Follicular Helper T Cells in Peripheral Blood of Patients With Rheumatoid Arthritis

Alicia Beatriz Costantino, a Cristina del Valle Acosta, a Laura Onetti, b Eduardo Mussano, b Ignacio Isaac Cadile, b Paola Virginia Ferrero b

a Laboratorio de Inmunología y Virología, Hospital Nacional de Clínicas, Facultad de Ciencias Médicas, Universidad Nacional de Córdoba, Córdoba, Argentina
b Servicio de Reumatología, Hospital Nacional de Clínicas, Facultad de Ciencias Médicas, Universidad Nacional de Córdoba, Córdoba, Argentina

ARTICLE INFO

Article history:
Received 16 February 2016
Accepted 8 July 2016
Available online 22 September 2017

Keywords:
Rheumatoid arthritis
Immunopathogenesis
Follicular helper T cells

ABSTRACT

Introduction: Rheumatoid arthritis (RA) is a chronic autoimmune disease that is characterized by the presence of different autoantibodies such as rheumatoid factor (RF) and anti-citrullinated protein antibodies. CD4 T cells expressing CXCR5, referred as follicular helper T cells (Tfh), collaborate with B cells to produce antibodies. Differential expression of CXCR3 and CCR6 within CD4+CXCR5+ T cells defines three major subsets: CXCR3+CCR6− (Tfh1), CXCR3−CCR6+ (Tfh2) and CXCR3−CCR6− (Tfh17). The aim of the study was to assess whether there is an association between the percentage of these cells and RA and whether there is a correlation with disease activity.

Material and methods: Twenty-four RA patients, 22 healthy controls (HC) and 16 undifferentiated arthritis (UA) patients were included. Percentage of CD4+CXCR5+ T cells and their subsets were analyzed by flow cytometry.

Results: No differences were found in the percentages of CD4+CXCR5+ T cells in the comparison of RA vs HC or RA vs UA patients. Tfh1, Tfh2 and Tfh17 subsets showed no differences either. There was no correlation between CD4+CXCR5+ T cells, Tfh1, Tfh2 and Tfh17, and Disease Activity Score in 28 joints (DAS28) or erythrocyte sedimentation rate. Surprisingly, there was a positive correlation between Tfh17 cells and C-reactive protein. Finally, there was no correlation between CD4+CXCR5+ T cells, or their subsets, and anti-mutated citrullinated vimentin, or between the cells and RF.

Conclusion: There were no differences between the percentages of CD4+CXCR5+ T cells and their subsets in peripheral blood of RA patients and the percentages of cells in the control groups. This finding does not rule out a pathogenic role of these cells in the development and activity of RA.

© 2016 Elsevier España, S.L.U. and Sociedad Española de Reumatología y Colegio Mexicano de Reumatología. All rights reserved.

Células T helper foliculares en sangre periférica de pacientes con artritis reumatoide

RESUMEN

Introducción: La artritis reumatoide (AR) es una enfermedad autoinmune y crónica caracterizada por la presencia de autoanticuerpos como factor reumatoide (FR) y anticuerpos antiproteínas citrulinadas. Una población de células T helper foliculares (Tfh), que expresan CD4+CXCR5+, colabora con las células B para la producción de anticuerpos. La expresión diferencial de CXCR3 y CCR6 dentro de las células CD4+CXCR5+ define 3 subpoblaciones mayores: CXCR3+CCR6− (Tfh1), CXCR3−CCR6+ (Tfh2) y CXCR3−CCR6− (Tfh17). El objetivo del estudio fue evaluar si existe asociación entre el porcentaje de estas células y la AR, y la correlación de las mismas con actividad de la enfermedad.

Material y métodos: Participaron 24 pacientes con AR, 22 controles saludables (CS) y 16 pacientes con artritis indiferenciada (AI). Los porcentajes de las células CD4+CXCR5+ y sus subpoblaciones fueron analizados por citometría de flujo.

Please cite this article as: Costantino AR, del Valle Acosta C, Onetti L, Mussano E, Cadile II, Ferrero PV. Células T helper foliculares en sangre periférica de pacientes con artritis reumatoidea. Reumatol Clin. 2017;13:338–343.

Corresponding author.
E-mail address: ferreropaola@hotmail.com (P.V. Ferrero).

© 2016 Elsevier España, S.L.U. and Sociedad Española de Reumatología y Colegio Mexicano de Reumatología. All rights reserved.
**Introducción**

La artritis reumatoide (AR) es una enfermedad crónica, inflamatoria, autoinmune que no afecta únicamente a los huesos y cartílagos, sino que está asociada con una serie de síntomas generales y con la muerte. Es caracterizada por la presencia de anticuerpos antifosfolípido, incluyendo antifosfolípido (RF) y anticitrullinado (ACR). Los autores han descrito recientemente una subpoblación de células TCD4⁺ CXCR5⁺, que se presentan en los linfoides dependientes de células B (B-cells), conocidos como células T helper tipo 2 (Tfh) y celular B, responsables de la producción de anticuerpos. Estas células expresan el quimiorreceptor CXCR5, que permite su migración al sitio del linfoides dependientes de células B en respuesta a su ligando CXCL13. La actividad de estas células Tfh puede ser medida a través de marcadores como ICOS. El objetivo de este estudio fue determinar si existían diferencias entre los perfiles de células TCD4⁺ CXCR5⁺ en pacientes con AR y los controles normales.

**Materiales y Métodos**

**Pacientes y Controles**

Se llevó a cabo un estudio de casos-control selectivo. Los pacientes fueron dados por el departamento de reumatología del Hospital Nacional de Clínicas, Universidad Nacional de Córdoba, Argentina, entre marzo de 2014 y febrero de 2015. Se incluyeron 24 pacientes que se diagnostican entre los 18 y 70 años, y se diagnosticaron de acuerdo con el criterio americano de reumatología/European League Against Rheumatism (ACR/EULAR). Se excluyeron 15 pacientes que no habían sido tratados, los que habían sido diagnosticados más de 12 meses antes de participar y los que habían disminuido o disminuido su medicación.

**Resultados**

No hubo diferencias en los porcentajes de células CD4⁺ CXCR5⁺ entre los pacientes con AR y los controles. No se observó correlación entre las células TCD4⁺ CXCR5⁺ y la determinación de actividad DAS28, así como tampoco con la velocidad de sedimentación globular. Sorpresivamente, hubo una correlación positiva entre las células Tfh17 y la proteína C reactiva. Finalmente, no se observó correlación entre las células TCD4⁺ CXCR5⁺ y cualquiera de las subpoblaciones antivirales citrulina utilizadas.

**Conclusión**

No se hallaron diferencias entre los perfiles de células TCD4⁺ CXCR5⁺ y sus subpoblaciones en sangre periférica de los pacientes con AR y las células de los grupos controles. Esto no descarta un papel patogénico de estas células en el desarrollo y actividad de AR.
components C3 (RV 80–180 mg%) and C4 (RV 17–40 mg%) were measured by means of single radial immunodiffusion.

**Determination of CD4⁺CXCR5⁺ T Lymphocytes and Their Distinct Subpopulations by Flow Cytometry**

Peripheral blood mononuclear cells (PBMC) were isolated by density gradient centrifugation using Ficoll-Paque™ Plus (GE Healthcare Bio-Sciences AB, Uppsala, Sweden). For immunostaining, we utilized 1.1⁶ PBMC and the following antibodies: anti-human CD4 fluorescein isothiocyanate (FITC) (clone RPA-T4), anti-human CD196 (CCR6) phycoerythrin (PE) (clone 11A9), anti-human CD183 (CXCR3) allophycocyanin (APC) (clone IC6/CXCR3) and anti-human CXCR5 peridinin chlorophyll protein (PerCP-Cy5.5) (clone RF8B2) (BD Pharmingen™, San José, California, United States). The data were acquired immediately after staining with a FACS Calibur 4-color flow cytometer (Becton-Dickinson, Boston, Massachusetts, United States). In each sample, we analyzed at least 50,000 CD4⁺ cells using the Infinicyt™ software package (version 1.7) (Citognos SL, Salamanca, Spain).

**Statistical Analysis**

The data were analyzed with the MedCalc software package (version 10.2.0.0) (MedCalc Software, Ostend, Belgium). The normal distribution of the data was assessed with the Kolmogorov–Smirnov test. For the variables with a normal distribution or whose logarithmic transformation had a normal distribution, we used parametric tests like one-way analysis of variance (ANOVA) and the Bonferroni correction, as well as the Pearson correlation coefficient. For the remaining variables, we employed nonparametric methods, such as the Kruskal–Wallis and Dunn tests, and the Spearman correlation coefficient. The categorical data were compared with the Chi-square test. A P value <.05 was considered to indicate statistical significance.

**Results**

**Characteristics of the study groups:** Table 1 shows the demographic and clinical characteristics of the study groups. The ages and sex distribution were homogeneous. As was expected, the ESR and CRP levels were higher in the patients with RA when compared to the HC. In all, 79% of the RA patients were positive for RF, whereas none of the controls had a positive test. Likewise, the anti-MCV autoantibody levels were significantly higher in RA patients when compared to the other 2 groups. The IgM levels were higher in patients with RA when compared to those of the group of HC, and those of IgA in the comparison with patients with UA. The levels of complement C4 were lower in RA patients when compared to those with UA, but remained in normal range in both groups. Most of the patients with RA had high disease activity (Table 2). Two patients, 1 of whom was in remission, had a low activity (DAS28 <2.6).²⁶

**Percentage of CD4⁺CXCR5⁺ T cells:** To identify the distinct subpopulations of these cells by flow cytometry, we applied the strategy detailed in Fig. 1. We compared the percentage of CD4⁺CXCR5⁺ T cells in the 3 groups, and found no significant differences (RA: 12.89% ± 7.73%, HC: 10.48% ± 3.90%, UA: 11.71% ± 5.04%; P = .66), as shown in Fig. 2. Likewise, there were no differences between the groups when we considered only the patients with high activity (DAS28 >5.1, n = 14; 12.84% ± 8%; P = .73).

**CD4⁺CXCR5⁺ T cell subpopulations:** We compared the percentages of the distinct subpopulations of CXCR3⁺CCR6⁻ (Tfh1), CXCR3⁻CCR6⁻ (Tfh2) and CXCR3⁻CCR6⁺ (Tfh17) cells within the CD4⁺CXCR5⁺ population (Fig. 3). We observed no differences in the Tfh1 subpopulation (RA = 12.75% ± 9.72%, HC = 11.21% ± 7.48%, UA = 12.81% ± 6.13%; P = .77), or in the Tfh17 subpopulation (RA = 37.94% ± 11.34%, HC = 40.79% ± 8.17%, UA = 37.34% ± 7.16%; P = .45). The difference observed in the Tfh2 subpopulation (RA = 32.66% ± 11.46%, HC = 39.53% ± 12.12%, UA = 27.56% ± 11.25%; P = .0092) was due to the lower percentage found in the UA group when it was compared to the HC (P < .01). No differences were observed when it was compared only with those patients who had high disease activity (n = 14) and the control groups (Tfh1 in RA was 12.74% ± 9.19%, P = .77; Tfh2 in RA was 35.03% ± 11.64%, P = .01; Tfh17 in RA was 36.21% ± 10.80%, P = .26), again, and as was expected, the difference in Tfh2 was due to the results on comparing UA vs HC (P < .01).

**Correlation of the study populations and disease activity:** We analyzed the correlation between the CD4⁺CXCR5⁺ population and its distinct subpopulations with the DAS28. We found no correlation between the CD4⁺CXCR5⁺ cells and the DAS28 (r = -.19, P = .37), or between the Tfh1, Tfh2 and Tfh17 subpopulations and the activity

**Table 1**

Summary of the Characteristics of the Controls and Patients Participating in the Study.

<table>
<thead>
<tr>
<th></th>
<th>RA (n = 24)</th>
<th>HC (n = 22)</th>
<th>UA (n = 16)</th>
<th>RA vs HC, P value</th>
<th>RA vs UA, P value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Sex, F/M</td>
<td>21/3</td>
<td>19/3</td>
<td>13/3</td>
<td>.77e</td>
<td>.93e</td>
</tr>
<tr>
<td>Age, years</td>
<td>51 ± 10</td>
<td>49 ± 10</td>
<td>52 ± 11</td>
<td>&gt;.05e</td>
<td>&gt;.05e</td>
</tr>
<tr>
<td>WBC, 10⁹/L</td>
<td>6.98 ± 1.85</td>
<td>7.23 ± 1.96</td>
<td>6.84 ± 1.74</td>
<td>&gt;.05e</td>
<td>&gt;.05e</td>
</tr>
<tr>
<td>Hb, g/L</td>
<td>12.6 ± 1.91</td>
<td>12.6 ± 0.98</td>
<td>12.96 ± 1.07</td>
<td>&gt;.05e</td>
<td>&gt;.05e</td>
</tr>
<tr>
<td>PLT, 10⁹/L</td>
<td>255 ± 80</td>
<td>254 ± 41</td>
<td>249 ± 59</td>
<td>&gt;.05e</td>
<td>&gt;.05e</td>
</tr>
<tr>
<td>ESR, mm/h</td>
<td>25 ± 21</td>
<td>9 ± 6</td>
<td>14 ± 12</td>
<td>&lt;.01e</td>
<td>&gt;.05e</td>
</tr>
<tr>
<td>CRP, mg/L</td>
<td>18 ± 14</td>
<td>9 ± 2</td>
<td>14 ± 8</td>
<td>&lt;.01e</td>
<td>&gt;.05e</td>
</tr>
<tr>
<td>RF, %</td>
<td>19 (79)</td>
<td>0 (0)</td>
<td>0 (0)</td>
<td>&lt;.001f</td>
<td>&lt;.001f</td>
</tr>
<tr>
<td>MCV, fL/μL</td>
<td>750 (77–530.0)</td>
<td>2.8 (2.3–5.3)</td>
<td>2.9 (2.5–3.8)</td>
<td>&lt;.001e</td>
<td>&lt;.001e</td>
</tr>
<tr>
<td>IgG, mg/L</td>
<td>1310 ± 355</td>
<td>1325 ± 257</td>
<td>1212 ± 336</td>
<td>&gt;.05e</td>
<td>&gt;.05e</td>
</tr>
<tr>
<td>IgM, mg/L</td>
<td>220 ± 88</td>
<td>169 ± 55</td>
<td>179 ± 60</td>
<td>&lt;.05e</td>
<td>&gt;.05e</td>
</tr>
<tr>
<td>IgA, mg/L</td>
<td>363 ± 126</td>
<td>313 ± 80</td>
<td>261 ± 114</td>
<td>&gt;.05e</td>
<td>&gt;.05e</td>
</tr>
<tr>
<td>C3, mg%</td>
<td>112 ± 33</td>
<td>119 ± 26</td>
<td>130 ± 30</td>
<td>&gt;.05e</td>
<td>&gt;.05e</td>
</tr>
<tr>
<td>C4, mg%</td>
<td>24 ± 9</td>
<td>26 ± 7</td>
<td>31 ± 8</td>
<td>&gt;.05e</td>
<td>&gt;.05e</td>
</tr>
<tr>
<td>DAS28</td>
<td>5.16 ± 1.35</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
</tr>
</tbody>
</table>

Anti-MCV, anti-mutated citrullinated vimentin antibodies; CRP, C-reactive protein; DAS28, Disease Activity Score in 28 joints; ESR, erythrocytes sedimentation rate; F, female; Hb, hemoglobin; HC, healthy controls; Ig, immunoglobulin; M, male; PLT, platelets; RA, rheumatoid arthritis; RF, rheumatoid factor; UA, undifferentiated arthritis; WBC, white blood cells.

1. Value expressed as mean ± standard deviation.
2. Value expressed as median and interquartile range (25th–75th percentile).
3. Chi-square test.
4. One-way analysis of variance (Bonferroni post-test).

Statistically significant P values (P < .05) are in boldface.
We detected no correlation between the CD4⁺CXCR5⁺ population and ESR or CRP (r = 0.18, P = .39 and r = 0.27, P = .20, respectively). We found no correlation between Th1, Th2 and Th17 and ESR (r = 0.080, P = .71; r = −0.0094, P = .97 and r = −0.25, P = .23, respectively), or between Th1 and CRP (r = 0.27, P = .20) or Th2 and CRP (r = 0.14, P = .51). Surprisingly, there was a positive correlation between Th17 and CRP (r = 0.47, P = .021) (Fig. 4).

**Discussion**

Although a number of immunopathogenic mechanisms have been proposed, the cause of RA is still unknown. As a consequence, the discovery of both the elements of the immune system that participate in its onset and the mechanisms involved is of great value for the development of more effective therapies. In our investigation, we studied a subpopulation of CD4⁺ T cells, Th cells, which have a very important role in the collaboration for the production of antibodies by the B cells. However, we found no significant differences in the percentage of CD4⁺CXCR5⁺ T cells in the blood of RA patients when compared with that of the HC or of patients with UA. The same can be said when we considered only patients with RA and high disease activity. The literature provides relatively variable results that may be associated with the manner of characterizing this population in peripheral blood. Our findings agree with those reported by Chakera et al., who found no differences when they compared Th cells such as CD4⁺CXCR5⁺, CD4⁺CD45RA⁻CXCR5⁺, CD4⁺CXCR5⁺ICOS⁺ and CD4⁺CXCR5⁺PD-1. The authors of another study observed similar percentages of CD4⁺CXCR5⁺CD45RA⁻ and CD4⁺CXCR5⁺CD45RA⁻CCR7⁺PD-1 high cells. Likewise, according to an article by Arroyo-Villa et al., no differences were found when CD4 and CXCR5 were employed to characterize them. However, when ICOS was added to the

| Table 2 Distribution of Disease Activity in Patients With Rheumatoid Arthritis. |
|-------------------------------------|-----------------|
| **DAS28**                           | **RA (n/%)**    |
| Low activity (<3.2)                 | 2/8.4          |
| Moderate activity (≥3.2 <5.1)       | 8/33.3         |
| Low activity (≥5.1)                 | 14/58.3        |

DAS28, Disease Activity Score in 28 joints; RA, rheumatoid arthritis.

index employed (Th1 and DAS28 r = 0.09, P = .68; Th2 and DAS28 r = −0.36, P = .09; Th17 and DAS28 r = −0.20, P = .35). When we took into account only those patients with high disease activity (n = 14), we observed no correlation between any of the subpopulations being studied and the DAS28 (CD4⁺CXCR5⁺ and DAS28 r = −0.32, P = .26; Th1 and DAS28 r = −0.17, P = .56; Th2 and DAS28 r = 0.33, P = .25; Th17 and DAS28 r = 0.024, P = .94).

**Correlation between Tfh cells and the different autoantibodies in patients with RA:** Given that Tfh cells collaborate with B cells for the production of antibodies, we proposed to determine whether CD4⁺CXCR5⁺ T cells and their distinct subpopulations could be related to the production of autoantibodies in RA patients. We found no correlation between the percentages of CD4⁺CXCR5⁺ T cells (r = 0.38, P = .066), or between the 3 subpopulations (Th1 r = −0.04, P = .84; Th2 r = −0.14, P = .51; Th17 r = −0.19, P = .37) and anti-MCV antibodies. The same result was observed when the percentages of these cell populations were correlated with RF (CD4⁺CXCR5⁺: r = 0.30, P = .15; Th1 r = −0.18, P = .38; Th2 r = −0.15, P = .46; Th17 r = 0.0051, P = .98).

**Correlation between Tfh cells and the distinct cell subpopulations with inflammation markers:** To establish whether Tfh cells and the different subpopulations had a role in the inflammatory process that occurs in RA patients, we correlated them with ESR and CRP.

![Fig. 1. Two-dimensional contour density plot showing the gating strategy in a representative experiment in a patient with rheumatic arthritis. FSC, forward scatter; SSC, side scatter.](image-url)
original labeling, a greater number of CD4+CXCR5+ICOS+ cells was observed in RA patients.21 Similarly, in contrast to our findings, the literature includes other studies that report an increase in the number of Tfh cells in RA patients after the addition of ICOS as a third cell marker, as occurred even with PD-1 when associated with the abovementioned markers.19,22,23 Previous studies of CD4+CXCR5+ T cells from mouse germinal centers identified a population that expresses the master regulators Bcl6 and Foxp3, with the capacity to prevent antibody-mediated autoimmunity, referred to as follicular regulatory T cells (TFg).24 Recently, this population was also reported in human blood and tonsils.25,26 As the Tfh and TFg cells fulfill opposing functions and share the expression of a number of markers, such as CXCR5, ICOS and PD-1, the addition of the use of Foxp3 as a marker would be crucial to be able to discriminate them. This would help to reconcile apparently contradictory findings in relation to the consequences of numerical changes in these populations observed in the course of autoimmune processes.

To define the Tfh cell subpopulations, we employed the combination of markers CCR6 and CXCR3, in accordance with the criteria employed by Morita et al. in 2011.10 That group of researchers and another team found that subpopulations Tfh2 and Tfh17 were effective collaborators of B cells for the induction of the synthesis of distinct immunoglobulin isotypes.10,21 As RA is a disease involving autoantibody production, we expected to observe an increase in Tfh2 and Tfh17 subpopulations; however, we detected no variation in either of the two, or in the Tfh1 subpopulation. In this respect, the literature also reports contradictory results in the study of patients with RA. The group of Arroyo-Villa et al. found a difference in the number of Tfh17 cells, but not in that of Tfh2,21 whereas another group reported no changes among any of the subpopulations in RA patients.20

In the study of the comparisons of the DAS28, CRP, ESR, RF and anti-MCV antibodies in the distinct populations, we detected no correlation in any case, except between Tfh17 and CRP. Similar to what occurred in this study, in the literature there are no previous reports of a correlation between Tfh cells and parameters of disease activity like the DAS28, CRP and ESR.22 In contrast, other groups observed a positive correlation between Tfh and the DAS2819,22 or between Tfh and anti-cyclic citrullinated peptide antibodies19,22 and Tfh and RF.19 With regard to the correlation between Tfh17 and CRP, an earlier article demonstrated that the Tfh17 subpopulation produces IL-21, IL-17A and IL-22.13 These cytokines could indirectly promote the production of CRP as they stimulate hepatocytes, keratinocytes and epithelial cells to produce IL-1β and IL-6,14 that act in the liver to induce CRP synthesis,3 which could be the mechanism employed in RA.

In conclusion, we found no numerical differences between Tfh cells and the distinct subpopulations in peripheral blood or any correlation with parameters of disease activity and inflammatory markers. Nevertheless, we cannot rule out the possibility that they play a pathogenic role in RA. This population could be recruited to the site of inflammation, the joint in this case, and collaborate with B cells for the production of autoantibodies in ectopic germinal centers, located in the synovial membrane of RA patients.

A clear phenotypic description of memory Tfh cell subgroups in the blood is important, not only to help understand their biological functions, but also for translational purposes, as these circulating cells could serve as potential biomarkers to monitor the deregulated responses of antibodies in autoimmune diseases like RA.
Ethical Disclosures

Protection of human and animal subjects. The authors declare that the procedures followed were in accordance with the regulations of the relevant clinical research ethics committee and with those of the Code of Ethics of the World Medical Association (Declaration of Helsinki).

Confidentiality of data. The authors declare that they have followed the protocols of their work center on the publication of patient data.

Right to privacy and informed consent. The authors have obtained the written informed consent of the patients or subjects mentioned in the article. The corresponding author is in possession of this document.

Funding

The funds required to perform the work for this report were provided by the Sociedad Argentina de Reumatología.

Conflicts of Interest

The authors declare they have no conflicts of interest.

Acknowledgments

We wish to thank biochemists Cecilia María Rodríguez and Melina Croquell for collaborating in the analysis of the flow cytometry samples.

References