Cytokine profile in relapsing-remitting multiple sclerosis patients and the association between progression and activity of the disease

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Abstract. Multiple sclerosis (MS) is a progressive immunemediated disease caused by demyelination of the central nervous system. Cytokines and their receptors have an important role in the evolution of MS lesions, and pro- and anti-inflammatory cytokine levels have been found to correlate with changes in MS disease activity. The aims of the present study were to evaluate the pro-inflammatory [tumor necrosis factor (TNF)- α and interleukin (IL) -1 β , -6 and -12], T helper (Th) 1 [interferon (IFN)-y], Th17 (IL-17) and Th2 (IL-4 and -10) cytokine serum levels in relapsing-remitting (RR)-MS patients and to evaluate the association between the cytokine profile and the progression and activity of the disease. Serum cytokine levels were assessed using enzyme linked-immunosorbent assays in 169 RR-MS patients in the remission clinical phase and 132 healthy individuals who were age-, gender-, ethnicity- and body mass index-matched. Disability and activity of the disease were evaluated using the Expanded Disability Status Scale and magnetic resonance imaging with gadolinium, respectively. IFN-γ and IL-6, -12 and -4 levels were higher in RR-MS patients compared to controls (P=0.0009, 0.0114, 0.0297 and 0.0004, respectively). IL-1 levels were higher in controls compared with RR-MS patients. IL-4 levels were higher in RR-MS patients with mild disability compared to those with moderate and severe disability (P=0.0375). TNF-α and IL-10 levels were higher in

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RR-MS patients with inactive disease compared with those with active disease. IL-17 levels showed a trend towards being higher in RR-MS patients with inactive disease compared to those with active disease (P=0.0631). Low TNF- α and high IFN- γ levels were independently associated with RR-MS (P=0.0078 and 0.0056, respectively) and also with the activity of the disease (P=0.0348 and 0.0133, respectively). Results indicated that RR-MS patients, even in the remission clinical phase, exhibit a complex system of inflammatory and anti-inflammatory cytokines that may interact to modulate the progression and activity of the disease.

Introduction

Multiple sclerosis (MS) is a progressive immune-mediated disease caused by demyelination of the central nervous system (CNS), with perivascular infiltrates of autoreactive T-cells that recognize and react against autoantigens, including myelin basic protein (MBP), proteolipid protein (PLP), myelin oligodendrocytic glycoprotein (MOG) and myelin-associated glycoprotein (1,2). Cytokines and their receptors are important in the evolution of MS lesions, and pro- and anti-inflammatory cytokine levels have been found to correlate with changes in MS disease activity (3).

MS is considered a T helper (Th) 1 lymphocyte-mediated disorder as levels of Th1-related molecules are increased in MS, however, the causes of the disease remain unclear. Th1 lymphocyte cells producing interferon (IFN)-γ and tumor necrosis factor (TNF)-α have been found in the perivascular infiltrates of MS patients (4). IFN-γ and interleukin (IL)-12 are increased in brain, cerebrospinal fluid (CSF) or peripheral blood of MS patients, particularly during acute exacerbations (5). By contrast, Th2 lymphocyte-related molecules, including IL-10 and -4, which downregulate Th1 cells and exert anti-inflammatory functions, are increased during the remission phases of MS (6). Th17, a subset of T cells that secrete IL-17, has been associated with the pathogenesis of

several inflammatory diseases, including MS and encephalomyelitis autoimmune experimental (EAE) (7,8). In EAE, Th1 and Th17 cells have been shown to contribute to disease pathogenesis, however, no absolute requirement for IL-17 or IFN- γ alone has been reported (9).

Cytokines have been identified as major regulators of the immune system and attempts have been made to correlate cytokine levels with MS disease activity, assessed using magnetic resonance imaging (MRI). There is evidence that acute dysregulation of the balance of cytokines is one of the key factors during acute relapse, leading to acute inflammatory lesions in MS patients. Elevated serum and CSF TNF- α and - β levels have been associated with onset of MS relapse (10). Disease progression in MS patients has also been correlated with high CSF TNF- α levels (11).

The clinical disability of MS patients is evaluated using the Expanded Disability Status Scale (EDSS) (12), while the disease activity is evaluated using MRI with gadolinium (Gd)-enhancing lesions, providing the most objective and sensitive tool for assessing the progression and activity of the disease in MS patients. MRI revealed disease activity 5-10 times more frequently than clinical measurements (13). Positive Gd-enhancing lesions are the earliest event in the formation of new MS lesions and are predictive of the course of disability in MS patients (14).

Cytokine changes have been investigated in MS patients from various populations worldwide. The majority of the studies enrolled a limited number of MS patients with various clinical forms of the disease during the remission or relapse phase (15-20). The aims of the present study were to evaluate the serum cytokine profile in RR-MS patients from a Southern Brazilian population and to evaluate the association between the inflammatory and anti-inflammatory cytokine profiles with clinical disability and disease activity of the patients.

Materials and methods

Study design and subjects. The case-controlled study was approved by the Institutional Research Ethics Committee of the State University of Londrina (Paraná, Brazil) according to the 1964 Declaration of Helsinki and its later amendments. Voluntary written consent was obtained from all individuals prior to the start of the study. The study enrolled 169 consecutive RR-MS patients from the Neurology Outpatients Service of the Outpatient Clinic Hospital, State University of Londrina (Paraná, Brazil), during 2011. RR-MS patients were diagnosed according to the revised McDonald Criteria (21,22). The RR-MS clinical form was determined according to the classification of Lublin and Reingold (23). Demographic, clinical and treatment data of the patients were recorded using a standard questionnaire and medical records. The age at onset of the disease was based on retrospectives data from medical records (21). All the RR-MS patients were in the remission clinical phase, defined as the period of recovery with no relapse episodes within the last three months prior to the time of enrollment in the study. The clinical disability was evaluated using the EDSS (12) at the time of patient enrollment and was categorized as mild (0.0-4.0), moderate (4.5-5.5) and severe (6.0-10.0), according to a previous study (24). The lesion location (brain, spinal cord and optic nerve) and the results of Gd contrast-enhancing lesions were obtained from the most recent MRI scan performed (±3 months from enrollment), as described previously (12,25,26). The control group consisted of 132 unrelated healthy blood donors with similar demographic characteristics and who were from the same geographical area as the patients. Blood samples were obtained from the Blood Bank of Londrina (Paraná, Brazil) All the healthy donors presented negative reactivity for serological screening tests for blood donation according to government procedures (27). Control subjects did not present either clinical or laboratory characteristics of autoimmune, infectious or inflammatory diseases. In addition, subjects reported that they were not using any anti-inflammatory drugs or drugs that affect the immune system.

RR-MS patients and control subjects were age-, gender-, ethnicity- and body mass index (BMI)-controlled. Anthropometric measurements were obtained by body weight, measured to the nearest 0.1 kg using an electronic scale, with individuals wearing light clothing and no shoes. Height was measured to the nearest 0.1 cm using a stadiometer. BMI was calculated as weight (kg) divided by height (m) squared.

Cytokine measurements. Blood samples for cytokine analysis were collected at the time of patient enrollment. The serum samples of MS patients were immediately separated, divided into aliquots and stored in a -80°C freezer for subsequent analysis, as described previously (28). Serum pro-inflammatory cytokine levels (IL-1 β , -12p70 and -6 and TNF- α) as well as Th1 (IFN- γ), Th2 (IL-4 and -10) and Th17 (IL-17) cytokines were measured with a sandwich enzyme-linked immunosorbent assay using a commercial immunoassay (eBioscience, San Diego, CA, USA), according to the manufacturer's instructions. Results were expressed in ng/ml. The limit of detection for the cytokines was defined according to the manufacturer's instructions (IL-1 β , -12p70 and -17, TNF- α and IFN- γ : 2.0 ng/ml and IL-6, -4 and -10: 1.0 ng/ml).

Statistical analysis. A database was created using the Excel Program of Microsoft Office and statistical analysis was performed using Graph Pad Prism v5.0 (Graph Pad Software, La Jolla, CA, USA). Categorical variables were analyzed using a Chi-square test (or Fisher's Exact test where appropriate) and were expressed in absolute number and percentage. Continuous variables were analyzed using the Mann-Whitney test and were expressed as the median, interquartile range (IQR; 25-75%) and range. Continuous variables of two groups or more were compared using the Kruskal-Wallis test with Dunn's post test. P<0.05 was considered to indicate a statistically significant difference. To determine which factors were independently associated with RR-MS, variables that presented P<0.20 in the univariate analyses and values with supposed clinical relevance or previous data in the literature were included in the multivariate logistic regression model. The multivariate analyses were evaluated using GraphPad Instat v3.0 (Graph Pad Software).

Results

Demographic, anthropometric and clinical characteristics. Demographic and anthropometric characteristics of the

Table I. Demographic and anthropometric characteristics of RR-MS patients and healthy controls in a population from South Brazil.

	RR-MS patients	Healthy controls		
Characteristics	(n=169)	(n=132)	P-value	
Gender, n (%)				
Female	123 (72.8)	104 (78.8)	0.2298^{a}	
Male	46 (27.2)	28 (21.2)		
Ethnicity, n (%)				
Caucasian	144 (85.2)	106 (80.3)	0.2604a	
Non-Caucasian	25 (14.8)	26 (19.7)		
Age, years				
Median (IQR)	42.0 (30.0-51.5)	38.5 (32.0-46.7)	$0.1090^{\rm b}$	
Range	14.0-85.0	26.0-62.0		
BMI, kg/m ²				
Median (IQR)	23.5 (21.5-27.2)	24.8 (22.0-27.9)	0.1120 ^b	
Range	16.9-44.0	16.4-40.2		

 a Chi-square test (P<0.05). b Mann Whitney test (P<0.05). RR-MS, relapsing-remitting multiple sclerosis; BMI, body mass index; IQR, interquartile range.

Table II.Clinical characteristics of RR-MS in a population from South Brazil.

Clinical characteristics	n	RR-MS patients
Age at onset, years	169	
Median (IQR)		33.0 (24.5-44.0)
Range		10.0-68.0
Disease duration, years	169	
Median (IQR)		7.0 (3.0-10.0)
Range		0.0-52.0
EDSS, n (%)	146	
Mild (0.0-4.0)		113 (77.2)
Moderate (4.5-5.5)		20 (13.8)
Severe (6.0-10.0)		13 (9.0)
Lesion location at MRI, n (%)	161	
Brain white matter		130 (80.8)
Spinal cord		54 (33.5)
Optic nerve		18 (11.2)
Disease activity at MRI, n (%)	161	
Positive Gd-enhancing lesions		35 (21.7)
Negative Gd-enhancing lesions		126 (78.3)
RR-MS treatment, n (%)	156	
IFN-β1a		37 (23.7)
IFN-β1b		81 (51.9)
Glatiramer acetate		21 (13.5)
Natalizumab		03 (1.9)
Treatment naïve		05 (3.2)
No adhesion		09 (5.8)
Corticosteroids (mg/day)		18 (10.7)

RR-MS, relapsing-remitting multiple sclerosis; BMI, body mass index; EDSS, Expanded Disability Status Scale; MRI, magnetic resonance imaging; IQR, interquartile range; IFN, interferon; Gd, gadolinium.

Table III. Serum cytokine levels obtained from RR-MS patients and healthy controls in a population from South Brazil.

Cytokine	RR-MS (n=169)	Healthy controls (n=132)	P-value ^a
IL-1β (pg/ml)			
Median (IQR)	2.0 (2.0-2.5)	2.0 (2.0-4.5)	0.0992
Range	2.0-40.7	2.0-63.2	
IL-6 (pg/ml)			
Median (IQR)	1.9 (1.0-3.9)	1.1 (1.0-3.4)	0.0114
Range	1.0-158.0	1.0-49.5	
TNF-α (pg/ml)			
Median (IQR)	2.0 (2.0-4.3)	2.0 (2.0-11.2)	0.1840
Range	2.0-68.4	2.0-384.5	
IL-12 (pg/ml)			
Median (IQR)	2.0 (2.0-2.0)	2.0 (2.0-2.0)	0.0297
Range	2.0-93.7	2.0-31.1	
IFN-γ (pg/ml)			
Median (IQR)	42.6 (23.4-74.9)	24.8 (11.7-80.8)	0.0009
Range	2.0-500.0	2.0-476.7	
IL-17 (pg/ml)			
Median (IQR)	2.0 (2.0-2.0)	2.0 (2.0-2.0)	0.2298
Range	2.0-53.2	2.0-88.2	
IL-4 (pg/ml)			
Median (IQR)	1.0 (1.0-4.0)	1.0 (1.0-1.0)	0.0004
Range	1.0-26.4	1.0-16.1	
IL-10 (pg/ml)			
Median (IQR)	4.7 (3.3-6.5)	4.0 (2.9-5.6)	0.1901
Range	1.0-104.9	1.0-28.2	

^aMann Whitney test (P<0.05). RR-MS, relapsing-remitting multiple sclerosis; IL, interleukin; TNF- α , tumor necrosis factor α ; IFN- γ , interferon γ ; IQR, interquartile range.

RR-MS patients and healthy individuals are shown in Table I. As expected, the RR-MS patients and controls did not differ in any of the controlled parameters (P>0.05). The clinical characteristics of RR-MS patients are evident in Table II. The median age at onset of the disease was 33.0 years, with a median disease duration of 7 years. The majority of patients presented mild EDSS and brain white matter lesions without activity. Data of MS therapy revealed the majority of RR-MS patients (75.6%) were treated with IFN- β and only 18 patients (10.7%) were treated with corticosteroids.

Serum cytokine level comparison between RR-MS patients and controls. Serum cytokine levels were compared between RR-MS patients and controls (Table III). With regards to the pro-inflammatory cytokine profile, IL-1 levels were higher among healthy controls compared with RR-MS patients (P=0.0992). IL-6 and -12 levels were higher in RR-MS patients compared with the controls (P=0.0114 and 0.0297, respectively). The univariate analysis demonstrated that serum TNF- α levels observed among RR-MS patients did not differ from those of the controls (median, 2.0 pg/ml and IQR, 2.0-4.3 vs. median, 2.0 pg/ml and IQR, 2.0-11.2; P=0.1840).

In the Th1 cytokine profile, IFN- γ levels were higher among RR-MS patients compared with controls (P=0.0009). However, the Th17 cytokine IL-17 did not differ between RR-MS and controls (P=0.2298).

Th2 anti-inflammatory cytokine profile results revealed that IL-4 levels were higher in RR-MS patients compared with controls (P=0.0004). Although IL-10 levels were higher in RR-MS patients compared with controls, this difference was not considered significant (P=0.1901).

Logistic regression analysis of the results obtained from 169 RR-MS patients and 132 controls included the variables IL-1, -6, -12, -4 and-10, TNF- α and IFN- γ and demonstrated that low TNF- α and high IFN- γ levels were independently associated with RR-MS (P=0.0078 and 0.0056, respectively).

Serum cytokine level according to EDSS disability scores of RR-MS patients. Serum cytokine levels were also evaluated according to the disability of RR-MS patients using categorized EDSS scores in mild, moderate and severe disability (Table IV). Factors, including gender (P=0.2357), ethnicity (P=0.2933), BMI (P=0.2998) and the use of corticosteroids (P=0.4333), were controlled. However, the age of RR-MS

Table IV. Serum cytokine levels obtained from RR-MS patients, according to the EDSS.

Characteristics	Mild EDSS	Moderate EDSS	Severe EDSS	P-value
MS patients, n	146	113	13	
Range	(0.0-4.0)	(4.5-5.5)	(6.0-10.0)	
Gender, n (%)				
Female	83 (56.9)	16 (11.0)	7 (4.8)	0.2357a
Male	30 (20.5)	4 (2.7)	6 (4.1)	
Ethnicity, n (%)				
Caucasian	93(63.7)	19 (13.0)	10 (6.8)	0.2933a
Non-Caucasian	20 (13.7)	01 (0.7)	03 (2.1)	
Age, years				
Median (IQR)	39.0 (28.0-48.0)	39.5 (34.3-56.0)	52.0 (39.5-57.0)	0.0034^{b}
Range	14.0-79.0	24.0-85.0	29.0-71.0	
BMI, kg/m ²				
Median (IQR)	23.4 (21.7-27.4)	24.8(20.5-27.3)	20.9 (19.9-23.9)	0.2998 ^b
Range	17.5-44.0	16.9-33.6	19.8-24.6	
Corticosteroids, mg/day ^c				
Median (IQR)	20.0 (20.0-20.0)	25.0 (20.0-30.0)	20.0 (20.0 -20.0)	0.4333 ^b
Range	2.0-40.0	20.0-30.0	20.0-20.0	
IL-1β, pg/ml				
Median (IQR)	2.0 (2.0-2.0)	2.0 (2.0-5.0)	2.0 (2.0-2.0)	0.4422 ^b
Range	2.0-40.7	2.0-15.9	2.0-2.0	
IL-6, pg/ml				
Median (IQR)	1.8 (1.0-3.8)	2.1 (1.0-3.9)	3.9 (1.3-4.9)	0.7408^{b}
Range	1.0-122.3	1.0-158.0	1.0-5.6	
TNF-α, pg/ml				
Median (IQR)	2.0 (2.0-5.0)	2.0 (2.0-3.3)	2.7 (2.0-28.8)	$0.0572^{\rm b}$
Range	2.0-50.1	2.0-4.7	2.0-30.0	
IL-12, pg/ml				
Median (IQR)	2.0 (2.0-2.0)	2.0 (2.0-2.0)	2.0 (2.0-4.5)	0.5083 ^b
Range	2.0-93.7	2.0-7.5	2.0-11.8	0.000
INF-γ, pg/ml				
Median (IQR)	41.2 (19.6-77.7)	55.5 (27.1-36.3)	52.6 (36.3-72.0)	0.6032 ^b
Range	5.0-500.0	9.8-500.0	27.0-500.0	0.0002
IL-17, pg/ml				
Median (IQR)	2.0 (2.0-2.0)	2.0 (2.0-2.0)	2.0 (2.0-5.5)	0.2359b
Range	2.0-53.2	2.0-4.2	2.0-13.7	0.2003
IL-4, pg/ml				
Median (IQR)	1.0 (1.0-9.6)	1.0 (1.0-1.4)	1.0 (1.0-1.0)	0.0375 ^b
Range	1.0-26.4	1.0-17.2	1.0-1.0	0.0313
IL-10, pg/ml	2.0 2011	1.0 1/12	1.0 1.0	
Median (IQR)	4.6 (3.2-6.6)	4.4 (3.4-6.8)	5.0 (3.0-5.2)	$0.8680^{\rm b}$
Range	1.0-21.9	2.2-16.5	2.9-5.4	0.0000

^aChi-square test (P<0.05). ^bKruskal-Wallis test with Dunn's Post test (P<0.05). ^cEighteen RR-MS patients were treated with corticosteroids at the time of enrollment in the study. RR-MS, relapsing-remitting multiple sclerosis; EDSS, Expanded Disability Status Scale; IL, interleukin; TNF- α , tumor necrosis factor α ; IFN- γ , interferon γ ; IQR, interquartile range.

patients that exhibited severe disability was higher than that of RR-MS patients with mild and moderate disability (P=0.0034).

Serum IL-1 levels did not differ according to the disability exhibited by the RR-MS patients (P=0.4422). Although IL-6, TNF- α and IFN- γ levels were higher among the RR-MS

Table V. Serum cytokine levels obtained from RR-MS patients according to Gd-enhancing lesions assessed by MRI.

	M		
Characteristics	Positive Gd	Negative Gd	P-value
RR-MS patients, n	35	126	
Gender, n (%)			
Female	24 (14.9)	93 (57.8)	0.5383a
Male	11(6.8)	33 (20.5)	
Ethnicity, n (%)			
Caucasian	29 (18.0)	107 (66.5)	0.7656a
Non-Caucasian	06 (3.7)	107 (66.5)	
Age, years			
Median (IQR)	32.0 (27.0-44.0)	44.0 (32.0-52.3)	0.0032^{b}
Range	14.0-79.0	15.0-85.0	
BMI, kg/m ²			
Median (IQR)	23.4 (22.1-27.7)	23.5 (21.1-27.1)	$0.6464^{\rm b}$
Range	19.3-35.5	16.9-44.0	
Corticosteroids, mg/day ^c			
Median (IQR)	20.0 (20.0-20.0)	20.0 (20.0-22.5)	0.9029^{b}
Range	20.0-40.0	5.0-30.0	
IL-1β, pg/ml			
Median (IQR)	2.0 (2.0-2.0)	2.0 (2.0-2.9)	0.4752 ^b
Range	2.0-18.0	2.0-40.7	
IL-6, pg/ml			
Median (IQR)	1.3 (1.0-3.2)	2.4 (1.0-4.0)	0.1129 ^b
Range	1.0-158.0	1.0-122.3	5.1123
TNF-α, pg/ml			
Median (IQR)	2.0 (2.0-2.1)	2.0 (2.0-5.6)	$0.0457^{\rm b}$
Range	2.0-17.5	2.0-68.4	0.0.157
IL-12, pg/ml			
Median (IQR)	2.0 (2.0-2.0)	2.0 (2.0-2.0)	$0.9089^{\rm b}$
Range	2.0-48.3	2.0-93.7	0.5005
IFN-γ, pg/ml			
Median (IQR)	52.8 (25.0-131.4)	39.5 (23.0-70.7)	0.1643 ^b
Range	6.4-500.0	2.0-500.0	0.1043
IL-17, pg/ml	0.1 500.0	2.0 300.0	
Median (IQR)	2.0 (2.0-2.0)	2.0 (2.0-2.6)	0.0631 ^b
Range	2.0-15.6	2.0-53.2	0.0031
IL-4, pg/ml	2.0 13.0	2.0 33.2	
Median (IQR)	1.0 (1.0-14.9)	1.0 (1.0-4.1)	$0.8980^{\rm b}$
Range	1.0-14.9)	1.0-26.4	0.0900
_	1.0-22.3	1.0-20.7	
IL-10, pg/ml Median (IQR)	3.4 (2.3-5.8)	4.9 (3.4-6.5)	$0.0533^{\rm b}$
	3.4 (2.5-3.8) 1.0-15.9	4.9 (3.4-6.5) 1.0-104.9	0.0333
Range	1.0-13.9	1.0-104.9	

^aChi-square test (P<0.05). ^bMann Whitney test (P<0.05). ^cEighteen RR-MS patients were treated with corticosteroids at the time of enrollment in the study. RR-MS, relapsing-remitting multiple sclerosis; BMI, body mass index; IL, interleukin; TNF-α, tumor necrosis factor α ; IFN- γ , interferon γ ; IQR, interquartile range; Gd, gadolinium.

patients with severe disability compared with those observed among the RR-MS patients with moderate and mild disability, the difference was not significant (P=0.7408, 0.6896 and

0.6032, respectively). No significant difference was found in IL-12 and -17 levels obtained among the RR-MS patients with mild, moderate and severe disability (P=0.5083 and

0.2359, respectively). Higher IL-4 levels were observed among RR-MS patients with mild disability compared with moderate and severe disability (P=0.0375). However, no difference was observed in IL-10 levels with respect to patient disability (P=0.8680). Following logistic regression analysis, variables presented no independent association with disease disability (P>0.05).

Serum cytokine level according to disease activity. Serum cytokine levels obtained from RR-MS patients were also evaluated according to the disease activity assessed by Gd-enhancing lesions observed during MRI (Table V). Factors, including gender (P=0.5383), ethnicity (P=0.7656), BMI (P=0.6464) and the use of corticosteroids (P=0.9029) were controlled. However, the age of RR-MS patients with positive Gd-enhancing lesions was lower compared with RR-MS patients with negative Gd-enhancing lesions (P=0.0032).

No difference was detected in IL-1, -6, -12 and -4 levels between RR-MS patients with positive and negative Gd-enhancing lesions (P=0.4752, 0.1129, 0.9089 and 0.8980, respectively). Although IFN- γ levels were higher among RR-MS patients presenting positive Gd-enhancing lesions compared with negative ones, no difference was found by univariate analysis (P=0.1643). TNF- α levels were higher in patients with negative Gd-enhancing lesions than positive (P=0.0572). Serum IL 17 levels showed a growing trend towards being higher in RR MS patients with inactive disease compared to those with active disease (P=0.0631). While serum IL-10 levels were higher among RR-MS patients with negative Gd-enhancing lesions than those with positive (P=0.0533). IL-4 levels did not differ between these patients (P=0.8980).

Logistic regression analysis between the results obtained from RR-MS patients and the disease activity, including the variables age, TNF- α , IFN- γ and IL-6, -17 and -10, demonstrated that low TNF- α and high IFN- γ levels were independently associated with disease activity (P=0.0348 and 0.0133, respectively).

Discussion

The present study evaluated the serum cytokine profile exhibited by RR-MS patients in a remission clinical phase from a population from South Brazil. Theoretically, CSF may reflect the relevant inflammatory process to a higher degree compared with other samples, due to its proximity to inflammatory lesions in the CNS. However, due to the flow pattern of CSF, it is unlikely that CSF in the lumbar cistern accurately reflects the production of inflammatory markers in the supratentorial region, where the majority of MS-related inflammation occurs (29). In addition, CSF collection is an invasive procedure, the intraparenchymal extracellular space may not necessarily communicate with the free CSF space (30) and it remains unclear whether CSF markers possess definite advantages over markers collected from blood and/or urine (29). Therefore, in the current study the cytokine profile was measured in serum samples.

To obtain more homogeneity of the cohort, only patients with the RR-MS clinical form were included in this study. The main result obtained in this study was elevated serum IFN- γ

and IL-6, -12 and -4 levels in RR-MS patients compared with controls. MS patients with mild EDSS presented elevated IL-4 levels. RR-MS patients with inactive disease exhibited elevated levels of TNF- α and IL-10 and -17. Logistic regression analysis revealed that low TNF- α and high IFN- γ levels were associated with MS and disease activity.

The demographic and clinical characteristics exhibited by the RR-MS patients are consistent with those of previous studies (18,24,31,34). The majority of RR-MS patients exhibited mild clinical disability, predominance of lesions in brain white matter and low disease activity. These results were expected considering all the RR-MS patients were in clinical remission and the majority of them were treated with IFN- β (1a or b) which has been demonstrated to decrease clinical disability and disease activity (18,35-37).

RR-MS patients and controls were age-, gender-, ethnicity- and BMI-matched for cytokine analysis. These parameters were controlled to minimize their effect of them on serum cytokine levels (38,39). IL-1 levels were higher among controls compared with RR-MS patients and may be explained by the hypothesis that IFN- β therapy decreases IL-1 β expression in the dendritic cells and monocytes of RR-MS patients (40). This pro-inflammatory cytokine is considered an important mediator for neuronal and axonal damage of the CNS in MS patients. One of the actions of IL-1 β in the brain is the induction of astrogliosis by astrocyte activation and IL-6 production, an effect that is markedly augmented in the presence of TNF- α , IFN- γ , IL-6 and the IL-6 receptor (41). However, IL-1 has also been revealed to induce proliferation of the nerve growth factor *in vitro* and may exhibit protective effects in MS patients (42).

The increased IL-6 levels identified among RR-MS patients compared with controls are consistent with previous studies. One study demonstrated that the number of IL-6 mRNA expressing mononuclear cells was enriched in CSF of MS patients (43). Increased IL-6 production was also observed in peripheral blood monocytes and macrophages stimulated with LPS from MS patients (44). IL-6 deficient mice were found to be resistant to induction of EAE (17) and also revealed a reduction of anti-MOG antibody titers, vascular cell adhesion molecule-1 (VCAM-1) and intercellular adhesion molecule-1 (ICAM-1) molecule expression, which may affect the entry of Th1 cells into the CNS (45). By contrast, IL-6 also presented beneficial effects, including anti-inflammatory properties. IL-6 inhibits IL-1β and TNF-α secretion in vitro (46) and induces circulating IL-1 receptor antagonist in vivo (47). Results of the current study are consistent with those of previous studies considering that elevated IL-6 levels were found in RR-MS patients, IL-1 levels were higher in controls and no difference was observed in TNF- α levels.

Navikas *et al* (48) demonstrated that MS patients present with high numbers of MBP-reactive cells in their blood which, following culture with MBP and myelin PLP, respond to production and secretion of additional cytokines, including IFN- γ , IL-4, transforming growth factor β (TGF- β), IL-10 and TNF- α . The authors of that study demonstrated that MS patients presented elevated MBP- and PLP-reactive IL-6 mRNA expressing cells in the peripheral blood. Cytokines are known to modulate production of other cytokines (49). Therefore, IL-6 functions with additional cytokines within the cytokine network.

TNF- α levels were lower in RR-MS patients than the controls, as demonstrated by the multivariate analysis. This result is consistent with previous studies reporting lower TNF- α levels in MS patients compared with controls (11,19). Decreased production of TNF- α by stimulated PBMCs from MS patients compared with controls was also demonstrated (3).

In the present study, increased IFN- γ levels obtained among MS patients compared with controls were also consistent with previous studies. Lymphocytes producing IFN- γ were identified in the perivascular infiltrates of MS patients (4). A previous study evaluated mRNA serum transcript in patients receiving various doses of IFN- β 1b and confirmed that IFN- β decreases the expression of Th1 inflammatory molecules, including IFN- γ and TNF- α . In addition, IFN- β increases the Th2-related molecule IL-10 (3).

Although IL-17 expression has been demonstrated to generate a highly pro-inflammatory environment and to induce severe pathological conditions (50), in the present study no difference was observed in IL-17 levels between MS patients and controls. A previous study (51) revealed that the proportion of Th17 cells in patients with relapses were significantly higher compared with patients during remission, in clinically isolated syndrome (CIS) and RR-MS clinical forms of MS. Another study (52) also revealed that IFN-β inhibits IL-23 which is involved in the differentiation of Th0 (naïve) to Th17 cells and, consequently, IL-17 production. In addition, IFN-β therapy inhibits the IL-27 cytokine which, in turn, inhibits Th17 cells directly and through the inhibition of IL-23, promotes development of these pathogenic T cells (8). IL-23 and -12 exert antagonistic effects. While IL-23 supports IL-17 production, IL-12 inhibits this production (53). IL-17, in turn, stimulates astrocyte IL-6 production and macrophage production of IL-6, TNF-α and IL-1β, cytokines that are involved in MS pathology (54).

The RR-MS patients enrolled in this study presented high levels of IL-12 which, in turn, antagonized IL-17 production and by this effect, reduced IL-1 β - and TNF- α -induced production by IL-17. In addition to the consumption of TNF- α in CNS lesions, the reduced production may explain, in part, why TNF- α levels did not differ among the patients and controls (54).

The higher IL-12 levels exhibited by RR-MS patients compared with controls enrolled in the present study are also consistent with previous observations that IL-12 levels were upregulated in MS patients (55). Overexpression of IL-12 in the CNS resulted in increased inflammation and cellular infiltrates (56). Although several studies have suggested a key pro-inflammatory role for IL-12 in the development of autoimmune responses, controversial results have been reported with respect to IL-12 levels obtained from MS patients and MS patients treated with IFN-β. A previous study (40) demonstrated that treatment with IFN-β1a significantly induces IL-12p35 gene expression in dendritic cells from RR-MS patients. By contrast, it was demonstrated that treatment with IFN-β inhibits IL-12 and decreases C-X-C motif chemokine 10, IL-18, IFN- γ and TNF- α (57). However, this therapy increases IL-10 and TGF-β levels (58).

In the Brazilian cohort enrolled in the present study, higher levels of IL-4 were found in MS patients compared with controls, a result consistent with previous studies (59,60).

IL-4 mRNA was previously revealed to be undetectable until disease remission in SJL mice immunized with PLP (61). IL-4 is a key cytokine produced by Th2 cells and inhibits Th1 cells by decreasing IL-1 and TNF- α cytokines. Administration of IL-4 in EAE exerts an ameliorating and protective effect. Studies demonstrated that IL-4 upregulation may reduce the severity of EAE, while its absence does not alter the course of the disease, possibly since in the absence of IL-4, an additional Th2 cytokine may substitute its functions and contribute to the tolerance of EAE (62).

Although IL-10 levels were higher in MS patients compared with controls, no statistically significant difference was found. It was previously suggested that IL-10 is important for recovery of EAE (63). In contrast to other Th2 cytokines (IL-4), the absence of IL-10 results in a more severe EAE, indicating that its function cannot be replaced by an additional Th2 cytokine.

The cytokine profile was also evaluated according to the EDSS of the RR-MS patients. Patients with mild disability were younger than those with moderate and severe disability, a result that was expected since at the beginning of the disease, low EDSS scores are detected among RR-MS patients and disability increases during disease progression (64). However, logistic regression analysis revealed no effect of age in the cytokine profile according to the EDSS. Although higher serum IL-6, TNF-α, IFN-γ and IL-12 levels were found in RR-MS patients with moderate and severe disability than those with mild disability, no significant differences were identified. As observed, the results of TNF- α levels are contradictory in various clinical disabilities. While low levels of TNF- α have been reported in specific MS patients (11,19), another study reported that TNF- α is the principal mediator of inflammatory response and is increased among MS patients, indicating that this cytokine is an important molecule in the pathogenesis and progression of MS (65).

In addition, serum IL-17 and -10 did not differ between EDSS categories. Several studies have reported that differential production of IL-10 may be a factor in the severity of MS (31,63). With regards to serum IL-4 levels, results demonstrate a significant increase of this cytokine among RR-MS patients suffering from mild disability compared with those with moderate or severe disability. This result confirms the important role of IL-4 as an anti-inflammatory modulator of MS lesion formation (6,19).

It was not possible to control age of RR-MS patients with regards to disease activity. However, logistic regression analysis revealed that age did not affect cytokine levels. There is evidence that an acute deregulation in the balance of cytokines is one of the key factors during an acute relapse leading to acute inflammatory lesions. RR-MS patients with active disease presented with lower TNF-α and higher IFN-γ levels compared with RR-MS patients without active disease, as demonstrated by multivariate analysis. IFN-y and IL-12 levels were increased in brain, CSF or peripheral blood from MS patients, particularly during acute exacerbation phases (5). Previously, Begolka and Miller (61) observed that IFN-y expression increases during the peak of the disease and decreases during the remission phase. Clinical attacks correlate with increased IFN-y production (66) and patients experiencing relapses have significantly increased IFN-γ production by PBMCs following stimulation with PHA

compared with patients in remission. However, this production was reduced following treatment with IFN- β (67). IFN- γ is produced within MS lesions by infiltrating lymphocytes and induces apoptosis in human oligodendrocytes (68).

Cytokines only function when bound to membrane receptors and the TNF- α receptor is present in almost all cells (6,61,69). In MS patients, TNF- α is produced and consumed at sites of inflammation, including lesions in brain white matter, spinal cord and optic nerve (70,71). TNF- α and - β cytokines were detected in astrocytes from demyelinating plaques (71), contributing to white matter destruction at the onset of MS (16). Results obtained in the present study are consistent with these previous findings. However, serum TNF- α levels measured in RR-MS patient samples of the present study may be the remaining expressed cytokines, neither acting at lesions on the CNS, nor binding to transmembrane or soluble receptors. Additional studies also demonstrated lower serum TNF- α levels in MS patients in remission compared with those in the relapse clinical phase (19,20).

Although IL-6 levels were higher in RR-MS patients with inactive disease, this difference was not identified to be statistically significant. This result is in agreement with a previous study (72) which demonstrated that macrophages within the MS lesion expressed IL-6 and the highest number of IL-6 expressing cells were found in inactive demyelination lesions. We hypothesize that IL-6 also has anti-inflammatory properties (46) that may affect disease activity.

A previous study has demonstrated that serum IL-12 levels correlate with active lesions (73), however, no significant difference was observed in serum IL-12 levels obtained from RR-MS patients with positive and negative Gd-enhancing lesions enrolled in the present study.

Increased levels of IL-17 were found in RR-MS patients with inactive disease. Th17 cell differentiation is orchestrated by multiple cytokines, including TGF- β and IL-6, -1- β , -21 and -23 (stimulating) and IFN- γ and IL-4, -12, -10 and -27 (inhibiting). The Th17 cell is a recently identified cell lineage that has been hypothesized to be critical for the development of the autoimmune response, in addition to Th1 cells (74). Studies from MS patients indicated that IL-17 mRNA is elevated in active MS brain lesions. IL-17 was also detected at lesions (75).

Serum IL-4 levels did not differ between RR-MS patients with active and inactive disease and this result was not consistent with previous studies that demonstrated high IL-4 levels in acute and chronic lesions (62). However, in the present study, serum IL-10 levels were high among RR-MS patients with inactive lesions. This result is consistent with previous observations of increased production of IL-10 and TGF- β by isolated lymphocyte clones of MS patients in remission (76). Decreased IL-10 levels were found prior to onset of exacerbation in RR-MS patients (15). IL-10 levels were also significantly decreased in MS patients with the secondary progressive form of MS compared with patients at four weeks prior to detection of MS by MRI and six weeks prior to clinical relapse (73).

Previously, Cannella and Raine (62) reported that the expression of adhesion molecules, including VCAM-1, ICAM-1 and their receptors on leukocytes, very late activation antigen-4 (VLA-4) and lymphocyte function-associated antigen-1

(LFA-1), respectively, together with a selection of pro-inflammatory and immunomodulatory cytokines including, IL-1, -2, -4 and -10, TNF- α , TGF- β and IFN- γ , are important for lymphocytic infiltration and interactions during tissue inflammation. In general, these molecules are not expressed by CNS cells. The authors identified high levels of these molecules in MS patients, particularly in those with chronic active lesions and also found a significantly increased expression of TNF- α and IL-4 in MS compared with non-inflammatory neurological diseases (62).

The present study demonstrates that RR-MS patients exhibit a complex system of cytokines which interact to modulate disease activity and the disabilities associated with it. Even in the remission phase of the disease and under immunomodulatory therapy, RR-MS patients present with an imbalance between pro-inflammatory, Th1, -2 and -17 cytokines. TNF- α was observed to be consumed in CNS lesions and activation of the cellular immune response was mediated by high levels of IFN- γ . These results are likely to contribute to an improved understanding of the role of the cytokine profile in RR-MS and may aid the development of new neuroprotective and neurorestorative therapeutic strategies.

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