

Exaggerated circulating Th-1 cytokine response in early rheumatoid arthritis patients with nodules

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Abstract

Background: Immunohistochemical studies of the rheumatoid nodule (RN) suggest it is a Th1 granuloma, with focal vasculitis, yet the pathogenesis remains unclear and little is known about circulating cytokines in these patients. **Objective:** We studied circulating cytokines in DMARD-naïve RA patients to investigate associations with subcutaneous RN. **Methods:** 149 DMARD-naïve adults with early RA (symptom duration ≤ 2 years) were assessed using the Simplified Disease Activity Index (SDAI), and hand and feet radiographs were scored using the modified Larsen method. Circulating cytokines and growth factors representative of T- helper cell 1(Th1) and Th2 cell, macrophages, and fibroblasts were measured using the Bio-Plex® suspension array system. **Results:** Of 149 patients, 34 (22.8%) had subcutaneous RN, and these patients had more severe disease with higher mean swollen joint counts ($p=0.02$), SDAI ($p=0.04$) and modified Larsen scores ($p=0.004$). There were no differences in Rheumatoid Factor or anti-cyclic citrullinated peptide antibody positivity between patients with RN and those without RN. Patients with RN showed significantly higher levels of circulating IL-12 ($p=0.02$), IL-2 ($p=0.048$), and VEGF ($p=0.033$) levels, with a trend towards higher levels of IL-7 ($p=0.056$), IFN- γ ($p=0.059$) and IL-8 ($p=0.074$) compared to those without RN. **Conclusions:** DMARD-naïve early RA patients with RN had more severe disease than those without RN, and showed an exaggerated circulating Th1 and macrophage cytokine profile.

Keywords: Rheumatoid nodules, extra-articular disease, circulating cytokines, VEGF

Rheumatoid nodules (RN) are the most common extra-articular feature of rheumatoid arthritis (RA), and are found most commonly in subcutaneous tissues at the site of recurrent mechanical irritation. Nodules are a feature of established RA, occurring in 30-40% of patients, with an average disease duration of eleven years (1). Joint inflammation and RN do not run the same clinical course: patients with low disease activity can still develop RN, and an increase in nodule size or formation of new nodules is well described with therapy that suppresses synovitis, such as methotrexate and anti-TNF drugs.

The RN has been described as “the most characteristic histopathological lesion in RA” and as such perhaps deserves more attention(2). Histopathologically, the RN is an immune granuloma consisting of a central area of necrosis, surrounded by a palisade of macrophages and fibroblasts, and a peripheral vascular area containing T lymphocytes and macrophages. A number of studies have drawn comparisons between RN and rheumatoid synovium, describing similarities between the tissues with respect to cellular components, immunohistochemical features and cytokine production(3, 4). There are also important differences including the absence of B lymphocytes and lymphoid follicles in the nodule, variations in expression of cell adhesion molecules and in cytokine gene expression(5).

There is increasing evidence of endothelial dysfunction and angiogenesis in rheumatoid synovium. One marker of angiogenesis is VEGF, and levels correlate with disease activity and radiographic progression(6). A recent study demonstrated higher VEGF levels in patients with extra-articular involvement and the authors concluded that VEGF may be a marker systemic of rather than joint inflammation in RA(7). Recently, IL-8 has been shown to play an important role in angiogenesis(8).

Better understanding of the cytokines present in serum of patients with RN might offer clues to the immunopathogenesis of RN and possibly other extra-articular complications of RA. To our knowledge there are no published studies on the relationship of circulating cytokines with RN. We therefore undertook a cross-sectional study of early RA patients who were disease modifying anti-rheumatic drug (DMARD)-naïve to determine the association of RN with disease activity, radiographic damage,

autoantibody status, the presence of the shared epitope (SE), and circulating cytokines. This study was approved by the University of the Witwatersrand Committee for Research on Human Subjects.

Patients and Methods

1.1 Patients

149 early RA (defined as disease duration ≤ 2 years) adults who met the American College of Rheumatology classification criteria for RA and were DMARD naïve were studied. Clinical assessments (Table 1) included a smoking history, the presence of subcutaneous RN, the Simplified Disease Activity Index (SDAI) and the health assessment questionnaire disability index (HAQ-DI). A chest x-ray (CXR) was performed and radiographs of hands and feet were scored using the modified Larsen score.

1.2 Auto-antibody and cytokine assays

Venous blood (30ml) was collected in endotoxin-free, silicone-coated vacutainers and allowed to coagulate at room temperature, followed by centrifugation (3000 rpm for 10 minutes). The serum was removed and aliquoted, and stored at minus 20°C until performance of the various assays described below. Rheumatoid factor (RF) was assayed by nephelometry (Siemens Healthcare Diagnostics, BN Prospec Nephelometer, Newark, USA) and anticyclic citrullinated peptide (aCCP) antibodies were measured by a second generation immunofluorometric procedure using the Immucap 250 system (Phadia AB, Uppsala, Sweden). Values of 15 IU/ml and 10 U/ml respectively were considered positive. Serum IL-1b, IL-1Ra, IL-2, IL-4, IL-6, IL-7, IL-8, IL-10, IL-12, IFN- γ , TNF, G-CSF, GM-CSF, CCL2, CCL4, and VEGF were measured using the Bio-Plex® suspension array system (Bio-Rad Laboratories Inc, Hercules, CA, USA) which utilizes Luminex® xMAP™ multiplex technology. The system uses an array of beads in liquid suspension, each containing different ratios of two spectrally distinct fluorophores, thereby assigning a unique spectral identity. The beads are conjugated with a monoclonal antibody specific for a target protein and incubated with the serum sample ($\frac{1}{4}$ dilutions), washed, followed by addition of a biotinylated detection antibody, washed

again, and finally incubated with streptavidin-phycoerythrin. A wide range of standards (0.38 – 91756.00 pg/ml) were used to enable quantitation of the individual cytokines using a Bio-Plex array reader with a dual laser detector and real time digital signal processing.

1.3 Assessment for the Shared Epitope

Genomic DNA was analysed for the presence of the SE using high-resolution rSSO PCR typing of the RAA amino acid motif at positions 72 to 74 of the third hypervariable region of the different human leucocyte antigen HLA-DRB1 allele, as described previously(9).

1.4 Statistical analysis

The Mann-Whitney test was applied to compare continuous variables, and in the case of categorical variables, the Pearson's Chi-Square test, or where indicated, the 2-tailed Fishers' Exact test was used. Log transformation was performed on cytokine levels to correct for non-normal distribution. Because the higher mean SDAI in the RN group compared to the non-RN group may explain the differences in cytokine levels, we further analysed our data adjusting for SDAI. As our dependent variable was binary (RN vs non- RN) we employed multiple logistic regression to determine how each of the cytokines changed when we introduced SDAI in the model. Stata 10 software (StataCorp, USA) was used. A p-value of <0.05 was considered significant, and p-values of <0.1 were considered to show a "trend" towards significance.

Results

2.1 Clinical features

The majority of patients were female (81%), Black Africans (93%) with mean symptom duration of 11.6 months. A high percentage of patients were RF and aCCP antibody positive (89 and 85% respectively), and the majority (93%) carried the SE.

Subcutaneous RN were observed in 34 patients (22.8%), and all of these occurred on the extensor surface of the forearm. No patients had nodules on CXR. As shown in Table 1, patients with RN had features of more severe disease with higher mean swollen joint counts, SDAI and modified Larsen scores compared to those without RN. A higher percentage of males were found in the RN group but this did not reach statistical significance, and smokers were no more likely to have RN than non-smokers.

Table 1: Clinical, radiographic and autoantibody features of 149 early RA patients with and without rheumatoid nodules (RN)

	RN (n=34)	no RN (n=115)	p value
Black Africans (%)	32 (94.1)	107 (93.0)	ns
Male (%)	10 (29.4)	19 (16.5)	ns
Age (years)- mean (SD)	46.2 (13.9)	46.8 (12.4)	ns
Duration (months)- mean (SD)	13.0 (7.67)	11.2 (7.0)	ns
Ever smoked (%)	7 (20.6)	25 (21.7)	ns
Swollen joint count - mean (SD)	16.1 (7.6)	12.5 (7.7)	0.02
CRP (mg/l)- mean (SD)	40.7 (38.6)	27.5 (37.5)	0.07
Simplified Disease Activity Index - mean (SD)	44.7 (14.8)	38.5 (16.5)	0.04
Health Assessment Questionnaire- mean (SD)	1.9 (0.8)	1.7 (0.7)	ns
modified Larsen Score- mean (SD)	28.3 (17.0)	20.7 (10.7)	0.004
Erosive disease (%)	20 (58.8)	59 (51.3)	ns
RF positive (%)	30 (88.2)	101 (87.8)	ns
aCCP positive (%)	28 (82.4)	87 (75.7)	ns

SD: standard deviation; **RF:** Rheumatoid Factor; **aCCP:** anticyclic citrullinated peptide; **CRP:** C reactive protein; **ns:** not significant

Table 2: Significant differences in cytokine and growth factor levels of RA patients with rheumatoid nodules (RN) compared to those without nodules

	Median (range)		Median (range) for log-transformed		p value*	p value adjusted for SDAI
	RN n = 34	no RN n=115	RN n = 34	no RN n=115		
IL-1b	7.6 (0.1-191.3)	6.2 (1.0-242.3)	2.0 (-2.3-5.3)	1.8 (0.1-5.5)	ns	ns
IL-1Ra	91.6 (-.3308.7)	89.6 (0-4838.2)	5.1 (3.2-8.1)	4.7 (1.1-8.5)	ns	ns
IL-2	6.5 (0.0-515.8)	0.0 (0.0-1319.9)	1.9(-6.9 – 6.2)	-6.9(-6.9 – 7.2)	0.04	0.048
IL-4	89.4 (0.0-4996.0)	59.2 (0.0-6825.1)	4.5(-6.9 – 8.5)	3.0(-6.9 – 8.8)	ns	ns
IL-6	37.7 (0-324.6)	26.1 (0-1077.7)	37 (-13-58)	3.3 (0.9-7.0)	ns	ns
IL-7	89.4 (0.0-4996.0)	19.2 (0.0-6825.1)	4.5(-6.9 – 8.5)	3.0(-6.9 – 8.8)	0.002	0.056
IL-8	12.7 (2.2-132.4)	8.7 (0.0-572.6)	2.5(0.8 – 4.9)	2.2(-6.9 – 6.4)	0.038	0.074
IL-10	13.2 (0-472.1)	13.0 (0-1172.1)	2.6 (-0.4-6.2)	2.6 (0.3-7.1)	ns	ns
IL-12	23.5 (1.3-3557.8)	11.7 (0.6-3104.6)	3.2(0.3 – 8.2)	2.5(-0.5 – 8.0)	0.02	0.016
IL-17	0 (0-228.7)	0 (0-38.6)	2.1 (-1.1-5.4)	2.0 (-0.2-37)	ns	ns
IFN- γ	110.2 (0.00-11151.7)	56.4 (0.0-10922.2)	7.3(1.2-9.3)	4.7(-6.9 – 9.3)	0.25	0.059
TNF	19.4 (29-1364.6)	4.2 (0.6-2951.7)	3.0 (1.1-7.2)	2.7 (-0.6-8.0)	ns	ns
G-CSF	17.0 (0-4817.4)	15.5 (0-8764.2)	4.7 (0-3726.2)	2.9 (-1.7-9.1)	ns	ns
GM-CSF	3.4 (0-1363.6)	1.1 (0-3726.2)	4.4 (0-1363.6)	3.6 (-2.3-8.2)	ns	ns
VEGF	407.9 (15.0-2886.4)	128.0(0.0-4503.2)	6.0(2.7 – 7.9)	4.9(-6.9 – 8.4)	0.011	0.033
CCL2	64.2 (0-237.3)	53.3 (0-417.3)	4.5 (1.6-5.5)	4.2 (1.4-6.0)	ns	ns
CCL4	129.2 (31.9-301.8)	107.6 (19.8-643.5)	4.9 (3.5-5.7)	4.7 (2.3-6.5)	ns	ns
IL12:IL-4 ratio	4.2 (0.8-123.7)	9.5 (0.6-2503.7)			0.08	0.15
IL12:IL10 ratio	3.2 (0.4-25.2)	2.6 (0.2-17.3)			0.036	0.051

Data are presented as pg/ml; **SDAI:** simplified disease activity index; **ns:** not significant

*p value for log transformed cytokine levels

2.2 Circulating cytokines

Compared to patients without RN, patients with RN had significantly higher T-helper cell 1(Th1) cytokine levels. As shown in Table 2, IL-12 and IL-2 were higher amongst RN patients even after correction for the SDAI, and there was a trend towards higher levels of IL-7 and IFN- γ . The Th1/Th2 cytokine ratio of IL-12 to IL-10, but not IL-12 to IL-4, was significantly higher in the RN group. In addition, VEGF levels were significantly elevated in the RN group, with higher levels of IL-8 that did not reach significance after correction for the SDAI. There were no significant differences between the two groups in the levels of IL-1 β and TNF, Th2 derived cytokines, nor any other cytokine or growth factor measured.

The occurrence of RF and aCCP antibodies was similar in the RN and the non-RN patients. There were no significant differences in the frequency of the SE between the RN and non-RN patients (56.0% vs 54.7% were homozygous for the SE allele, and 32.0% vs 38.9% heterozygous).

Discussion

In this study of early RA patients, we observed increased serum levels of Th1 and macrophage derived cytokines in the subgroup of patients with RN, independent of disease activity. These findings are consistent with immunohistochemical studies of RN describing a lymphocytic and macrophage infiltrate, with cytokines IFN- γ , IL-2, TNF and IL-1 β extracted from RN (4). Furthermore, levels of VEGF and IL-8 were significantly higher in patients with RN. Elevation of these pro-angiogenic factors might imply that angiogenesis is a prominent feature in RN, in keeping with early and recent descriptions of vascular proliferation in RN (10, 11). However, it could be argued that the circulating VEGF and IL-8 levels are a marker of the overall increased angiogenesis in active RA synovitis (12), rather than indirect evidence of angiogenesis in RN.

The pathogenesis of the RN remains unclear, but local vasculitis has been suggested(13), and our findings supporting a T-cell driven process are compatible with

this. Possibilities include sensitization of vascular endothelium by circulating cytokines such as TNF and IFN- γ , as well as CRP. This may result in hyper-responsiveness to minor mechanical trauma to pressure points favouring adhesion of monocytes, localized activation and differentiation of these cells, generation of T cell chemoattractants and activation of these cells by monocyte/macrophage/fibroblast cytokines (IL-7, IL-12) in either the absence or presence of citrullinated autoantigens. Citrullinated proteins have been identified in RN, raising the question of the role of aCCP antibodies in the pathogenesis of the RN(14).

In this study, circulating IL-7 levels were significantly elevated amongst RN patients. Previously, we and others have shown RA patients to have higher IL-7 levels compared to healthy controls(15, 16) , although reduced IL-7 levels in RA patients compared to healthy controls or osteoarthritis patients have been reported elsewhere(17).

The black South African RA population has a high frequency of the SE allele, and this is borne out in this study where 93% of patients carried the SE. We have previously shown an association between the SE and aCCP antibodies, with higher disease activity scores and higher levels of circulating cytokines occurring in these patients(9), perhaps suggesting a predilection to more severe disease. However, in the present study the SE was not a risk for RN. This is in keeping with a large meta-analysis that shows no significant association between SE and nodules(1). Others have reported an association between the SE and RN, and in particular the SE-containing alleles DRB1*0401(18). The low frequency of the SE and of HLA DRB1*0401 in Afro-Americans and Africans has been suggested as an explanation of the low incidence of RN in these populations(19), but seemingly this is not the case in Southern Africans.

Nearly a quarter of patients in this cohort had RN, and this is higher than the 10% prevalence described elsewhere in patients with early disease(20). As in other studies, RN was associated with higher disease activity and radiological damage scores(20). Males have a higher risk of RN(1, 21), however in the present study although more males than females had RN, this did not reach statistical significance. Smoking was not

associated with RN in the present study, whereas others have reported a strong association(21). The small size of our cohort is one possible explanation.

The chief limitation of this study was that we did not study the relationship between circulating cytokines and the immunohistochemistry of RN. In addition, we were unable to assess the IL-23/IL-17 pathway, as neither IL-21 nor IL-23 was available on the cytokine bead array system at the time of the study. A major strength of this study is that all patients had early disease and were DMARD-naïve; hence therapy was not a confounder in cytokine production or RN formation.

In conclusion, DMARD-naïve RA patients with RN had severe disease, with an exaggerated Th1 and macrophage cytokine profile. Further work is needed to better define the role of these cytokines and IL-17 pathway in the pathogenesis of RN by investigating their relationship with histopathology and immunohistochemistry changes of RN.

Conflict of interest statement

The authors declare no conflicts of interest.

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