ABSTRACT

Objectives: This study aims to investigate the distribution of myeloid-derived suppressor cells (MDSCs) in patients with primary or secondary Sjögren’s syndrome (SS) or rheumatoid arthritis (RA) in order to better understand MDSCs significance in the pathogenesis of these autoimmune diseases.

Patients and methods: We examined the frequency and calculated absolute counts of overall MDSCs (human leukocyte antigen-antigen D related (HLA-DR)low/cluster of differentiation (CD) 33+/CD11b+) and monocytic MDSCs (HLA-DRlow/CD33+/CD11b+/CD14+) subset in peripheral blood samples of 23 RA (5 males, 18 females; mean age 57 years; range 41 to 81 years), 25 primary Sjögren’s syndrome (pSS) (1 male, 24 females; mean age 56 years; range 32 to 77 years), 17 secondary Sjögren’s syndrome (sSS) (1 male, 16 females; mean age 60 years; range 49 to 73 years) and 23 nonautoimmune sicca syndrome (nSS) (23 females; mean age 59 years; range 44 to 92 years) patients by flow cytometric analysis.

Results: Analysis revealed that the frequency of overall MDSCs increased in RA group (46.5±3.4) compared with nSS group (35.6±3.2; p=0.0322). An increase of absolute count of overall MDSCs was most evident in both RA (4383±456.8) and sSS groups (3890±495.7) compared with pSS (2447±275.1; p=0.0002 and 0.0067) and nSS groups (2025±218.1; p<0.0001 and p=0.0012). The highest absolute count of monocytic MDSCs also manifested in RA group (195.4±39.0), compared with all the other groups (86.0±24.9; p=0.0002 [pSS], 128.5±53.4; p=0.0076 [sSS], 83.7±19.0; p=0.0136 [nSS]).

Conclusion: To summarize, we have determined that the most prominent increase of both total and monocytic MDSCs was evident in RA and sSS groups, which leads us to believe that MDSCs are associated with rheumatic processes.

Keywords: Flow cytometry; myeloid-derived suppressor cells; rheumatoid arthritis; Sjögren’s syndrome.

In recent years, more attention has been paid to myeloid-derived suppressor cells (MDSCs) and their importance in cancer and other illnesses. Increasing numbers of studies suggest that the expansion of these regulatory cells may be a common response to various forms of inflammation.1

Myeloid-derived suppressor cells are defined as heterogeneous cell population and they include myeloid progenitor and immature myeloid cells. In steady state, MDSCs reside mostly in bone marrow, but in presence of various pathological conditions, they can expand and be detected in peripheral lymphoid and cancerous tissues, blood
stream, the spleen and inflammatory sites. 2,3 Due to their heterogeneity both in morphology and function, MDSCs lack one specific marker of identification. 4

In murine models, researchers characterize MDSCs by the expression of granulocyte-differentiation antigen (Gr-1) and Crohn’s disease (CD11b) markers which represent a mixture of immature myeloid cells, myeloid progenitor cells, monocytes-macrophages, immature granulocytes and dendritic cells. 5,6 Both murine and human MDSCs can be subdivided into more monocytic and more granulocytic subtypes. Human MDSCs are divided into CD14+ monocytic and CD15+ granulocytic subtypes. Both of the subtypes express myeloid specific markers CD11b and CD33, but lack the expression of human leukocyte antigen-antigen D related (HLA-DR) and other mature myeloid cells markers. 4

A majority of research on MDSCs is performed in terms of cancer; however, it is already known that MDSCs are involved in various autoimmune diseases such as systemic lupus erythematosus, 7 autoimmune type 1 diabetes, 8 autoimmune hepatitis, 9 and also in viral, 10 bacterial 11 and parasitic 12 infections, sepsis, 13 inflammation 14 or other pathological conditions. 15 There are limited data available concerning MDSCs in case of rheumatoid arthritis (RA) and no data in case of rheumatoid arthritis (SS). Fujii et al. 16 showed that in a collagen-induced arthritis (CIA) mouse model, MDSC lowered the number of T helper cells and suppressed the progression of CIA. The findings of MDSCs as immune response suppressors, functioning directly or indirectly through the induction of regulatory T cells, are summarized in a review article of Greten et al. 4 Therefore, in this study, we aimed to investigate the distribution of MDSCs in patients with primary or secondary SS or RA in order to better understand MDSCs significance in the pathogenesis of these autoimmune diseases.

PATIENTS AND METHODS

Patients who participated in the study were recruited from the Rheumatology Centre of Vilnius University Hospital Santaros Klinikos between September 2014 and December 2016. Of the patients included, 25 (1 male, 24 females; mean age 56 years; range 32 to 77 years) had primary Sjögren’s syndrome (pSS), 17 (1 male, 16 females; mean age 60 years; range 49 to 73 years) had secondary Sjögren’s syndrome (sSS) due to RA, 23 (5 males, 18 females; mean age 57 years; range 41 to 81 years) had RA and 23 (23 females as control group; mean age 59 years; range 44 to 92 years) had nonautoimmune sicca syndrome (nSS). The diagnoses of pSS and sSS were based on the criteria defined by the American-European Consensus Group criteria for SS. 17 RA was diagnosed according to 2010 American College of Rheumatology (ACR)/European League Against Rheumatism (EULAR) criteria for RA. 18 None of the patients in the control group met the SS classification criteria and they were classified as patients with nSS. All study participants underwent extensive serological evaluation (Table 1). None of the patients was on immunosuppressive medications at the time of the study. A written informed consent was obtained from each subject. The study protocol was approved by the Ethics Committee for Biomedical Research in Vilnius region (2014-05-20, No. 158200-14-733-248). The study was conducted in accordance with the principles of the Declaration of Helsinki.

Multicolor flow cytometric analysis was carried out to determine the frequency of MDSC. The following anti-human monoclonal antibodies were used in the study: anti-CD14 (phycoerythrin [PE]), anti-CD11b (fluorescein isothiocyanate [FITC]), anti-CD33 (allophycocyanin [APC]), anti-HLA-DR (peridinin chlorophyll protein complex [PerCP]) (all from BioLegend, San Diego, CA, USA). 50 µL of heparinized venous blood was stained with appropriate amounts of monoclonal antibodies (according to manufacturer’s recommendations) in flow cytometric test tubes. Cell dyeing protocol was carried out as described previously 19 and analyzed using FACSCalibur flow cytometer (BD Biosciences, San Jose, CA, USA) and CELLQuest software (BD Biosciences, San Jose, CA, USA). Flow cytometric gating strategy of total and monocytic MDSCs (mMDSCs) in one patient is presented in Figure 1.

Serum samples of study participants were stored at -80°C until analysis. Serum anti-Sjögren’s syndrome-related antigen A (anti-Ro/SSA) and anti-Sjögren’s syndrome-related antigen B (anti-La/SSB) specific
Table 1. Clinical and serological characteristics of study participants

<table>
<thead>
<tr>
<th></th>
<th>RA (n=23)</th>
<th>pSS (n=25)</th>
<th>sSS (n=17)</th>
<th>nSS (n=23)</th>
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<tbody>
<tr>
<td></td>
<td>n %</td>
<td>Mean±SD</td>
<td>Mean Range</td>
<td>n %</td>
</tr>
<tr>
<td>Age (year)</td>
<td>57 41-81</td>
<td></td>
<td></td>
<td>56 32-77</td>
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<tr>
<td>Disease duration</td>
<td>13±12</td>
<td>8±5</td>
<td>14±10</td>
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<td>4.9±1.3</td>
<td>-</td>
<td>5.7±2.0</td>
<td>-</td>
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<tr>
<td>Ocular symptoms</td>
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<td>2.2±0.9</td>
<td>2.2±1.1</td>
<td>1.9±1.0</td>
</tr>
<tr>
<td>Oral symptoms</td>
<td>0.4±0.7</td>
<td>1.7±0.6</td>
<td>1.4±0.8</td>
<td>0.9±0.7</td>
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<tr>
<td>Duration of dryness (year)</td>
<td>1±2</td>
<td>5±4</td>
<td>4±4</td>
<td>5±6</td>
</tr>
<tr>
<td>Schirmer’s I test positive (≤5 mm/5 min)</td>
<td>2 8.7</td>
<td>21 84</td>
<td>17 100</td>
<td>8 34.8</td>
</tr>
<tr>
<td>Schirmer’s I test (mm/5 min)</td>
<td>20.0±10.3</td>
<td>4.3±4.6</td>
<td>3.1±1.3</td>
<td>7.0±5.5</td>
</tr>
<tr>
<td>Unstimulated salivary flow positive (≤1.5 mL/15 min)</td>
<td>0 0</td>
<td>20 80</td>
<td>10 58.8</td>
<td>3 13</td>
</tr>
<tr>
<td>Unstimulated salivary flow (mL/15 min)</td>
<td>3.8±1.4</td>
<td>1.5±1.0</td>
<td>1.7±1.0</td>
<td>2.6±1.3</td>
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<tr>
<td>Focus score (number of lymphocytic foci/4 mm²)</td>
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<td>1.8±0.7</td>
<td>-</td>
<td>-</td>
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<td>Positive autoantibodies</td>
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<td></td>
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<tr>
<td>RF</td>
<td>23 100</td>
<td>10 40.0</td>
<td>17 100</td>
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<tr>
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<td>17 73.9</td>
<td>1 4</td>
<td>8 47</td>
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<tr>
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<td>18 72</td>
<td>4 23.5</td>
<td>-</td>
</tr>
<tr>
<td>Anti-Ro/SSA⁺</td>
<td>1 4.3</td>
<td>16 64</td>
<td>3 17.6</td>
<td>1 4.3</td>
</tr>
<tr>
<td>Anti-La/SSB⁺</td>
<td>- -</td>
<td>14 56</td>
<td>2 11.8</td>
<td>-</td>
</tr>
<tr>
<td>Anti-Ro/SSA/Anti-La/SSB⁺</td>
<td>- -</td>
<td>14 56</td>
<td>2 11.8</td>
<td>-</td>
</tr>
</tbody>
</table>

RA: Rheumatoid arthritis; pSS: Primary Sjögren’s syndrome; sSS: Secondary Sjögren’s syndrome; nSS: Nonautoimmune sicca syndrome; SD: Standard deviation; DAS28: Disease activity score 28; RF: Rheumatoid factor; ACCP: Anti-cyclic citrullinated peptide; ANA: Antinuclear antibodies; Anti-Ro/SSA: Anti-Sjögren’s syndrome-related antigen A; Anti-La/SSB: Anti-Sjögren’s syndrome-related antigen B.
antibody levels were measured by enzyme-linked immunosorbent assay (ELISA) by use of the Anti-SSA (Ro) antibodies enzyme immunoassay and Anti-SSB (La) antibodies enzyme immunoassay kits (BioSystems S.A., Barcelona, Spain) according to manufacturer’s recommendations. The absorbance of samples was read at 450 nm with microplate reader (BioTek Instruments Inc., Winooski, VT, USA). The concentration of antibodies present in the sample was calculated by interpolating the absorbance in four parametric calibration curve using Gen5 Microplate Data Collection & Analysis Software (BioTek Instruments Inc., Winooski, VT, USA). Concentration of anti-Ro/SSA and anti-La/SSB antibodies in serum sample greater than 12.5 U/mL was considered positive.

**Statistical analysis**

Mann-Whitney U test was performed to determine the statistical differences using GraphPad Prism 6.0 software (GraphPad Software, San Diego, CA, USA). The data are expressed as mean ± standard error of the mean. Correlations were determined with Spearman rank correlation tests. P values less than 0.05 were considered statistically significant.

**RESULTS**

The median frequency of total MDSCs within leukocytes in RA group (46.5±3.4) was significantly increased compared to control

![Flow cytometric analysis of total and monocytic MDSC](image-url)
(nSS) group (35.6±3.2; p=0.0322) (Figure 2a). However, in case of absolute counts of total MDSCs in peripheral blood of study participants, more significant differences arose. Although the increase of MDSCs in RA group remained (4383±456.8; p<0.0001), the absolute count of MDSCs increased in sSS group (3890±495.7) in comparison to control group (2025±218.1; p=0.0012). The absolute counts in both groups were also significantly higher compared to pSS group (2447±275.1; p=0.0002 and p=0.0067, respectively) (Figure 2b).

The highest frequency of mMDSCs was observed in RA group (2.1±0.4), but significant differences were established only with pSS and sSS groups (1.3±0.3; p=0.0141 and 1.2±0.4; 0.0047, respectively) (Figure 2c). In case of absolute count of mMDSCs, the tendency remained the same - the highest absolute count was observed in RA group (195.4±39.0), but in this respect, the increase was significant compared to all the other study groups (86.0±24.9; p=0.0002 [pSS], 128.5±53.4; p=0.0076 [sSS], 83.7±19.0; p=0.0136 [nSS]) (Figure 2d).

We also analyzed the distribution of both total and mMDSCs in anti-Ro/SSA and anti-La/SSB positive patients (Table 1), instead of all study participants in pSS group. Even though the p values changed slightly (Figure 3a-d), no additional significant changes were observed.

In our previous study, we analyzed plasmacytoid (pDC) and conventional dendritic cells (cDC), cytotoxic T lymphocytes (CTL), natural killer (NK) and natural killer T (NKT) cells in peripheral blood of the same study subjects as in this study. Consequently, we recalculated correlations and established associations.
between MDSCs and other peripheral blood cell populations. For this purpose, only the same patient data as for MDSC analysis were selected and correlations were calculated both in frequencies and absolute counts.

We determined that in RA group, the frequency of total MDSCs negatively correlated with cDC (p=0.0028, r = -0.5948). Both the frequency and absolute count of mMDSCs also moderately correlated with cDC (p=0.0156, r=0.4977 and p=0.0362, r=0.4387, respectively), however, the correlations were positive. Furthermore, correlations between the frequency of total MDSCs and CTL were established (“classical” memory cells CD8$^{\text{high}}$/CD45RA$^{-}$ p=0.0126, r= -0.5115; effector memory CTL subtype CD8$^{\text{high}}$/CD45RA$^{-}$ p=0.0353, r=0.4407).

In sSS group, the frequency and absolute count of total MDSCs positively correlated with CD8$^{\text{high}}$/CD45RA$^{-}$ CTL subtype (p=0.0406, r=0.5007 and p=0.0267, r=0.5357, respectively), while the frequency of mMDSCs negatively correlated only with CD8$^{\text{high}}$/CD45RA$^{-}$ CTL subtype (p=0.0143, r= -0.5816).

In pSS group, both the frequency and absolute count of mMDSCs positively correlated with CD8$^{\text{high}}$/CD45RA$^{-}$ CTL subtype (p=0.0425, r=0.4088 and p=0.0249, r=0.4474, respectively). The absolute count of mMDSCs also correlated with cDC (p=0.0066, r = 0.5285). Correlations with NK and NKT cells were not established in all of the study groups (RA, pSS and sSS); however, in control group, mMDSCs positively correlated with NK cells (p=0.0410, r=0.4494).

**Figure 3.** The distribution of total and monocytic MDSC in the peripheral blood of RA, anti-Ro/SSA and anti-La/SSB positive pSS*, sSS and nSS patients. (a, c) Plots indicate frequencies and (b, d) absolute counts of total and monocytic MDSC, respectively. Each point represents one patient; horizontal lines represent mean value ± SEM. Analysis was performed by Mann-Whitney U test. MDSC: Myeloid-derived suppressor cells; RA: Rheumatoid arthritis; anti-Ro/SSA: Anti-Sjögren’s syndrome type A; anti-La/SSB: Anti-Sjögren’s syndrome type B; pSS: Primary Sjögren’s syndrome; sSS: Secondary Sjögren’s syndrome; nSS: Nonautoimmune sicca syndrome; SEM: Standard error of the mean.
**DISCUSSION**

The bone marrow is stimulated to release MDSCs into the bloodstream to protect the host from harmful excessive immune stimulation in case of acute or chronic infection, and to limit the formation of an autoimmune response to tissue antigens released during injury. The role and molecular mechanisms of MDSCs in human autoimmune diseases are complicated and still unclear. Even though most MDSC studies are carried out in murine models,9,20,21 the number of human MDSC studies is increasing.22,23 It is already known that MDSCs can inhibit functions of various T cell populations in many ways. Zhu et al.24 demonstrated that in experimental autoimmune encephalomyelitis model, mMDSCs exhibit strong suppressive effect on activated T cells and participate in T cell inhibition by producing nitric oxide. Nitric oxide production by MDSCs result in nitrosylation of cysteine residues, which destabilize messenger ribonucleic acid, thus preventing the production of cytokines necessary for T cell proliferation.25,26

There is limited research on the distribution of MDSCs in peripheral blood of patients with autoimmune diseases. In some cases, authors report that in animal models, MDSCs decrease the severity of autoimmune processes,27,28 while other studies show that MDSCs are associated with a worse prognosis of the disease.29,30 Jiao et al.23 determined that in RA patients, circulating MDSC population increased significantly compared to healthy individuals and these findings coincide with our results. Furthermore, they established a negative correlation between MDSCs and Th17 cells, which confirms the association between them. We have also calculated the correlations between MDSCs and other peripheral blood cell populations and found that these cells correlate with cDC and a few subpopulations of CTL, which also confirms the association between MDSCs and T cells.

For a long time, it has been assumed that T cells play the main role in SS pathogenesis; however, more recent research claim that B cells are not only crucial in SS pathogenesis, but are also the main cells participating in the development of the disease.31 It is also known that there are almost five times more B cells than T cells in salivary gland infiltrate of pSS patients.32 The fact that B cells play the main role in pSS disease may be the reason why we observed significantly increased frequency of total MDSCs only in RA and a significant increase of absolute count of total MDSCs in RA and sSS study groups. In pSS group, the absolute count of both total and mMDSCs was almost the same as in our control group. It is known that anti-Ro/SSA and anti-La/SSB autoantibodies are associated with a more severe course of the disease: earlier onset, decreased salivary flow, more intense eye symptoms, worse Schirmer's test etc.33,34 However, after narrowing down pSS group to only anti-Ro/SSA and anti-La/SSB positive patients, results did not change. Nevertheless, we have established a correlation between mMDSC and cDC, as well as a CTL subtype.

Secondary Sjögren’s syndrome is a common manifestation in patients with RA, its prevalence varying from 4 to 50%.35 This subgroup of patients has distinct clinical, immunological and genetic profiles.36 Observation studies indicate that RA and sSS have different outcomes and patients with sSS have two-fold higher risk of non-Hodgkin’s lymphoma and higher mortality rate.37 In our study, we have determined that total MDSCs in sSS increased, compared to pSS and nSS and were comparable to RA; however, mMDSCs in sSS increased only slightly and were comparable to pSS and nSS. It is known that granulocytic and monocytic MDSCs have distinct molecular properties and distinct gene expression profiles but also opposing effects on tumor cells.38 This can also be assumed in terms of autoimmune diseases. Our presumption is that granulocytic MDSCs play a more significant role in both RA and sSS. Moreover, in this group, we established correlations between total and mMDSCs and different CTL subpopulations.

One of the downfalls of our study is that we only measured the distribution of MDSCs and mMDSCs in RA, pSS and sSS study groups. We observed increased frequency of total MDSCs in all study groups compared to control group, yet the frequency of mMDSCs increased only in RA group. Due to the differences between total and mMDSC populations, further and more extensive studies of MDSC subpopulations should be conducted, distinguishing granulocytic CD15+ MDSCs and other subpopulations. Furthermore, larger sample sizes are required. Different factors
(prostaglandins, cyclooxygenase-2, interleukin 6, granulocyte-macrophage colony-stimulating factor, macrophage colony-stimulating factor, interferon gamma, transforming growth factor beta and tumor necrosis factor) are responsible for MDSC expansion, proliferation and inhibition of differentiation into mature cells.\textsuperscript{39-42} Assessment of serum factor levels of study patients might give us an insight as to why there is such a difference in MDSC distribution between these two autoimmune diseases (RA and pSS).

In conclusion, we have determined that the most prominent increase of both total and mMDSCs was evident in RA and sSS groups, which leads us to believe that MDSCs are associated with rheumatic processes. Furthermore, to the best of our knowledge, this is the first study to analyze the distribution of MDSCs in peripheral blood of pSS and sSS patients and to compare the results with RA.

Declaration of conflicting interests
The authors declared no conflicts of interest with respect to the authorship and/or publication of this article.

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