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Post-streptococcal glomerulonephritis: Immunopathological aspects and perspectives for *in* s*ilico* experiments

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Acute post-streptococcal glomerulonephritis (APSGN) is an acute inflammatory disease involving the kidney glomeruli that causes diffuse proliferative lesions, oropharyngeal or skin infections. Clinically, it is a typical example of nephritic syndrome, with the occurrence of hematuria, edema, and hypertension, which may evolve to acute renal failure. Besides the consultation of important textbooks in the field, this article was based on a literature review using DeCS (Health Sciences Descriptors) web tool, which included searches in the databases LILACS (Latin American Literature in Health Sciences), PubMed (U. S. National Library of Medicine), and SciELO (Scientific Electronic Library Online). This simulator was constructed with the main objective of investigating autoimmune diseases, and it has been consistent with results presented in the literature. For the investigation of APSGN, our work comprises the extension of *AutoSimmune* to include important environments such as the kidney tissue as well as agents representing *S. pyogenes* and also the complement system. There are still many unclear points relevant to medical treatment of APSGN. In this context, *in silico* research might open important new possibilities of how to observe and test the immune response to *S. pyogenes* and their multiple interactions and outcomes. This review presents the current comprehension on APSGN and provides an important basis for the development of the proposed computational approach for the disease scrutiny.

Key words: Autoimmune diseases, AutoSimmune, Glomerulonephritis, in silico, Streptococcus pyogenes.

INTRODUCTION

Some species of bacteria usually belong to the normal microbiota of the respiratory, gastrointestinal, and genital tracts. In certain contexts, they can cause sore throat, skin infections, neuroinfections, sepsis, endocarditis, and pneumonia, which are closely related to the direct action of microorganisms. Furthermore, there are disorders related to the production of toxins – which is the case of scarlet fever – and also the emergence of autoimmune post-infectious syndromes and toxic shock syndrome, especially rheumatic fever and post-streptococcal glome-rulonephritis (APSGN) (Baldy, 1997).

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PSGN is an acute inflammatory disease involving the kidney glomeruli that causes diffuse proliferative lesions (Alpers, 2010), and is a consequence of oropharyngeal or skin infections. Clinically, it is a typical example of nephritic syndrome, with the occurrence of hematuria, edema, and hypertension, which may evolve to acute renal failure. Nephritic syndrome by APSGN generally appears one to four weeks after infection of the pharynx (pharyngitis) or skin (impetigo), usually attacking children aged between six and ten years. It should be highlighted the immunological mediation by immune complexes (IC) (Alpers, 2010; Bisno, 2000; Bisno and Stevens, 2000). Several cationic antigens - such as the NAPIr, SpeB, zSpeB – may be related to nephritogenicity (Bisno and Stevens, 2000; Batsford et al., 2005). Laboratory tests indicated low levels of complement in serum due to consumption and increased anti-Streptococcal antibodies – the anti-streptolysin O –, which confirms the previous occurrence of infection by this agent. Light microscopy showed granular immune deposits, confirming the IC mechanism. The glomerulus is enlarged and shows hypercellular shape, due to infiltration, proliferation and, in severe cases, formation of crescents. The swelling and proliferation of endothelial cells lead to the obliteration of capillary lumens and interstitial edema (Alpers, 2010). Over 95% of the patients recover with conservative therapy. Still, prognosis is not good in cases of prolonged and significant proteinuria and abnormal glomerular filtration rate (GFR).

The present work aims to present a review on APSGN etiopathogenesis, highlighting the research prospects for the nosological entity, providing, additionally, a direction for a computer modeling approach to study the disease.

METHODS

This article was based on a wide literature examination. The search strategy (Table 1) in DeCS (Health Sciences Descriptors) included the terms "autoimmunity", "immunology", "glomerulonephrytis", "kidney", and *Streptococcus pyogenes*, in accordance with the following strategies:

Strategy 1 – Streptococcus pyogenes + glomerulonephritis;

Strategy 2 – *Streptococcus pyogenes* + autoimmunity + kidney;

Strategy 3 – Streptococcus pyogenes + immunology + kidney.

The search for articles was conducted in the databases of LILACS (*Latin American Literature in Health Sciences*), PubMed (*U. S. National Library of Medicine*), and SciELO (*Scientific Electronic Library Online*), additionally to textbooks. Complete publications – In English or Portuguese – were selected according to the content of their title or abstract, i.e., if there was reference to the post-infectious or post-streptococcal glomerulonephritis. Table 1 (page 21).

The final selection for the articles was performed as follows: reading of the title and abstract as well as detailed assessment of the article. Publications that did not refer to human beings were excluded. The preference was given to those that prioritized aspects related to etiopathogenesis and experimentation.

RESULTS AND DISCUSSION

POST-STREPTOCOCCAL GLOMERULONEPHRITIS: EPIDEMIOLOGICAL ASPECTS AND RELEVANCE FOR PUBLIC HEALTH

The APSGN epidemiological pattern has changed significantly over the last three decades. Furthermore, this disease has been rarely found in developed nations. However, it still occurs in developing countries, where 9.5 to 28.5 new cases per 100,000 people are recorded per year (Rodríguez-Iturbe and Musser, 2008).

The disease has almost disappeared from central Europe, where the elderly population is most affected. In Italy, occurrence rate in people over 60 is 0.9 patients per million inhabitants, while for people aged below 60 years, it is 0.4 (Rodríguez-Iturbe and Musser, 2008; Montseny et al., 1995; Coppo et al., 1998). Still, APSGN is relatively common in rural and Aboriginal communities, and places presenting lower social-economic development, reaching 21% (4.6-51.6%) of the children with kidney failure admitted to hospitals (Rodríguez-Iturbe and Mezzano, 2005). In India, for example, morbid condition represents 73% of cases of acute glomerulonephritis in the elderly (Pakash et al., 2001). Carapetiset and colleagues Carapetis et al. (2005), in a study with 11 populations, reported that the incidence in children - in the least developed countries - was around 24.3 cases per 100,000 people a year. For sick people aged over 15 years, two cases per 100,000 people a year were observed. According to Italian data, in developed countries, the APSGN drops to 0.3 cases per 100,000 people per year. Worldwide, out of the 472,000 APSGN cases per year, about 456,000 are found in developing countries (Rodríguez-Iturbe and Musser, 2008; Carapetis et al., 2005).

Table 2, adapted from Rodríguez-Iturbe and colleagues (Rodríguez-Iturbe and Musser, 2008), compares the APSGN load in developing countries, based on two studies. Although using different methodologies, this research suggests two close estimated incidences. Table 2 (page 22)

In Brazil, the occurrence of colonization by S. pyogenes is described as usual in groups of patients with persistent tonsillitis, aging between 3 and 16 years, who are susceptible to infection and complications. Amelia and colleagues (Maciel et al., 2003) analyzed the prevalence of Group A β-hemolytic S. pyogenes in a sample group of 735 students from the city of Recife. Prevalence of 0.8% of the agent was observed in asymptomatic carriers. A similar rate was reported in Switzerland in young adults and 6% in school children (Maciel et al., 2003; Hoffmann, 1985), and still 7% and 18.8% among school children in the United States and India, respectively (Hoffmann, 1985; Karoui et al., 1982; McMillan et al., 1986). It is noteworthy that 30% of acute cases of pharyngotonsillitis are caused by Streptococcus pyogenes (McMillan et al., 1986).

ETIOLOGY: Streptococcus pyogenes

The *Streptococcus pyogenes* is one of the important species of the genus *Streptococcus*. *S. pyogenes* is a facultative anaerobe bacterium that grows in crops, in pairs or small chains (Bisno and Van de Rijn, 2000), and

Table 1 – Strategy for literature search.

	BASES CONSULTED*			ARTICLES SELECTED ACCORDING TO THE
SEARCH STRATEGIES	LILACS	PUBMED	SCIELO	INCLUSION CRITERIA (total number)**
Strategy 1 Streptococcus pyogenes + glomerulonephritis	10	342	2	
Strategy 2 Streptococcus pyogenes + autoimmunity + kidney	0	1	0	30
Strategy 3 Streptococcus pyogenes + immunology + kidney	0	54	0	

* In SCIELO and LILACS, the search was conducted with the descriptors written in Portuguese; PUBMED descriptors were written in English. ** Some selected articles were cited in both SCIELO and PUBMED; thus, we prefer to mention here just the total number of the selected articles.

produces β -hemolysis, classified in Lancefield group A (Lancefield, 1933). It should be also said that it is a nutritionally fastidious germ that requires the addition of blood to the media.

PATHOPHYSIOLOGICAL ASPECTS OF POST-STREPTOCOCCAL GLOMERULONEPHRITIS

Based on the observations of Clemens von Pirquet, from the early twentieth century, it was possible to correlate streptococcal infection prior to the onset of the manifestations of renal problems. At the time, von Pirquet proposed a process of "altered immune reactivity" (Rodríguez-Iturbe and Batsford, 2007). Later, Seegal and Earle (1941) introduced the concept of nephritogenic and rheumatogenic strains of streptococci, and reported that the most serious complications of infection - rheumatic fever and glomerulonephritis - could not coexist in the same patient. Today, it is known that this observation is not true, since these diseases can coexist, albeit rarely (Tokura et al., 2008). It should be highlighted that the Lancefield strains of the types M 1, 2, 4, 12 and 25 recovered from the upper airways were nephritogenic, as well as the strains M 49, 55, 57 and 60, which are associated with impetigo (Rodríguez-Iturbe and Batsford, 2007). Then, these strains were sorted into classes I and II, respectively, as rheumatogenic and nephritogenic, according to the result of the action by lipoproteinase on the M protein.

The pathogenic mechanisms of the disease are still under investigation. The first hypotheses formulated deserve special attention, even after the emergence of molecular biology. There is almost a consensus that the pathological base of the lesion happens due to the local inflammatory response related to the deposition of immune complexes. Doubtlessly, immune complexes can be formed prior to deposition or originate from *in situ* reaction with "planted" antigens or by cross-reactivity to autologous antigens (Alpers, 2010; Rodríguez-Iturbe and Musser, 2008; Rodríguez-Iturbe and Batsford, 2007). Regarding autoimmunity, an important role is played by the cell injury immune-mediated by cells and by the complement activation, with directly chemotactic and cytolytic action. Investigations with knockout mice for complement proteins indicated the importance of leukocyte Fc receptors and kidney cells themselves as mediators of the process, showing that injuries are not exclusively mediated and amplified by the complement binding (Oda et al., 2010; Burova et al., 2012; Zipfel et al., 2006).

Under light microscopy, the lesions show leukicyte infiltration and proliferation of endothelial and mesangial cells (Oda et al., 2010; Burova et al., 2012). Regarding the distribution of glomerular deposits, some conjectures can be made about its location, which can be: (1) in the mesangium, with neutral charges; (2) in the subendothelial space, highly anionic; (3) and/or in the subepithelial space, highly cationic.

When immune complexes are very large, they lose the ability to enter the glomerular basement membrane (GBM). They are eliminated by the mononuclear phagocyte system and are not nephritogenic. The pattern of deposits can be distinguished as follows: (i) granular, characteristic in nephritis by circulating immune complexes, and in situ, and (ii) linear, characteristic of the classic anti-GBM disease. Once deposited, the immune complexes can be degraded by infiltration of inflammatory macrophages/monocytes. Thus, the reaction can be contained, as in cases of short and limited antigenic exposure (Rodríguez-Iturbe and Musser, 2008; Rodríguez-Iturbe and Batsford, 2007; Oda et al., 2010; Burova et al., 2012). Repeated cycles of exposure and response lead to the most chronic membranopro-

Study	Population	Estimate of annual cases	Estimate of annual incidence (cases per 100,000 people)
Carapetis <i>et al.</i>	4,876,709,000	456,000	9.3
Rodriguez-Iturbe et al.	29,967,989	2,850 to 8,550	9.5 to 28.5

 Table 2. APSGN load in developing countries: a comparison between studies.

Adapted from Rodriguez-Iturbe et al., 2006.

liferative GN type. Damage to the epithelial cells occurs by antibodies targeted to them, as well as by toxins and cytokines. Such disturbances lead to changes in the visceral epithelial cells, with destruction of the pedicular processes, detachment of MBG cells and, ultimately, changes in the filtration process (Alpers, 2010; Rodríguez-Iturbe and Batsford, 2007; Uchida et al., 2011).

These are some of the effectors of injury: (1) neutrophils and monocytes attracted by chemotaxis, resulting from complement activation - especially C5a and Fc-mediated activation; these cells release proteases which degrade GBM, reactive oxygen species, metabolites of arachidonic acid, contributing to TGF reduction. There is still infiltration of macrophages, T lymphocytes, and NK cells (natural killer); (2) platelets, which release eicosanoids and growth factors (PDGF); (3) resident glomerular cells – mesangial – also release inflammatory mediators, cytokines, chemokines, growth factors, endothelin, which, even in the absence of immune infiltration, can start inflammation; (4) chemotactic components of the complement and its final via - the membrane attack complex (MAC) -, which causes cell lysis and stimulates mesangial cells to produce other mediators; it is worth noting that the action of MAC itself is enough to trigger proteinuria; (5) All mediators produced by infiltration, including monocyte chemoattractant protein 1 and CCL5, which are lymphocyte attractant. The growth factors PDGF, TGFand Fibroblast growth factor are related to mesangial cell proliferation, extracellular matrix deposition and hyalinization (Alpers, 2010).

Once the injuries are started, mechanisms emerge and culminate with progression. It starts with adaptive change – characterized by compensatory hypertrophy – of the remaining glomeruli. Any renal disease which destroys the nephrons at a GFR rate of 30% to 50% of the normal rate will progress gradually to renal failure (Alpers, 2010). This is associated with hemodynamic changes, increased arterial blood flow, filtration and transcapillary pressure, and often with systemic hypertension. This context is believed to lead to injury of endothelial and epithelial cells, accumulation of protein in the mesangial matrix, proliferation with progressive injury. TGF-□ causes, chronically, sclerosis and loss of the proliferative ability of

podocytes (Alpers, 2010; Rodríguez-Iturbe and Batsford, 2007; Uchida et al., 2011).

Homo sapiens sapiens / Streptococcus pyogenes INTERACTION: THE NUANCES OF AUTOIMMUNITY

After the clear definition of the relation between S. pyogenes and APSGN, it was conducted a search for bacterial antigens that would be the main originators or triggers of the disease. Indeed, until now, there are numerous speculations and uncertainties relating to such structures. In the 60s and 70s, much was discussed about the deposition of circulating immune complexes. In fact, in 1976, the possibility of in situ reaction was also questioned, due to the impossibility - according to the ideas prevailing at the time - of the existence of subepithelial deposition by preformed complexes. This concept was reviewed by Vogt and colleagues (Vogt et al., 1990), who highlighted the possibility of translocation, dependent on size and charge of the compound by glomerular membrane (Rodríguez-Iturbe and Batsford, 2007). Complement activation is recognized as a key feature in the immunopathogenesis of the disease, with emphasis on the different mechanisms by which it occurs. The immunoglobulin-binding proteins on the bacterial surface attach to C4BP (C4-binding protein), thus affecting the activation classical pathway; there is predominance of activation by the alternative pathway. Besides, there is the lectin pathway initiated by the recognition of glucosamine residues of the bacterial wall by mannose binder. In the pathogenic process, much has been discussed about the cellular immune mechanisms. Rastaldi and colleagues (Rastaldi et al., 1996) related the glomerular intensity of and tubulointerstitial macrophagocytic infiltration with the expression of adhesins and proteinuria.

Similarly, during the course of inflammation, it has been observed the occurrence of increased circulating levels of IL-6, IL-8, TNF-a, monocyte chemotactic protein and cellular products (Soto et al., 1997).

There is a long search for antigens of *S. pyogenes* that can mimic glomerular structures.

Kefalideset and colleagues (Kefalides et al., 1986) described antibodies against laminin and collagen in some sick people. Furthermore, it is known that protein M12 epitopes are shared by the glomerular basement membrane, vimentin and mesangial proteins, but Kraus and Beachey (Kraus and Beachey, 1988) were responsible for the identification of epitopes with autoimmune potential.

McIntosh et al. demonstrated, also in the 70's, that group Α streptococci type M12 possessed а neuraminidase that altered the structure of IgG by desialyzation, which allowed an anti-IgG response that would lead to glomerular injury (McIntosh et al., 1978; Rodríguez-Iturbe et al., 1980; Marin et al., 1995). This is corroborated by the fact that anti-IgG deposits were described in 19 of the 22 kidneys biopsied at the time (McIntosh et al., 1978; Rodríguez-Iturbe et al., 1980). The ability to produce this enzyme is shared by the rheumatogenic strains. In the desialyzation mechanism, it is known that there is predominantly a IgG binder in the wall of the microorganism, with high avidity for the Fc portion.

There is also a surface protein H which, in interaction with the Fc of IgG, is released into the serum by action of SpeB protein. In serum, this protein H causes complement activation with dose-dependent cleavage of C3, while the H-IgG complex allows anti-IgG response once more.

The main antigenic structures investigated are presented below in order to establish the molecule of streptococcal origin involved in the events of autoimmunity.

MOLECULAR STRUCTURES ALLEGEDLY INVOLVED IN THE DEVELOPMENT OF **APSGN**

The molecules probably involved in the immunopathogenesis of APSGN are the *M* protein, streptokinase, NAPIr (Nephritis-Associated Streptococcal Plasmin Receptor) and SpeB (Streptococcal pyrogenic exotoxin B).

M Protein

This peptide is an important anti-phagocytosis virulence factor to distinguish more than 150 different types of *Streptococcus* of group A. Currently, M 1, 2, 4, 12, 25, 49, 55, 57, 60 are recognized as having nephritogenic potential. M protein can be located in the glomeruli and it must be highlighted that glomerulus lesions can be induced by repeated injections of M protein either alone or in combination with fibrinogen (Rodríguez-Iturbe and Batsford, 2007). Notice that if the M protein actually had a central role in the nephritogenic process, then APSGN recurrence, which is still rare, would be even less likely, because the pattern of response to ubiquitinated antigens generates long-term immunity, which does not occur by exposure to M protein (Rodríguez-Iturbe and Batsford, 2007; Calish and Vaz, 2009). Furthermore, it is known that other groups – not the group A – of *Streptococcus* have nephritogenic potential, including *Streptococcus zooepidemicus*, of Group C. However, cross-reactivity between epitopes of M protein and glomerular components has been demonstrated.

In recent research with animal experiments, published by Burova and colleagues (Burova et al., 2012; Burova et al., 2003), it was observed the effect of the IgGFc binding protein, which belongs to the M protein family. In the strain M22, used in that test, it was observed that the deletion of a gene that encoded two variables of that IgG binder maintained the nephritogenicity of the agent, but the deletion of both genes annihilated its potential. In this investigation, for the purpose of comparison, IgGFc ligands of other bacterial isolates, Staphylococcus aureus protein A and protein G from group G of Streptococcus, were used and none of them caused renal injury. On the other hand, circulating anti-IgG antibody was found in seven of the ten models with experimental lesions. Finally, this experiment demonstrated that therapy with Fc fragments at the beginning of renal injury prevents the advance of glomerulonephritis also due to reduced stimulus to anti-IgG production and its consequences. Therefore, similarity between the injury caused by the bacterium and the isolated component studied reveals that this would be a primary factor causing glomerulonephritis, which contradicts some hypotheses primary other antigens about also accepted. Nevertheless, it is known the existence of others IgG binding proteins, such as the protein H member of M-like protein family described above. All of them and their contribution to renal injury are important subjects for investigation.

Streptokinase

The presence of this compound in the list of possible nephritogenic antigens is controversial. Its inclusion was due to its observation in biopsy, in selective inoculation experiments and in comparison to nephritogenicity loss after gene deletion (Norstrand et al., 2000). However, later on, the same group of researchers decided to exclude it. Nowadays, it is known that streptokinase converts plasminogen into plasmin, which could cause direct degradation of extracellular matrix - for example. fibronectin activation and laminin of metalloproteinases, collagenases, complement, and coagulation factors (Castellino and Bajaj, 1997; Grella and Castelino, 1997). All this etiopathogenic scenario triggers local injury and predisposes to deposition of immune complexes.

NAPIr (Nephritis-Associated Streptococcal Plasmin Receptor)

NAPIr is a structure homologous to GAPDH (glyceraldehyde-3-phosphate dehydrogenase), which is

capable of binding to plasmin, keeping the proteolytic activity of the latter, preventing the physiological inhibitory action of alpha₂-antiplasmin (Oda et al., 2008). Although presenting the same characteristic of plasmin activator as *Streptococcal pyrogenic exotoxin B* (discussed below), its amino acid sequences as well as structural and functional properties are quite different. Furthermore, NAPIr is isolated from the cytoplasm of the agent, while SpeB is isolated from the external environment.

In all the patients, deposits of glomerular NAPIr are found from the initial stage of the disease, mainly in neutrophils, endothelial, and mesangial cells. Only the pattern and intensity differ, according to Oda and colleagues (Oda et al., 2010). They differ in the location of deposits of C₃ and IgG, but coincide with the locations of active plasmin. There is also similarity when compared with SpeB by double immunofluorescence. On the other hand, in the same work of Oda et al., no NAPIr mark is found in any of the sick people with rapidly progressive GN. Double staining for NAPIr and collagen IV revealed prevalence of the former inside the glomerular tuft, with lower marking in the parietal cells of Bowman's capsule. This is consistent with the predominance of endocapillary glomerular inflammation in the disease (Yoshizawa et al., 2004).

Regarding the preferential localization of NAPIr in neutrophils and the consequent hyper proteolytic activity observed, it is assumed that this association plays a role in the *in situ* destruction (Yamakami et al., 2000). The urokinase-type plasminogen activator receptor, a neutrophilis CAPDH receptor, is one of the mechanisms that explains this interaction. There is also the possibility of phagocytosis of NAPIr as antigen, which does not occur with macrophages, by a mechanism still unclear.

SpeB (Streptococcal pyrogenic exotoxin B)

The exotoxin SpeB - a cationic cysteine proteinase derivative from zymogen zSpeB - is produced in vivo by Streptococcus during the course of infection, behaving as a plasmin ligand receptor. Evaluation, by Bastford and colleagues (Batsford et al., 2005), of NAPIr and SpeB in serum and biopsy of patients from different countries and continents revealed deposits of SpeB in 12 out of the 17 biopsies and anti-SpeB antibodies in all sera investigated. The complement deposits were co-located with SpeB, while the anti-NAPIr antibodies were described in only five out of the 47 samples, with a single occurrence in biopsies. In any case, this antigen and the previous one are found in all isolates of S. pyogenes, but only a small proportion of those infected develop nephritis. In addition, Streptococcus zooepidemicus does not have the gene encoding SpeB. Even so, S. zooepidemicus was associated with an epidemic in Brazil (1988) (Sesso and Pinto, 2005). Another mechanism of action of SpeB, described by Wei and colleagues (Wei et al., 2005), showed that microbial pathogens often exploit human complement regulatory proteins such as factor H (FH) and factor H-like protein 1 (FHL-1) for immune evasion (Thurman and Holers, 2006; Vernon et al., 2012). Notice that Fba is an FH and FHL-1 binding protein expressed on the surface of the human bacterium Streptococcus pathogenic pyogenes, а common agent of pharyngeal, skin, and soft-tissue infections. Fba has been shown to contribute to phagocytosis resistance, intracellular invasion, and virulence in mice. Analysis of a speB confirmed that SpeB accounts for the loss of Fba from the cell surface, suggesting that the protease may modulate FH and FHL-1 recruitment during infection (Norstrand et al., 1999).

SCIENTIFIC PERSPECTIVES: IN SILICO EXPERIMENT

Experimental medicine has new alternatives for investigating the immune system, besides the animal research model, among which the *in silico* process of experimentation, namely, the use of computer simulation, has demonstrated to be very promising (Folcik et al., 2007; Macal and North, 2009; Li et al., 2009; Possi et al., 2011; Possi et al., 2010; Bastos et al., 2011; Silva et al., 2012; Siqueira-Batista et al., 2012; Gomes et al., 2012).

Animal experiment models for APSGN have not been considered adequate for a long time, perhaps due to the specificity of the encounter between *S. pyogenes* and *H. sapiens sapiens*. Nordstram and colleagues (Norstrand et al., 1999) developed a model using a steel cage and osmotic pumps, subcutaneously, in mice and rabbits. The device was loaded with bacterial isolate associated with nephritis, which allowed it to cause renal injury. Nowadays, the humanized mice have been successfully used to simulate some biological interactions (Sun et al., 2004; Poyart et al, 2005).

The *in silico* simulation of human immune system (SI) may greatly help the research work on APSGN (Gomes et al., 2012). SI is quite complex and its complete comprehension is considered one of the most challenging issues in biology (Folcik et al., 2007; Li et al., 2009; Possi et al., 2011; Bastos et al., 2011). However, knowledge of its mechanisms is fundamental to the advancement of several areas of science, including medicine and computer science (Macal and North, 2009; Li et al., 2009; Possi et al., 2011; Possi et al., 2010; Bastos et al., 2011), which makes this biological system a mandatory field of research.

The methods used for SI *in silico* experiments include the multi-agent systems (MAS) approach, based on autonomous agents, and has been successfully applied in many complex systems, allowing the assessment of hypotheses of interaction between cells and the behaviors emerging from these interactions claim that this is possible because MAS allows exploring the complex and deterministic macroscopic behavior that emerges from stochastic microscopic interactions (Li et al., 2009). For this reason, MAS is considered by many authors as the best approach for modeling SI. Its main disadvantage, however, is the high computer cost when a large number of agents is used (Li et al., 2009). It must be observed that when this approach is applied to system modeling, agents, i.e., the level of granularity, must be defined, and then rules concerning their behavior must be established. As a result, the system behavior emerges from the interaction among agents as well as between agents and the environment. Notice that the granularity level should be established in such a way that the obtained model can suitably represent the system in study and, at the same time, the resulting computational complexity does not make the investigation infeasible. In other words, it is necessary to represent only the entities that are essential for a good representation of the system (Li et al., 2009).

BIS, also known as "The Basic Immune Simulator" (Folcik et al., 2007), is among the agent-based SI simulators. Its model studies the interactions between cells of innate immunity and adaptive immunity cells. *AutoSimmune* (Possi et al., 2011), which is an extension of BIS, is another important tool for SI simulation and was initially aimed to test hypotheses related to autoimmune diseases. Since then, *AutoSimmune* has been extended to include important SI mechanisms such as the simulation of the mast cell behavior (Silva et al., 2012).

We are working with the prospect of in silico investigation of the APSGN pathophysiology. For this purpose, the AutoSimmune simulator developed by Possi et al. (Possi et al., 2011) is being used as the basis for our simulation (Silva et al., 2012). For the recognition of S. pyogenes, which is an initial step towards the investigation of post-streptococcal glomerulonephritis, it is necessary to include a kidney tissue environment in the model, simulating the kidneys, as well as agents representing the complement system in its activation pathways, especially considering the classical pathway which is activated in the presence of immune complexes (IgG or IgM linked to specific antigens) until the formation of the membrane attack complex (MAC) that causes the lysis of antigen. Furthermore, the molecules SpeB (Streptococcal pyrogenic exotoxin B) and NAPIr (Nephritis-Associated Streptococcal Plasmin Receptor), probably involved in the immunopathogenesis of the APSGN, are important targets to be investigated and modeled with respect to their behavior and interactions. Notice that Oda and colleagues (Oda et al., 2010) reported that NAPIr glomerular deposits are found in the earliest stages of the disease, mainly in neutrophils, endothelial, and mesangial cells - with variations only in pattern and intensity. Besides, according to Bastford and collaborators, in the evaluation of the biopsy of patients from different countries and continents, NAPIr and SpeB are found in all isolates of S. pyogenes.

At the end of the work, we intend to validate the simulator, comparing it with the findings in literature, to evidence the consistent behavior of the model to be presented.

CONFLICT OF INTEREST.

Work done in the Department of Medicine and Nursing and the Department of Computer Science, Universidade Federal de Viçosa (UFV). There is no conflict of interest.

CONCLUSION

After an exhaustive search for a determinant antigen, there is a current understanding that the participation of the host – H. sapiens sapiens – in the course of infection - in strict correlation with renal involvement - is fundamental for understanding APSGN etiopathogenesis. Undoubtedly, much remains to be clarified about the emergence of the morbid condition. Currently, S. pyogenes is one of the best known organisms in the genomic field, which provides a large number of scientific perspectives. Hitherto, the deposition of immune complexes has been adopted as the starting point for the disease and the recruitment of several mechanisms of injury. The best diagnostic suggestion is still given by high antibody titers against SpeB/zSpeB or NAPIr. In this context, in silico investigations may offer new possibilities for observing and testing the immune response to S. pyogenes and its multiple interactions and outcomes.

CONTRIBUTION OF AUTHORS

CA Bastos and EP Santos wrote the first version of the article, under the guidance of AP Oliveira. AP Gomes, LA Santana and R Siqueira-Batista designed search strategy. A detailed review of article was performed by AP Gomes, AP Oliveira, FR Cerqueira, LA Santana, RS Ferreira and R Siqueira-Batista.

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