

MANNOSE BINDING LECTIN IN SERA AND SYNOVIAL FLUIDS OF PATIENTS WITH RHEUMATOID ARTHRITIS

Zoran Mijušković¹, Miodrag Čolić², Ivan Zgradić³, Ljiljana Rackov³,
Goran Radunović⁴, Nada Pilipović⁴, Radmila Nikolajević¹, Jelica Stojanović⁵

¹Institute of Medical Biochemistry,

²Institute of Medical Research,

³Department of Clinical Rheumatology and Immunology,
Military Medical Academy, Belgrade,

⁴Institute of Rheumatology, Belgrade, Yugoslavia,

⁵Faculty of Natural Sciences and Mathematics, Kragujevac, Yugoslavia

Summary: It has been supposed that low levels of mannose binding lectin (MBL) may be involved in protection of rheumatoid arthritis associated with agalactosil forms of GOIgG immunoglobulins found in rheumatoid synovial fluid and lead to enhanced complement activation and joint cartilage damages. In order to examine the possible link between MBL and mediated complement activation, its concentration was measured in sera and synovial fluids of 59 patients classified in moderate and severe activity RA groups and compared with the levels in the same biological fluids of the control group with menisci lesions. No patients had undetectable or lowered MBL in both RA groups. Mean values were either normal or elevated depending on the severity of the disease as a response to inflammation. This finding does not exclude a possible role of MBL-GOIgG mediated complement activation, but provides no support to mitigating RA severity.

Key words: rheumatoid arthritis, mannose binding lectin, complement, agalactosil IgG

Introduction

Human mannose-binding lectin (MBL) is a macromolecule protein with an apparent M_w from 270 to 650 KD and consists of about 9 to 18 identical subunits-trimers of 32 KD. Each of them has carbohydrate-recognition domain (CRD) at the COOH-terminus, a collagen like domain (CLD) and cystein rich domain at the NH₂-terminal portion. CLD associated with 32 KD trimers is a structure unit which can further assemble into higher multimers forming dimeric to hexameric units. So, the gross structure of MBL is remarkably homologous with C1q complement component. MBL is Ca²⁺ dependent, but unlikely to C1q, functions in antibody as independent host defence against pathogens. Its ability to bind carbohydrate structures (mannose or N-acetylglucosamine) is wide-

ly present on yeasts, bacteria and viruses can thereby activate the complement system (1). Deficiency of MBL has been linked to recurrent infections in infants, older children and adults (2–4). The concentration of MBL in sera is largely under genetic control and depends primarily on genetic polymorphism within the first exon of the structural gene (5). A point of mutation at codon 54 gives rise to a mutant allele which has a gene frequency (GF) of 0.11–0.19 in European populations (1, 6). Two other mutant alleles at codon 52 (GF: 0.02–0.05) and 57 (GF: 0.07–0.29) are known and are relatively common in African and Eskimo populations. It has been shown that local activation of complement may play a role in mediating synovial inflammation (7) in rheumatoid arthritis (RA). Synovial fluid (SF) in RA contains high concentrations of immunoglobulin and rheumatoid factors (RF) which, through their ability to form immune complexes, may promote complement activation. In addition, distinctive glycosylation pattern of these synovial immuno-

Address for correspondence

Zoran V. Mijušković
Military Medical Academy
Institute of Medical Biochemistry
Crnotravska 17, 11002 Belgrade, Yugoslavia
e-mail: zmijusb@YUIBC.net

* Invited paper presented on 13th Congress of Medical Biochemistry and Laboratory Medicine, May 14–18, 2002, Niš, Yugoslavia

globulins have been reported, including a raised proportion of IgGs which contain galactose deficient oligosaccharides on side chains in hinge region of CH₂ domain (8, 9). Thus, such IgG is called agalactosil IgG (G0IgG), which terminates with mannose or N-acetylglucosamine becomes accessible for binding to MBL (10) and can mediate opsonization and lectin »pseudo-classical« complement pathway (11). Some evidence confirmed that G0IgG has been implicated in pathogenesis of RA in an animal model (12). It has also been suggested that lower levels of MBL would limit the tissue damage mediated by complement activation and might prevent excessive uptake of pathogens or mitigate the severity of chronic inflammatory disorders. In addition, some authors (13, 14) have hypothesized that lower MBL levels and activity might be protective against rheumatoid disorders. Another author (15) suggests that it would have been of interest to compare MBL levels in patients with active RA with those in patients in remission. MBL concentrations from the so called »pseudoclassical« complement pathway and their possible relations with circulated immune complexes (CIC) and RF isotypes were compared with complement C3 and C4 components of a classical pathway. To test all mentioned observations, taking into account that RA is predominately an articular disease, it has been interesting to compare MBL levels in SFs and sera of different RA activity groups and compare them with that of the control group (CG) without synovitis.

Patients and Methods

In this multicentric study 59 RF positive sera and SFs were examined in RA patients for the level of MBL and compared with the same biological fluids of a control group of 15 patients with lesion menisci. All RA patients satisfied American Rheumatism Association criteria (16) and were classified in two groups: One with moderate (n=24) disease activity and the other with severe (n=35) disease activity. Activity of RA was determined according to adequate clinical (number of painful and swollen joints, Steinbrocker's radiographic changes, Ritchie's articular index, Disease Activity Score) and biochemical (erythrocyte sedimentation rate, fibrinogen, C-reactive protein and serum amyloid A concentration) parameters.

Concentrations of MBL, CRP, SAA, CIC and IgMRF in both fluids were examined by latex immunonephelometry (»DADE Behring«) method, while IgGRF and IgARF were measured by enzyme immuno (»CogentDiagnostics«) assays. Before testing all SF samples were pretreated with hyaluronidase (*Streptomyces hyalurolyticus*).

Statistical analyses were performed using Basic Statistic software for comparison of means (Mann-Whitney U test), standard deviation and data distribution.

Results

All patients were IgMRF, IgGRF or IgARF positive, alone or in RF isotype combinations, but there

Table I Comparison of C3c, C4 complement components, MBL and CIC in sera and SFs in three tested groups

CONTROL GROUP N=15		$\bar{x} \pm SD$ (g/L)	$\bar{x} \pm SD$ (g/L)	statistical significancy	% of \bar{x} C3, C4, CIC in SF vs. serum
SERUM	vs. SF				
C3c	vs. C3c SF	1.12 ± 0.26	0.28 ± 0.16	p < 0.001	- 75 %
C4	vs. C4 SF	0.25 ± 0.09	0.07 ± 0.06	p < 0.001	- 72 %
CIC	vs. CIC SF (IU/mL)	1.81 ± 0.76	2.42 ± 0.66	p < 0.05	+ 34 %
MBL	vs. MBL SF (mg/L)	2.05 ± 1.03 r.i.: 0.09-5.6	1.11 ± 0.76	p < 0.05	- 54 %
MODERATE ACTIVITY of RA N=24		serum	SF		
C3c	vs. C3c SF	1.22 ± 0.37	0.41 ± 0.28	p < 0.001	- 66 %
C4	vs. C4 SF	0.25 ± 0.08	0.12 ± 0.18	p < 0.001	- 52 %
CIC	vs. CIC SF (IU/mL)	3.01 ± 2.92	5.33 ± 7.53	p < 0.05	+ 77 % *
MBL	vs. MBL SF (mg/L)	2.56 ± 1.51	0.99 ± 0.89	p < 0.05	- 39 %
SEVERE ACTIVITY of RA N=35		serum	SF		
C3c	vs. C3c SF	1.33 ± 0.36	0.44 ± 0.3	p < 0.001	- 67 %
C4	vs. C4 SF	0.29 ± 0.13	0.10 ± 0.1	p < 0.001	- 69 %
CIC	vs. CIC SF (IU/mL)	7.43 ± 13.4	13.5 ± 21.1	p < 0.01	+ 82 % *
MBL	vs. MBL SF (mg/L)	3.34 ± 1.9	1.46 ± 0.9	p < 0.05	- 44 %

Abbreviations: CIC – circulated immune complexes; SF – synovial fluid; vs. – versus; N = number of patients; \bar{x} – arithmetic mean; SD – standard deviation; p – statistical significancy; r.i. – reference interval; * p < 0.05 vs. adequate value in control group (t – test proportions).

Table II Comparison of C3c, C4 complement components, MBL and CIC in sera and SFs in three tested groups

ACTIVITY of RA groups SEVERE vs. MODERATE N = 35 vs. N = 24			$\bar{x} \pm SD$ (g/L)	$\bar{x} \pm SD$ (g/L)	statistical significancy
C3c	vs.	C3c	1.33 ± 0.40	1.22 ± 0.37	NS
C3cSF	vs.	C3c SF	0.44 ± 0.30	0.41 ± 0.28	NS
C4	vs.	C4	0.29 ± 0.10	0.25 ± 0.08	NS
C4 SF	vs.	C4 SF	0.10 ± 0.10	0.12 ± 0.18	NS
CIC	vs.	CIC (IU/mL)	7.43 ± 13.40	3.01 ± 2.92	p < 0.05
CIC SF	vs.	CIC SF (IU/mL)	13.47 ± 21.1	5.33 ± 7.53	p < 0.001
SEVERE N = 35	vs.	CONTROL N = 15			
C3c	vs.	C3c	1.33 ± 0.40	1.12 ± 0.26	NS
C3cSF	vs.	C3c SF	0.44 ± 0.30	0.28 ± 0.16	NS
C4	vs.	C4	0.29 ± 0.10	0.25 ± 0.08	NS
C4 SF	vs.	C4 SF	0.10 ± 0.10	0.07 ± 0.06	NS
CIC	vs.	CIC (IU/mL)	7.43 ± 13.40	1.81 ± 0.76	p < 0.001
CIC SF	vs.	CIC SF (IU/mL)	13.47 ± 21.1	2.42 ± 0.66	p < 0.001

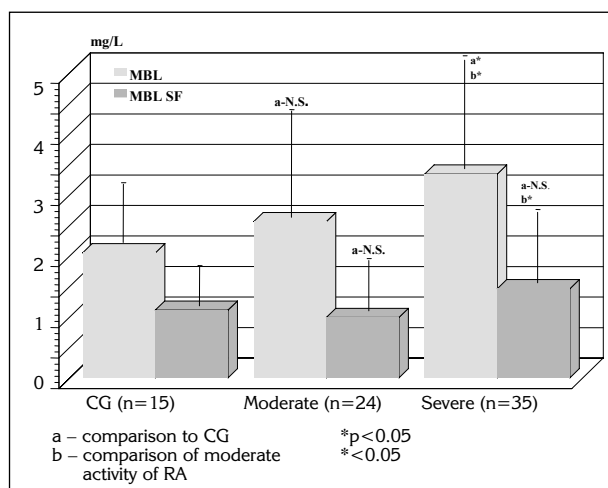


Figure 1. Comparison of MBL concentration in sera and SFs among control group and groups with moderate and severe RA activity

was no correlation ($r < 0.03$; $p > 0.05$) between RF and CIC levels, with MBL concentration neither in sera nor in SF. MBL sera concentrations were either normal or elevated in about 11% of patients in both MA and SA RA groups. In SF MBL group they were 39%, 44% and 54% lower than sera levels in MA, SA and CG respectively, but these differences were not statistically significant (Table I). Similar results were obtained for C3 and C4 components of complement which were lower in SF (between 55% and 75%). Marked elevation was only found in CIC levels in SF in both RA activity groups in spite to CG; this finding

was statistically significant regarding the t-test proportions.

An important result was that no sample of sera of both RA groups had undetectable MBL concentrations. Mean average MBL values in sera, but not in SF, were significantly higher ($p < 0.05$) in SA group compared with MA group, but such differences were not detected in groups MA and CG (Figure 1).

In both cases of the disease activity the SM filtration rate was the same, while CIC concentration in SF was significantly higher in contrast to sera levels. CIC levels in sera and SF compared among SA vs. MA were higher but less than in SA group vs. CG (Table II). No similar differences were found for the level of C3 and C4.

Discussion

It was expected that patients with RA would have an increased average of MBL concentrations, and conversely, that low MBL concentration would be absent or under-represented. In fact, the obtained results showed statistically higher mean values in sera of SA vs. MA RA groups, but not regarding concentrations in SF. A higher MBL concentration in SF of CG than in sera in RA groups could be explained by a possible more severe SM destruction and, consequently, worse filtration rate in patients with lesion menisci in comparison to SM destruction provoked by inflammation in RA groups.

Results regarding MBL level appeared as a consequence of severe inflammation among the tested groups in relation to MBL as an acute phase reacted

(APR) production. More over, data that none of 59 RA patients had undetectable or significantly lower MBL levels even in MA RA group indicated that the frequency of MBL codon 54 mutation was rather low in RA patients. Result from this study confirmed no hypothesis that in RA, MBL deficiency might mitigate the effects of complement damage mediated by a pathway involving GOIgG, MBL binding and complement activation (10), and so be associated with less severe disease. The others showed no relationship between the presence of codon 54 mutation allele and other indicators of severe RA, such as RF (14). Our previous results revealed a connection between RA severity and high level of different RF isotype combinations (17, 18). In all, this study also suggests that MBL deficiency does not protect against severe RA. Similar conclusions have been reported in a study of MBL protein levels only in sera of RA patients, but were not classified according to the severity of the disease (15).

Unclear aetiopathology of RA up to date, involves a large spectrum of biological molecules with several antigen or autoantigen and adequate antibodies, producing high levels of CIC, which are

also important for the severity of RA. The obtained higher CIC level in SF than in serum implies its local production, which could be seen from the result obtained in this study. However, which part of CIC levels belongs to GOIgG complexes and complement activation via MBL is an open question. Complement activation or consumption of C3 and C4 in sera is hardly to detect for the same reason as MBL, because all mentioned molecules belong to APR family. Their production in response to the inflammation possibly masks the known consumption. Some authors, therefore suggest other complement degradation of split products as more relevant for complement activation in RA (19).

In conclusion, the findings presented here do not exclude the putative role of MBL-GOIgG interaction in complement activation in RA, but there is also no evidence of supporting the opinion that such complexes play a major role in the pathogenesis, possible prevention or perpetuation of RA. Measuring MBL as well C3 and C4 in a laboratory routine practice is therefore, time consuming without marked relevant data.

MANOZA VEZUJUĆI LEKTIN U SERUMU I SINOVIJALNOJ TEČNOSTI BOLESNIKA SA REUMATOIDNIM ARTRITISOM

Zoran Mijušković¹, Miodrag Čolić², Ivan Zgradić³, Ljiljana Rackov³,
Goran Radunović⁴, Nada Pilipović⁴, Radmila Nikolajević¹, Jelica Stojanović⁵

¹Institut za medicinsku biohemiju

²Institut za medicinska istraživanja

³Klinika za reumatologiju i kliničku imunologiju
Vojnomedicinska akademija, Beograd

⁴Institut za reumatologiju, Beograd

⁵Prirodno-matematički fakultet, Kragujevac

Kratak sadržaj: Pretpostavljalo se da niska koncentracija manoza-vezujućeg lektina (MBL) može biti povezana sa zaštitom prema ispoljavanju reumatoidnog artritisa (RA), zajedno sa gliko formom agalaktozil GOIgG koji je ustanovljen u sinovijalnoj tečnosti. Ovakav kompleks olakšava aktivaciju sistema komplementa u procesu razgradnje zglobne hrskavice. U cilju iznalaženja povezanosti aktivacije komplementa posredstvom MBL, određivana je koncentracija ovog lektina u serumu i sinovijalnoj tečnosti kod 59 bolesnika koji su razvrstani u grupe sa umerenom i visokom aktivnošću RA. Rezultati su upoređeni sa koncentracijom MBL u oba biološka uzorka iz kontrolne grupe pacijenata sa povredama meniskusa. Nijedan ispitanik nije imao nemerljivu ili sniženu koncentraciju MBL. Srednje vrednosti su bile u okviru fizioloških granica ili pak povišene zavisno od stepena aktivnosti RA, kao proteinski odgovor na proces inflamacije. Ovi rezultati ne isključuju moguću ulogu kompleksa MBL-GOIgG u aktivaciji komplementa, ali ne idu u prilog podrške mišljenju da MBL sudeluje u ublažavanju jačine ispoljavanja simptoma bolesti.

Cljučne reči: manoza vezujućii lektin, reumatoidni artritis, komplement, agalaktozil IgG

References

1. Turner MW. Mannose binding protein. *Biochem Soc Trans* 1994; 22: 88–94.
2. Sumiya M, Super M, Tabona P et al. Molecular basis of opsonic defect in immunodeficient children. *Lancet* 1991; 337: 1569–70.
3. Summerfield JA, Ryder S, Sumiya M et al. Mannose binding protein gene mutations associated with unusual and severe infections in adults. *Lancet* 1995; 345: 886–9.
4. Garred P, Madsen HO, Hofmann B, Svejgaard A. Increased frequency of homozygosity of abnormal mannan binding protein alleles in patients with suspected immunodeficiency. *Lancet* 1995; 346: 941–3.
5. Madsen HO, Garred P, Thiel S et al. Interplay between promotor and structural gene variants control basal level of mannan binding protein. *J Immunol* 1995; 155: 3013–20.
6. Snowden N, Stanworth S, Donn R et al. Mannose-binding protein genotypes and recurrent infection. *Lancet* 1995; 346: 1629–30.
7. Ollier WER, MacGregor A. Genetic epidemiology of rheumatoid disease. In: Saklatvala J, Walport MJ, eds. *Immunology of rheumatoid disease*. *Br Med Bull* 1995; 51: 267–85.
8. Parekh RB, Dwek RA, Sutton BJ et al. Association of rheumatoid arthritis and primary osteoarthritis with changes in the glycosylation pattern of total serum IgG. *Nature* 1985; 316: 452–7.
9. Ezekowitz RA. Agalactosyl Ig and mannose-binding proteins: Biochemical nicety or pathophysiological paradigm? *Nature Med* 1995; 1: 207–8.
10. Malhotra R, Wormald MR, Rud PM, et al. Glycosylation changes of IgG associated with rheumatoid arthritis can activate complement via mannose-binding protein. *Nature Med* 1995; 1: 237–43.
11. Turner MW. Mannose-binding lectin: the pluripotent molecule of the innate immune system. *Immunol Today* 1996; 17: 532–40.
12. Rademacher TW, Williams P, Dweek RA. Agalactosyl glycoforms of IgG autoantibodies are pathogenic. *Proc Natl Acad Sci USA* 1994; 91: 6123–7.
13. Thomson C. Protein proves to be a key link in innate immunity. *Science* 1995; 269: 301–2.
14. Stanworth JS, Donn PR, Hassall A, Dawes P, et al. Absence of an association between mannose-binding lectin polymorphism and rheumatoid arthritis. *Br J Rheumatol* 1998; 37: 186–88.
15. Kilpatrick DC. Mannan binding protein in sera positive for rheumatoid factor. *Br J Rheumatol* 1997; 36: 207–9.
16. Arnett FC, Edworthy SM, Bloch DA et al. The American Rheumatism Association 1987 revised criteria for the classification of rheumatoid arthritis. *Arthritis Rheum* 1988; 31: 315–24.
17. Mijušković VZ, Zgradić I, Nikolajević R, Popović M, Nedović J, Lazarević M, Stojanović J. Rheumatoid factor isotypes as the prognostic markers in Rheumatoid arthritis. *Proceedings of the 10th International Congress of Immunology 1-6 Nov. 1998; New Delhi, India, Monduzzi Editore Sp. A – Bologna, 1251–1255.*
18. Mijušković VZ. Significant parameters of humoral immunity in sera and synovial fluid for the course and activity of RA. *Dissertation. Military Medical Academy, Belgrade, 2002.*
19. Mollines TE, Lea T, Melbye OJ, Pahle J, Grand O, Harboe M. Complement activation in rheumatoid arthritis evaluated by C3dg and the terminal complement complex. *Arthritis Rheum* 1986; 29: 715–26.

Received: November 25, 2002

Accepted: January 15, 2003