO. Furthermore, increased AnxV- EMPs have been found in patients with x-ray-confirmed lung fibrosis compared to cases without (frequency of ILD was higher in lSSc group).<sup>53</sup> The significantly increased concentration of both PMPs and monocytes-derived microparticles (mMPs) has been found in SSc patients with interstitial pneumonia (IP). PMPs-enhanced rsCD40L may stimulate the activation of monocytes and promote the production of mMPs from THP-1 suggesting the role of MPs in pathophysiology of progressive SSc with IP.<sup>52</sup> Furthermore, oxidized extracellular HMGB1, soluble or associated to PMPs, may amplify activation of neutrophils. Activated leukocytes and membrane HMGB1 are elevated in SSc patients with PAH or with diffuse subtype of disease.<sup>111</sup> PAH is associated with a specific pattern of platelets activation and higher fraction of HMGB1 + PMPs.<sup>131</sup>

## Conclusion

Although the knowledge about the role of MPs in fibrosis has recently advanced considerably, this research area still presents a great number of challenges. There is now accumulating evidence of the multiple faces of MPs as conveyors of cell information with major role in inflammation, thrombosis, and angiogenesis. MPs are undoubtedly implicated in immunopathogenesis. At present, the attention may be focused for the first time on the fact that MPs may have different behaviors. In fact, they can be antifibrotic and profibrotic as well. These particles may contribute to EndoMT and EMT via different mediators such as TGF- $\beta$ , PDGF, and HMGB1 protein. Furthermore, they may directly produce ROS through Nox-dependent processes, leading to the development and maintenance of oxidative stress, which is an important trigger of fibrosis. MPs are implicated in microangiopathy and clinical features of SSc. On the contrary, MPs contain proteolytic enzymes, such as MMPs involved into ECM degradation. Moreover, these particles may induce the synthesis in fibroblast of some MMPs, thus enhancing the ECM degradation

process. Expressing VEGF, TF, and the protein sonic Hedgehog, MPs may promote new vessel formation. The proven antifibrotic activity of MPs might serve as new therapeutic targets, opening new research avenues. More studies are warranted to provide novel insights into the world of MPs, to disclose their real potential as factors with a regenerative as well as an antifibrotic role.

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