Systemic sclerosis (SSc) is an autoimmune disease characterized by vasculopathy and uncontrolled cutaneous and internal organs fibrosis. Diagnosis of SSc in an early phase can be difficult because of a lack of typical symptoms. The delay in diagnosis and treatment of SSc may lead to uncontrolled progression of the disease, thus identification of possible early indicators of skin and organ involvement to prevent their further damage is necessary. The aim of this study is to review the latest biomarkers of organ involvement in SSc. In patients with lung fibrosis lung-epithelial-derived surfactant protein (SP-D), the glycoprotein Krebs von den Lungen-6 (KL-6), and chemokine ligands 2, 4 and 18 (CCL2, CXCL4, CCL18) are elevated, while in patients with skin fibrosis serum levels of heat shock protein 27 (Hsp27), interleukin 16 (IL-16), and IgG-galactosylation ratio are increased. Adiponectin concentration is inversely correlated with the intensity of cutaneous fibrosis. Skin gene profiling also seems very promising. In patients with heart involvement increased serum levels of brain natriuretic peptide (BNP) are present, as well as raised Midkine and Follistatin-like 3 (FSTL3) proteins, ratios of Cu/Se and ceruloplasmin (CP)/Circulating selenoprotein P (SELENOP) and higher whole blood viscosity level. Elevated calprotectin levels are found in individuals with gastrointestinal involvement. Increased levels of chemerin and ARA autoantibodies are associated with renal involvement, whereas high levels of adhesion molecules are found in patients with scleroderma renal crisis (SRC). Currently there are no biomarkers in use that can specifically identify the early involvement of organs.

**KEY WORDS:** systemic sclerosis, fibrosis, autoantibodies, biomarker

**ABSTRACT**

Systemic sclerosis (SSc) is an autoimmune disease characterized by vasculopathy and uncontrolled cutaneous and internal organs fibrosis. Diagnosis of SSc in an early phase can be difficult because of a lack of typical symptoms. The delay in diagnosis and treatment of SSc may lead to uncontrolled progression of the disease, thus identification of possible early indicators of skin and organ involvement to prevent their further damage is necessary. The aim of this study is to review the latest biomarkers of organ involvement in SSc. In patients with lung fibrosis lung-epithelial-derived surfactant protein (SP-D), the glycoprotein Krebs von den Lungen-6 (KL-6), and chemokine ligands 2, 4 and 18 (CCL2, CXCL4, CCL18) are elevated, while in patients with skin fibrosis serum levels of heat shock protein 27 (Hsp27), interleukin 16 (IL-16), and IgG-galactosylation ratio are increased. Adiponectin concentration is inversely correlated with the intensity of cutaneous fibrosis. Skin gene profiling also seems very promising. In patients with heart involvement increased serum levels of brain natriuretic peptide (BNP) are present, as well as raised Midkine and Follistatin-like 3 (FSTL3) proteins, ratios of Cu/Se and ceruloplasmin (CP)/Circulating selenoprotein P (SELENOP) and higher whole blood viscosity level. Elevated calprotectin levels are found in individuals with gastrointestinal involvement. Increased levels of chemerin and ARA autoantibodies are associated with renal involvement, whereas high levels of adhesion molecules are found in patients with scleroderma renal crisis (SRC). Currently there are no biomarkers in use that can specifically identify the early involvement of organs.
Diagnosis of SSc in an early phase is difficult because of the lack of typical signs and symptoms. The early detection and treatment of patients who have a high risk of disease progression is crucial to avoid organs disability in future. Unfortunately currently there are no biomarkers in use that can identify the early involvement of organs.

THE AIM

The aim of the study is to review the latest biomarkers of organ involvement in SSc reported in the literature.

REVIEW

BIOMARKERS IN SYSTEMIC SCLEROSIS

SKIN

Nowadays, the gold standard in the assessment of skin involvement is the modified Rodnan skin score (mRSS). It consists of palpation and pinching of the skin within 17 anatomical areas and assessment of skin thickness on a four-stage scale (0 to 3) within each of them. It is a non-invasive and inexpensive method, however it is considered to be subjective and depends on experience of the examining doctor. There is a need to find biomarkers that correlate with mRSS and can be objectively measurable.

Studies regarding heat shock protein 27 (Hsp27), which serves as a pro-inflammatory signaling molecule and helps the cell to survive under conditions of stress have been conducted. In Japanese study, serum levels of Hsp27 in 67 SSc patients were investigated, it was found that the serum levels of Hsp27 correlated positively with mRSS and were notably higher in patients with dcSSc than in those with lcSSc or healthy controls [10].

Another potential biomarker which has recently been investigated is the IgG–galactosylation ratio (IgG-Gal ratio), which has been found to be increased in immune diseases and certain cancers. In Chinese patients with SSC and healthy controls the IgG-Gal ratio was measured. Results showed that the IgG-Gal ratio was positively associated with mRSS and thus may be useful in assessing the severity of skin fibrosis and helpful in distinguishing between SSc subtypes [11].

Another marker worthy of attention is adiponectin, a hormone produced and secreted into the blood by activation of the nuclear receptor PPAR-γ which affects many metabolic processes. It was discovered that the level of adiponectin is inversely proportional to skin fibrosis and correlates negatively with the modified Rodnan skin score [12, 13].

Expression of IL-16 in SSc lesions and patients sera has also been examined. The correlation between serum IL-16 levels and the clinical symptoms of SSc in the skin was investigated. According to the results, the percentage of IL-16-positive cells was higher in patients with dcSSc than in healthy individuals. Patients with dcSSc also had increased serum levels of IL-16. Consequently, it has been suggested that IL-16 serum concentration may serve as biomarker to determine the severity of skin fibrosis [14].

In order to discover novel biomarkers scientists have started to study patients genes. Thus far the expression of genes regulated by TGFβ and IFNγ has been taken into account. Analysis of skin samples from patients with dcSSc revealed a four-gene biomarker panel consisting of THBS1, COMP, SIGLEC1, and IFI44; their expression correlated moderately well with mRSS [15, 16]. Another study discovered a two-gene pharmacodynamic biomarker (THBS1, MS4A4A) which may prove useful in the longitudinal assessment of skin involvement [17]. Recent studies have delved into the topic of epigenetics, including long noncoding RNAs (lncRNAs), increased plasma levels of TINCR, HOTTIP, and SPRY4-IT1 and decreased levels of ANCR were observed in SSc patients in comparison with the control group. Additionally, HOTTIP and SPRY4-IT1 were positively correlated with mRSS. ANCR and SPRY4-IT1 also correlated with PAH. Plasma SPRY4-IT1, HOTTIP, ANCR, and TINCR appear to be candidates for SSc biomarkers and SPRY4-IT1 may be used to predict the risk of SSc and define subtype [18].

These studies suggest that there is scope for more effective risk stratification and treatment selection for patients with SSc based on the expression of skin genes profiles.

LUNGS

Several molecules have been considered as biomarkers of lung involvement and many studies indicate that they may be very useful in the future.

It is important to mention endothelial microparticles (EMP) – submicron vesicles released in healthy and disease conditions from different cells which may modulate cell-cell signaling in vascular diseases. It was discovered that CD144+ EMP levels were significantly higher in the SSc-PAH patients. Moreover, the group of EMP molecules in SSc patients was more distinct [19]. EMP (especially CD144+) are potentially promising biomarkers of SSc-PAH and may be involved in its pathogenesis.

According to one study, in which the levels of lung-epithelial-derived surfactant protein (SP-D) and the glycoprotein Krebs von den Lungen-6 (KL-6) were assessed in SSc patients serum samples, SP-D is a relevant diagnostic biomarker of SSc-associated interstitial lung disease, whereas KL-6 could assess the severity of lung fibrosis [20].

The large groups of molecules which may be considered as biomarkers of lung changes are chemokine ligands and interleukins. Chemokine (C-C motif) ligand 2 (CCL2), also called monocyte chemotactant protein-1 (MCP-1), is released by a variety of immune cells and participates in the migration and activation of monocytes and T cells. It was reported that elevated serum CCL2 levels are associated with the presence of ILD and correlated with its severity [21]. Chemokine (C-X-C motif) ligand (CXCL4), known as platelet factor 4 (PF-4), plays a key role in cell migration, inflammation and wound repair; plasma CXCL4 levels were elevated in SSc patients and were associated with lung fibrosis and PAH [22]. Chemokine (C-C motif) ligand 18 (CCL18) is selectively chemotactic for T cells and has been shown to directly stimulate collagen production in fibroblasts [20]. It has been reported that serum CCL18 levels were elevated in SSc patients – according to a number of studies CCL18 appeared as a strong prognostic
marker for SSc-associated ILD progression as well as predictive biomarker for worsening of the lung disease in SSc [3, 20].

Results of some analyses have indicated that serum OX40L levels were elevated in SSc patients, especially in those diagnosed with dcSSc [23], but the latest studies suggest there is no significant association [20].

It has been suggested that the most solid biomarkers for diagnosis, classification, predicting organ involvement and prognosis in SSc are antinuclear autoantibodies. Each type of antibody is connected with a different SSc subset and is strongly predictive of the specific organ manifestations [24]. Anti-topoisomerase I antibodies are associated with ILD – it was reported that the presence of these antibodies correlated with a decline in differential forced vital capacity (FVC) levels [25]. Focusing on advances in the SSc antibody-biomarker field, recent studies have shown monospecific anti-Ro52/ tripartite motif-containing 21 (TRIM21) antibodies are associate with ILD and poor survival in SSc [26]. Moreover, many functional autoantibodies against cell surface receptors have also been identified in SSc. Specifically autoantibodies against the angiotensin II type 1 receptor (AT1R) and the endothelin-1 type A receptor (ETAR) have been reported as being predictive and prognostic biomarkers of SSc-PAH [27].

GASTROINTESTINAL TRACT
In SSc, only the skin is affected more often than the gastrointestinal tract [28]. Recent studies reported the role of fecal calprotectin (FC) as a potential biomarker of small intestinal bacterial overgrowth (SIBO). FC levels are elevated during active inflammation of the intestines. Measurement is uncomplicated and non-invasive, which makes it a very useful prospective test. A cohort study of 125 patients showed that elevated FC levels (>50mcg/g) were related to the severity of clinical digestive symptoms, delayed gastric emptying, and esophageal motor involvement. A strong association was reported between the presence of SIBO and a highly elevated FC levels (>275mcg/g). This test may help to identify a group of high-risk SSc patients, who require a glucose breath test – the assessment traditionally used to diagnose SIBO. The measurement of FC levels can also be used to evaluate the eradication of SIBO [29].

In patients with interstitial lung disease, the concentration of FC has also been found to increase. The role and usefulness of this observation in the early diagnosis of ILD are currently being investigated [30].

Presently available treatment of GI involvement is symptomatic and has short-time efficacy. No disease-modifying drugs are yet known. Identification of novel biomarkers could be helpful both for earlier diagnosis and the research of effective medications [6].

KIDNEYS
The kidneys are commonly involved in dcSSc. It was discovered that there is a strong association between scleroderma renal crisis (SRC) and anti-RNA polymerase III autoantibodies. Genetic susceptibility and altered protein expression were analyzed in kidney biopsy specimens from anti-RNAP III-positive SSc patients. The expression of two proteins – GPATCH2L and CTNND2 were increased in SRC in comparison with control samples. This may indicate the potential role of these proteins in the pathogenesis of SRC [31]. Findings reported by Liu et al. also demonstrated that anti-RNA polymerase III antibodies have predictive value for SSc-related renal crisis [32].

Additionally, elevated serum levels of circulating molecules: intercellular adhesion molecule-1 (ICAM-1), vascular cell adhesion protein-1 (VCAM-1) and E-selectin have also been found in patients with SRC [1].

It is also important to mention that biomarkers of activation of complement pathways were significantly elevated in SSc in comparison to healthy patients, and were similar to those found in other rheumatic diseases. Deposits of C3b molecules in some patients with SRC were also found [33]. Studies have shown that elevated levels of chemerin were related to an increased risk of renal injury, this may be due to direct damage of the kidneys or decreased chemerin clearance both in patients with SSc as well as in general population [13].

HEART
Right heart catheterization (RHC) is required to diagnose PAH. Due to invasiveness of this procedure, it is usually performed only in high-risk patients. Identification of possible indicators of heart involvement that would be appropriate for non-invasive screening is necessary. That will help to reduce the need for RHC and may lead to better therapeutic outcomes [34]. In clinical practice, the DETECT algorithm for PAH detection is commonly used, one of its components is the measurement of serum levels of the N-terminal fragment of pro-brain natriuretic peptide (NT-proBNP). This marker is elevated in SSc patients with PAH, but it is not specific for PAH or right ventricular dysfunction. It can also be increased in left heart dysfunction or in renal insufficiency [3, 35]. NT-proBNP appears to be a credible and independent predictor of mortality in patients with SSc [36], the level of this marker is also associated with skin fibrosis and is higher in patients with the diffuse rather than limited type of SSc [8].

Moreover, a negative correlation between the level of miRNA let-7d and PAH has been observed, suggesting a potential role of miRNA let-7d as a biomarker for the presence and severity of PAH in SSc [37].

In addition, two proteins, Midkine and Follistatin-like 3 (FSTL3) combined, may act as SSc-PAH indicators and have been identified as potential drug targets [34].

Recent studies have shown that patients with SSc-related PAH have elevated ratios of Cu/Se and ceruloplasmin (CP)/Circulating selenoprotein P (SELENOP) when compared to controls – this implies that these parameters may help to identify the risk of SSc-related PAH. Furthermore, a shortage of selenium in patients with skin fibrosis may indicate a deficiency of selenoenzymes needed to slow down the pathogenesis of scleroderma and successful control of oxidative stress [38].

A recent cohort study showed that higher whole blood viscosity levels in SSc patients can serve as an independent indicator for PAH development. However, this needs to be confirmed by further research [39].
The main aim of this review is to discuss the latest biomarkers of organ changes in SSc and underline their importance. Taking into account the results of the search for biomarkers correlating with mRSS the most recent are heat shock protein 27, IgG–galactosylation ratio, and IL-16 [10, 11, 14]. It also appears to be very promising to study skin gene profiles of SSc patients, including epigenetics (Table 1) [15-18].

Results of one of the latest studies appear to show that the serum level of KL-6 may reliably assess ILD severity to optimize further patient’s management, while the serum level of SP-D is useful for early identification of SSc patients having developed ILD. The same study also showed that there is no confirmation that the serum level of OX40L may be associated with lung fibrosis [20]. Many studies have proven the levels of chemokines (CCL2, CCL18, CXCL4) are reliable biomarkers for activity as well as severity of the lung fibrosis (Table 2) [20-22]. Measurement of fecal calprotectin level seems to be useful in identifying SSc patients at high risk of gastrointestinal tract involvement [29].

In case of renal involvement, even though SRC has become quite a rare complication in SSc patients, most of the research is focused on this life-threatening complication [7, 40]. The latest research supports a potential role for altered Wnt signaling in SRC pathogenesis and indicates that anti-RNA polymerase III autoantibodies seem to be one of the most promising biomarker of renal involvement in SSc [31, 32] as well as chemerin, which high level is associated with renal function impairment [13].

In diagnosis of patients with PAH the measurement of NT-proBNP is already in use [35]. However, the search for more specific markers is in progress. The most recent studies have concerned expression of miRNA let-7d, Midkine and Follistatin-like 3 (FSTL3), ratios of Cu/Se and CP (ceruloplasmin)/SELENOP(Circulating selenoprotein P), and whole blood viscosity level [34, 37-39]. (Table 1)

**Table 1. Potential biomarkers of organ involvement in SSc patients.**

<table>
<thead>
<tr>
<th>ORGAN</th>
<th>BIOMARKER</th>
</tr>
</thead>
<tbody>
<tr>
<td>SKIN</td>
<td>Heat shock protein 27</td>
</tr>
<tr>
<td></td>
<td>IgG–galactosylation ratio</td>
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<tr>
<td></td>
<td>Adiponecin</td>
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<tr>
<td></td>
<td>Interleukin 16</td>
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<tr>
<td></td>
<td>THBS1, COMP, SIGLEC1 and IFI44 four-gene signature</td>
</tr>
<tr>
<td></td>
<td>THBS1 and MS4A4A two-gene signature</td>
</tr>
<tr>
<td></td>
<td>SPRY4-IT1, HOTTIP, ANCR, and TINCR</td>
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<tr>
<td>GASTROINTESTINAL TRACT</td>
<td>Fecal calprotectin</td>
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<tr>
<td>HEART</td>
<td>NT-proBNP</td>
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<tr>
<td></td>
<td>miRNA let-7d</td>
</tr>
<tr>
<td></td>
<td>Midkine and Follistatin-like 3 (FSTL3)</td>
</tr>
<tr>
<td></td>
<td>Ratios of Cu/Se and CP (ceruloplasmin)/SELENOP(Circulating selenoprotein P)</td>
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<tr>
<td></td>
<td>Whole blood viscosity level</td>
</tr>
<tr>
<td>LUNGS</td>
<td>CD144+ EMP</td>
</tr>
<tr>
<td></td>
<td>SP-D and KL-6</td>
</tr>
<tr>
<td></td>
<td>CCL2, CCL18, CXCL4</td>
</tr>
<tr>
<td></td>
<td>IL-6</td>
</tr>
<tr>
<td></td>
<td>Endothelin-1</td>
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<tr>
<td></td>
<td>Autoantibodies: Anti- Ro52/TRIM, Agonistic anti-AT1R, Agonistic anti-ETAR, Anti-topoisomerase I, Anti-Th/To</td>
</tr>
<tr>
<td></td>
<td>anti-RNAP III autoantibodies</td>
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<tr>
<td></td>
<td>ICAM-1, VCAM-1</td>
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<tr>
<td>KIDNEYS</td>
<td>GPATCH2L and CTNND2</td>
</tr>
<tr>
<td></td>
<td>C3b molecules</td>
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<td></td>
<td>Chemerin</td>
</tr>
</tbody>
</table>

NT-proBNP - N-terminal fragment of pro-brain natriuretic peptide; EMP- endothelial microparticles; SP-D - lung-epithelial-derived surfactant protein; KL-6 - glycoprotein Krebs von den Lungen-6; CCL2 - Chemokine (C-C motif) ligand 2 = monocyte chemoattractant protein-1 (MCP-1); CXCL4 - Chemokine (C-X-C motif) ligand = platelet factor 4 (PF-4); CCL18 - Chemokine (C-C motif) ligand 18; Anti- Ro52/TRIM - anti-Ro52/tripartite motif-containing 21 antibodies; Agonistic anti-AT1R - autoantibodies against the angiotensin II type 1 receptor; Agonistic anti-ETAR - autoantibodies against the endothelin-1 type A receptor; anti-RNAP III autoantibodies - anti-RNA polymerase III autoantibodies; ICAM-1 - intercellular adhesion molecule-1; VCAM-1 - vascular cell adhesion protein.
CONCLUSION

The outcomes of this review suggest that there are many molecules that can serve as biomarkers in the diagnosis, assessment of disease activity, and prediction of internal organs complications in SSc. Moreover, we may potentially classify patients into different risk groups, to improve treatment implementation and its efficacy. Nevertheless, advanced research including bigger groups of participants is needed for validation.

REFERENCES


Table 2. SSc lung changes with potential biomarkers.

<table>
<thead>
<tr>
<th>Biomarkers</th>
<th>Association</th>
<th>Value of the biomarker</th>
</tr>
</thead>
<tbody>
<tr>
<td>Microparticles</td>
<td></td>
<td></td>
</tr>
<tr>
<td>EMP CD144+</td>
<td>PAH</td>
<td>Predictive</td>
</tr>
<tr>
<td>Proteins</td>
<td></td>
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</tr>
<tr>
<td>SP-D</td>
<td>Lung fibrosis</td>
<td>Diagnostic</td>
</tr>
<tr>
<td>KL-6</td>
<td>Lung fibrosis</td>
<td>Severity</td>
</tr>
<tr>
<td>Chemokines</td>
<td></td>
<td></td>
</tr>
<tr>
<td>CCL18</td>
<td>Lung fibrosis</td>
<td>Predictive, Prognostic</td>
</tr>
<tr>
<td>CCL2</td>
<td>Lung fibrosis</td>
<td>Severity, Activity</td>
</tr>
<tr>
<td>CXCL4</td>
<td>Lung fibrosis, PAH</td>
<td>Predictive</td>
</tr>
<tr>
<td>Interleukins</td>
<td></td>
<td></td>
</tr>
<tr>
<td>IL-6</td>
<td>Lung fibrosis</td>
<td>Predictive</td>
</tr>
<tr>
<td>Endothelin-1</td>
<td>PAH</td>
<td>Severity</td>
</tr>
<tr>
<td>Autoantibodies</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Anti-Ro52/TRIM21</td>
<td>ILD</td>
<td>Prognostic</td>
</tr>
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<td>Agonistic anti-AT1R</td>
<td>PAH</td>
<td>Predictive, Prognostic</td>
</tr>
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<td>Antithoposomerase I</td>
<td>ILD, Lung fibrosis</td>
<td>Predictive</td>
</tr>
<tr>
<td>Anti-Th/To</td>
<td>ILD, Lung fibrosis</td>
<td>Predictive</td>
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</table>

EMP- endothelial microparticles; SP-D - lung-epithelial-derived surfactant protein; KL-6 - glycoprotein Krebs von den Lungen-6; CCL2 - Chemokine (C-C motif) ligand 2 = monocyte chemoattractant protein-1 (MCP-1); CXCL4 - Chemokine (C-X-C motif) ligand = platelet factor 4 (PF-4); CCL18 - Chemokine (C-C motif) ligand 18; IL-6 - Interleukin 6; Anti-Ro52/TRIM - anti-Ro52/tripartite motif-containing 21 antibodies; Agonistic antiAT1R - autoantibodies against the angiotensin II type 1 receptor; Agonistic anti-ETAR - autoantibodies against the endothelin-1 type A receptor; PAH – pulmonary arterial hypertension; ILD – interstitial lung disease


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Conflict of interest

Authors declare no conflict of interest.

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