Different T cells’ distribution and activation degree of Th17 CD4+ cells in peripheral blood in patients with osteoarthritis, rheumatoid arthritis, and healthy donors: preliminary results of the MAGENTA CLICAO study

Alfredomaria Lurati1
Antonella Laria1
Arianna Gatti2
Bruno Brando2
Magda Scarpellini1

1Rheumatology Unit, Fornaroli Hospital, Magenta, Italy; 2Transfusional Centre, Legnano Hospital, Legnano, Italy

Objective: To determine distribution of T cells and activation degree of Th CD4+ cells in peripheral blood of patients with osteoarthritis (OA), rheumatoid arthritis (RA), and healthy donors.

Methods: Patients with established diagnosis of RA according to American College of Rheumatology/European League Against Rheumatism 2010 criteria, knee or hip OA according to American College of Rheumatology criteria, and healthy blood donor volunteers were eligible. Multi-channel flow cytometry and monoclonal antibodies against CD3, CD4, CD8, CCR6, CD38, CXCR3, and HLA DR were used to distinguish and evaluate T cells’ subpopulation.

Results: We analyzed blood samples of 15 patients with well-defined RA, 56 with hip or knee OA, and 20 healthy age matched controls. Blood samples from RA patients showed significantly higher counts of CD4+ CD38+ DR+ (activated CD4 T cells) and Th17 (CCR6+ CXCR3−) cells as compared to OA patients and control group (P<0.01). Furthermore the samples from the OA patients showed a higher percentage of activated CD4 T cells and Th17 cells as compared to control group (P<0.05). Interestingly there was no difference between Th1 (CD4+ CXCR3+ CCR6−) and Th2 (CD4+ CXCR3− CCR6−) between the three groups (P>0.1).

Conclusion: According to the latest view of OA disease pathogenesis, our preliminary results support the hypothesis that OA may also be a disease with an immunological/inflammatory involvement like RA. It seems that there is a quantitative but non-qualitative difference in Th17 cells’ profile, including the expression of activation markers, between RA and OA.

Keywords: Th17, flow cytometry, osteoarthritis

Introduction
Osteoarthritis (OA) is a chronic, painful, disabling condition affecting the whole joint (bone, synovia, and cartilage), well defined clinically and radiologically, but its etiology is largely poorly understood.1–4 In a pro-inflammatory milieu, chondrocytes become metabolically active and initiate inflammatory processes that secrete several inflammatory cytokines that work synergistically to stimulate synthesis of enzymes that break down cartilage. Immunohistochemical studies have confirmed that mononuclear cell infiltration, lymphocytes and macrophages, and production of pro-inflammatory cytokines and mediators of joint damage are common synovial membrane (SM) characteristics in patients with OA.5–7 The study of OA synovial fluid (SF) has rendered similar findings.5–13 Mononuclear cell infiltrates in synovial tissues

Correspondence: Alfredomaria Lurati
Rheumatology Unit, Fornaroli Hospital, Via Donatore
Sangue 50, 20013 Magenta, Italy
Tel +39 02 9796 3843
Email alfredomaria.lurati@ao-legnano.it
The Th1 subset of T cells was found to be approximately five times more than Th2 cells and higher levels of Th1 cytokines, IL2, and INF-gamma, were detected in most OA patients. Data from literature show that CD4+/CD8+ ratio in OA ST is approximately 5:1 compared to normal ST where the ratio is 2:1 and the higher ratio is comparable to rheumatoid arthritis (RA) ST. Furthermore, T cells’ CD4+ levels are higher in early OA compared to late stage OA. Studies on IL-1 levels in SF specimens from different arthropathies, including OA, have shown that the range of IL-1 levels in OA are similar than in RA, a chronic inflammatory disease known to be characterized by the presence of a prominent cellular infiltrate in the SM and the SF. T lymphocytes represent the most important cell type for immune functions and are responsible for specific immunity involved in the arthritic process. T cell populations are not homogenous and consist of different subgroups, each with a specific function and structure. In addition to common surface antigens (CD) found in all T lymphocytes (such as CD2, CD3, and CD5), other surface molecules also exist that are used to discriminate between different T lymphocyte subgroups (such as CD61, CCR6, CCR4, CXCR3). CD4+ Th cells are activated by the antigenic stimulation of T cell receptors and differentiated into different subsets of effector Th cells. Among these cells, INF-gamma-producing Th1 cells are predominant in RA. Recent reports have suggested that IL-17-producing Th17 cells are a new subset of cells critical to the pathogenesis of RA. IL-17 induces the production of inflammatory cytokines such as IL-1, IL-6, IL-8, and TNF-α and it has been detected in the serum, SF, and synovium of patients with RA. Beyond Th17 and Th1 cells, there is a third pathogenetic Th cell population in RA, namely Th17/1 cells. This population expresses a cytokine phenotype intermediate between Th17 and Th1. On their surface, Th17/1 cells express CCR6 and CD61, suggesting a common origin of Th17 lymphocytes. However, Th17/1 are CCR4 negative, different from Th17. This intermediate population is CXCR3 and T-bet positive (like Th1 cells) and produce both IL-17 and INF-gamma.

On the other hand, reports about T cell subtypes in OA SM or SF are scarce. In particular, CD4+ T lymphocytes and their cytokines play a major role in the initiation of inflammation in this disease. The objective of this study was to analyze, by means of various CD markers’ expression, the means level and activation degree of T cells particularly Th17 lymphocytes in peripheral blood samples from OA patients and to compare them with samples from patients with RA and healthy controls.

Methods

Patients with established diagnosis of RA according to American College of Rheumatology/European League Against Rheumatism 2010 criteria, knee or hip OA according to American College of Rheumatology criteria, and healthy blood donor volunteers were eligible. Other inclusion criteria were: DAS28 (disease activity score based on 28 joints) between 3.2 and 5.1 or a Western Ontario and McMaster Universities Arthritis Index (WOMAC) score of more than 50 respectively in patients with RA or OA, age between 40 and 60 years old, a Kellgren Lawrence index of 2–3. Patients with RA were naïve to disease-modifying antirheumatic drugs therapy including methotrexate and steroids. Patients with OA were naïve to treatment with nonsteroidal anti-inflammatory drugs and disease-modifying osteoarthritis drugs. Exclusion criteria were the presence of other autoimmune diseases, tumors or secondary OA. Finally, no intra-articular treatment with steroids or hyaluronic acid was allowed from 3 months before enrollment. Multichannel flow cytometry and monoclonal antibodies against CD3, CD4, CD8, CCR6, CD38, CXCR3, and HLA DR were used to distinguish between and evaluate T cells’ subpopulation. Participants were informed about the aim of the project and gave their written consent. A second sample to confirm the results was obtained 1 month later and the same analysis was performed. The Fornaroli Hospital Local Ethics Committee approved the project. All patients signed informed consent to participate in the study. We collected blood samples from February 2014 to November 2014.

Statistical analysis

The Shapiro-Wilk test was used to study the normality of variables. Data are displayed as mean and standard deviation. In view of non-normal distribution, variables were compared using the Mann–Whitney test for two groups and the Kruskal–Wallis test for multiple groups. Values less than 0.05 were considered significant. SPSS version 14 for Windows was used.

Flow cytometry

Freshly drawn ethylenediaminetetraacetic acid blood samples were analyzed by 8-color flow cytometry (FACSCanto II; BD, Franklin Lakes, NJ, USA) with the following conjugated antibody panel: CD45-FITC; CXCR3-PE; CD4-PerCP-Cy5.5; CCR6-PE-Cy7; CD38-Alexa 647; CD8-APC-H7; CD3-V450; HLA DR-V500 at the appropriate concentrations. After 20-minute staining in the dark, 2 mL of ammonium chloride lysing was added for 10 minutes. After centrifugation at 1,500 rpm for 7 minutes, the pellet was re-suspended in 200
microliters of cold phosphate buffered saline and immediately analyzed. At least 50,000 lymphocytes (defined as CD45+++, SSCLow cells) were acquired. The gating strategy included the parallel capture of CD4+/CD3+ and CD8+/CD3+ cells in two separate downstream hierarchies. Each parent subset was then further dissected into functional subpopulations, namely CD4+ T cells as Th1 cells (CD4+ CXCR3+ CCR6−), Th2 cells (CD4+ CXCR3− CCR6−) and Th17 cells (CD4+ CXCR3− CCR6+), respectively according to Maeker et al.21 Both CD4+ and CD8+ cells were divided into quiescent (CD38− HLADR−) or activated elements (CD38+ and/or HLADR+). Functional subset percentages were calculated over the total lymphocyte population and over the parent CD4+ or CD8+ subsets, respectively, and all values were also recorded as absolute levels per microliter on the basis of total lymphocyte count (DXH800; Beckman Coulter, Brea, CA, USA).

Results
We analyzed blood samples of 91 subjects (75 females, 16 males): 15 patients with well-defined RA, 56 with hip or knee OA, and 20 healthy age matched controls. Median age was 43 (range 34–52) years old (P>0.05 between groups). The median duration of disease in OA and RA patients was 3.8 (range 3.1–5.7) and 3.1 (range 2.8–4.5) years respectively (P>0.05). Blood samples from the RA patients had a significantly higher count of CD4+ CD38+ DR+ (activated CD4 T cells) and Th17 (CCR6+ CXCR3−) cells as compared to OA patients and control group (P<0.01). Furthermore the samples from the OA patients had a higher percentage of activated CD4 T cells as compared to control group (P<0.05) (Figures 1–6). Interestingly there was no difference between Th1 (CD4+ CXCR3+ CCR6−), Th2 (CD4+ CXCR3− CCR6−), and Th17 levels between the three groups (P>0.1). Analog results were obtained in the second sample performed 1 month later, with no significant variations between groups (P>0.05).

Discussion
Circulating markers of inflammation, such as CRP, may be elevated in OA compared with control populations without disease.22–24 Histological examination of synovia frequently demonstrates inflammatory cell infiltration, involving macrophages and lymphocytes, increased cell turnover and angiogenesis.25,26 CD4+ Th cells are activated by the antigenic stimulation of T cell receptors and differentiated into different subsets of effector Th cells.18 It is well known that RA has been characterized by a Th1 response. Among these cells, INF-gamma-producing Th1 cells are predominant in RA. However, the etiopathogenetic role in RA of a new Th subset (Th17) has appeared to be very important in the last years. These Th17 cells and Th17/1 are more important than Th1 in the contribution to pathogenesis in early RA.27,28 IL-17 induces the production of inflammatory cytokines such as IL-1, IL-6, IL-8, and TNF-α and it has been detected in the serum, SF, and synovium of patients with RA and OA.19 Previous reports including tissue samples from patients with OA and RA had demonstrated infiltrating mononuclear cells and pro-inflammatory cytokines in both diseases but consistently less abundant in OA. Gallelli et al recently described levels of IL-6, TNF-α, and VEGF in SF of patients with knee OA.29–33 The pro-inflammatory cytokine IL-17 is implicated in both the induction of local inflammation and the joint destruction that characterize the ST of RA. IL-17 is able to stimulate fibroblast-like synoviocytes, macrophages, chondrocytes, and osteoclasts.34,35 IL-17 is present at increased concentrations in serum, SF, and syno-
ovium samples from patients with RA and has been shown to induce and perpetuate chronic inflammation, cartilage damage, and bone erosion. In RA, the number of Th17 cells in peripheral blood correlates with disease activity.\textsuperscript{36} Many studies have demonstrated the Th17 cell heterogeneity, high degree of phenotypic instability and plasticity, responsible for induction of a Th1-like phenotype.\textsuperscript{36–38} Ghoreschi et al demonstrated that Th17 cells can be divided into pathogenic and non-pathogenic phenotype according to their different cytokine secretion. In particular, IL-17, IL-21, IL-22, IL-2, and IFN-gamma are secreted by pathogenic cells (including Th17/1); while non-pathogenic Th17 release IL-17, IL-21, IL-9, and IL-10. According to cytokine milieu, Th17 can induce differentiation in Th17/1 lymphocytes or into Th1 phenotype, in the presence of IL-12 or IL-23 in the absence of TGF-beta in vitro, respectively.\textsuperscript{37–40} Th17/1 cells secrete IL-17 and INF-gamma while Th1 cells do not release IL-17 but only INF-gamma. OA shares several characteristics with RA, including joint destruction and synovitis. It can be assumed that tissue damage in OA is responsible for the chemotaxis of inflammatory cells, while in RA joint destruction is considered to be a consequence of inflammation. In addition, IL-17 has been shown to be present in SF from patients with OA. However, in literature there are few data about Th17/1 in OA, today.\textsuperscript{41} In this report, we showed that patients with OA have a different expression of activated Th cells, with a marked tendency of higher expression than healthy controls. There was no significant qualitative evidence but only quantitative differences between RA and OA Th sub-populations.
thus indicating that lymphocytes in OA and RA exhibited a similar Th phenotype with evident activation and involving the same pro-inflammatory cytokines. In particular, we found the same levels in the OA and RA subgroups of Th1 cells, Th2 cells, and Th17 cells. Also, the levels of CD38+DR+ and DR+ cells were similar in OA and RA subgroups, significantly higher than in the healthy control group. These results indicate that, despite a mechanical contribution to its pathophysiology, OA might be characterized as a low-grade inflammatory condition, similar to other rheumatic diseases with a well-known inflammatory etiology such as RA.

**Disclosure**

The authors have no conflicts of interest to disclose.

**References**


