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Blockade of gC1qR/p33, a receptor for C1q, inhibits adherence of *Staphylococcus aureus* **to the microvascular endothelium**

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Abstract

Endovascular infections with *Staphylococcus aureus* (*S. aureus*) are associated with high mortality. gC1qR/p33 (gC1qR), a receptor for the complement component C1q expressed on endothelial cells, interacts with protein A of *S. aureus* and gC1qR blockade reduces *S. aureus* colonization during infective endocarditis. The aim of this study was to analyze *in vivo* whether this observation is due to a decreased interaction of *S. aureus* with the microvascular endothelium. A dorsal skinfold chamber was prepared in Syrian golden hamsters, which were treated with the monoclonal antibody (MAb) 74.5.2 directed against gC1qR or vehicle. The interaction of fluorescein isothiocyanate (FITC)-labeled *staphylococci* and leukocytes with the endothelium was analyzed under physiological conditions as well as after TNF-α-induced inflammation using intravital fluorescence microscopy. Administration of MAb 74.5.2 significantly reduced adherence of *S. aureus* to the endothelium in untreated and TNF-α-exposed tissue. In addition, we could demonstrate *in vitro* that *S. aureus* adherence to human endothelial cells was inhibited by MAb 74.5.2. Blockade of gC1qR did not affect leukocyte-endothelial cell interaction. In conclusion, our findings indicate that immunological inhibition of gC1qR may be therapeutically used to decrease the interaction of *S. aureus* with the microvascular endothelium.

Keywords

Staphylococcus aureus; endothelium; gC1qR/p33; monoclonal antibody; adherence; infective endocarditis; dorsal skinfold chamber; intravital fluorescence microscopy

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Introduction

Staphylococcus aureus (*S. aureus*) is a highly virulent pathogen causing a wide array of community- and hospital-acquired infections (Lowy, 1998). Today, antimicrobial resistance is a major issue in the treatment of invasive *S. aureus* disease, as methicillin-resistant *S. aureus* (MRSA) is pandemic worldwide (Westh et al., 2006; Wisplinghoff et al., 2004) and as the advent of community-acquired MRSA in many parts of the world now represents an additional serious public health problem (Boyle-Vavra and Daum, 2007; Fridkin et al., 2005; Stam-Bolink et al., 2007).

S. aureus-induced endovascular infections are of particular importance as they are associated with a high mortality rate (Madani, 2002; Yeaman and Bayer, 2000). Pathogenetically, endovascular *S. aureus* disease is a multistep process involving primary bacterial adherence to the endothelium, establishment and persistence of the infection, bacterial proliferation usually associated with tissue damage and finally dissemination of septic emboli and organic spread (Foster and McDevitt, 1994; Hartleib et al., 2000; Herrmann et al., 1988; Moreillon et al., 2002).

During the last few years, it has been reported that *S. aureus* expresses a number of bacterial cell wall-anchored proteins, which mediate bacterial adherence to host cells as well as extracellular matrix (ECM) components. These include FnBPA and FnBPB, which recognize fibronectin and fibrinogen (Foster and Hook, 1998; Wann et al., 2000), or ClfA and ClfB binding to fibrinogen (McDevitt et al., 1994; Ni et al., 1998). Another important adherence protein of *S. aureus* is protein A (Hartleib et al., 2000), which has been shown to bind to von Willebrand factor (vWF) and to gC1qR/p33 (gC1qR) (Hartleib et al., 2000; Nguyen et al., 2000).

The cellular protein gC1qR is expressed on activated platelets (Peerschke et al., 2003) and on endothelial cells (Guo et al., 1999). Furthermore, gC1qR is present in the extracellular matrix (Hasan et al., 1998) and circulates in a soluble form in the blood plasma (Peterson et al., 1997; van den Berg et al., 1997). Interestingly, recent studies have demonstrated that gC1qR enhances *S. aureus*-fibrinogen interactions and that administration of monoclonal antibodies (MAbs) directed against gC1qR inhibit *S. aureus* tissue colonization during infective endocarditis (Peerschke et al., 2006). Accordingly, gC1qR may play a crucial role in the pathogenesis of *S. aureus* endovascular infections and, thus, may represent a novel target for the establishment of therapies preventing colonization and subsequent infection.

To test this hypothesis, we used the dorsal skinfold chamber model in Syrian golden hamsters, which allows to analyze *in vivo* the interaction of fluorescein isothiocyanate (FITC)-labeled *staphylococci* with the microvascular endothelium by means of intravital fluorescence microscopy (Kerdudou et al., 2006; Laschke et al., 2005a; Roller et al., 2008). Using this animal model, we were able to study the effects of gC1qR blockade on *S. aureus*endothelial cell interaction, which represents the initial step in the pathogenesis of *S. aureus*associated endovascular diseases.

Materials and Methods

Animals

The experiments were conducted in accordance with the German legislation on protection of animals and the *NIH Guide for the Care and Use of Laboratory Animals* (Institute of Laboratory Animal Resources, National Research Council, Washington, USA), and were approved by the local governmental animal care committee.

Eight- to ten-week-old male Syrian golden hamsters with a body weight of 60–80g were used for the study. The animals were housed one per cage and had free access to tap water and standard pellet food (Altromin, Lage, Germany) throughout the experiment.

Preparation of the dorsal skinfold chamber

Dorsal skinfold chambers were prepared in Syrian golden hamsters as described previously in detail (Menger et al., 2002). In brief, animals were anesthetized using sodium pentobarbital anesthesia (50 mg/kg body weight i.p.) and two symmetrical titanium frames were implanted on the extended dorsal skinfold of the hamsters, so that they sandwiched the double layer of skin. One layer of skin was then removed in a circular area of ~15mm in diameter, and the remaining layers (consisting of striated skin muscle, subcutaneous tissue and skin) were covered with a removable cover slip incorporated into one of the titanium frames (Figs. 1A and B). After the procedure, the animals were allowed to recover from anesthesia and surgery for at least 48h before the microcirculatory analyses.

Preparation of bacteria

For this study, we used the wild-type strain *S. aureus* Cowan 1 (ATCC 12598). Bacteria were prepared as described previously (Laschke et al., 2005a). To obtain 6h (late exponential phase) bacteria, *S. aureus* were cultured in 10 mL of Muller Hinton broth under constant rotation and then washed. This suspension yielded approximately $5 \times 10^8 - 1 \times 10^9$ cfu/mL. The exact determination of each experiment's bacterial number was performed by viable cell counting after vigorous vortexing of the washed, resuspended microorganisms. Subsequently, the bacteria were stained with FITC (20mg/mL; Sigma, Deisenhofen, Germany), washed again and fixed in 1% formaldehyde for a time period of 3h. Bacteria were fixed in order to i) ascertain synchronization of bacteria in the late exponential phase, which has been shown in previous studies to be optimal for the *in vivo* interaction of bacteria with the endothelium (Kerdudou et al., 2006), and ii) to exclude surface adhesinindependent effects on endothelial and blood cells, which might be mediated by a panoply of factors released from viable organisms. The fixed FITC-labeled *staphylococci* were then washed three times in PBS and the inoculum was adjusted to 10^9 cells/mL sterile PBS / 1% BSA (extrapolated from the number of viable counts prior to fixation). The sample was stored for $2-3$ days at 4° C in the dark p rior to application, and 1 mL of this suspension was injected into each animal.

Intravital fluorescence microscopy

For the *in vivo* microscopic analysis of the microcirculation, the animals were anesthetized and a fine polyethylene catheter (PE10, 0.28mm internal diameter) was inserted into the A. carotis for application of bacteria and fluorescent dyes. Subsequently, the hamsters were put in lateral decubital position on a Plexiglas pad and the dorsal skinfold chamber preparation was attached to the microscopic stage. Intravital fluorescence microscopy was performed after i.a. injection of 10^9 FITC-labeled bacteria. At the end of the experiment, i.e. after the analysis of intravascular bacterial adherence, 0.1mL 5% FITC-labeled dextran 150,000 were additionally injected for contrast enhancement by staining of blood plasma in order to determine microhemodynamic parameters (Fig. 1C) and 0.1mL 0.1% rhodamine 6G (Sigma) for direct staining of white blood cells.

The microscope used for our study was a Zeiss Axiotech microscope (Zeiss; Oberkochen, Germany) with a 100W mercury lamp attached to an epi-illumination filter block for blue (450–490nm excitation; >520nm emission wavelength), green (530– 560nm; >580nm) and ultraviolet (330– 390nm; >430nm) light. The microscopic images were recorded by a charge-coupled device video camera (FK6990; Pieper, Schwerte, Germany) and transferred to a DVD system for off-line evaluation (Panasonic AG-7350-SVHS; Matsushita, Tokyo,

Japan). By means of 5x (numerical aperture $(n.a.) = 0.15$), $20x (n.a. = 0.4)$ and $50x (n.a. =$ 0.55) objectives, magnifications of \times 115, \times 430 and \times 1150 were achieved on a 14- inch video screen (KV-14CT1E; Sony, Tokyo, Japan).

Microcirculatory analysis

Quantitative off-line analysis of the DVDs was performed using the computer-assisted image analysis system CapImage (Zeintl; Heidelberg, Germany). In each animal, analyses were performed in 8–10 postcapillary or collecting venules. To exclude that differences in bacteria-endothelial cell interaction between MAb 74.5.2-treated animals and respective controls were caused by changes in local microhemodynamics, we additionally analyzed the diameters, centerline velocity, volumetric blood flow, and wall-shear rate of these vessels as described previously (Laschke et al., 2005a).

Adherent *staphylococci* were defined as bacteria which did not move or detach from the endothelial lining of venules during an observation period of 20s and are given as the number of adherent cocci/mm² of endothelial surface. Leukocytes were classified according to their interaction with the microvascular endothelium as rolling or adherent cells (Laschke et al., 2009). Rolling leukocytes were defined as cells moving with a velocity less than twofifths of the centerline velocity, and are given as number of cells/min, passing a reference point within the microvessel. Adherent leukocytes were defined as cells that did not move or detach from the endothelial lining within a 20s observation period, and are given as number of cells/mm² of endothelial surface.

In addition, macromolecular leakage as a parameter of microvascular permeability and, thus, endothelial integrity was assessed after intravenous injection of the macromolecular fluorescent dye FITC-labeled dextran 150,000 by determining densitometrically grey levels in the tissue directly adjacent to the venular vessel wall (E_1) , as well as in the marginal cell free plasma layer within the vessel (E₂). Extravasation (E) was then calculated as $E = E_1/E_2$ (Laschke et al., 2005b).

Experimental protocol

A dorsal skinfold chamber was prepared in 25 Syrian golden hamsters. After 48h, animals received either an i.p. injection of $100mg/kg$ MAb 74.5.2 (1.6 mL, n = 13), which was kindly provided by the Cell Culture/Hybridoma Facility, Department of Molecular Genetics and Microbiology, Stony Brook University (Stony Brook, NY, USA), or serum free medium control (1.6 mL, n = 12) according to prior studies (Peerschke et al., 2006). Then, 48h later, the tissue of the dorsal skinfold chamber was exposed to tumor necrosis factor (TNF)-α (topical application for 30 min, 2000 U dissolved in 100μl PBS; Boehringer Mannheim, Germany) in order to induce inflammation. In a second set of experiments, the chamber tissue was solely exposed to 100μl PBS (topical application for 30 min). Subsequently, intravital fluorescence microscopy was performed at 5, 10, 20, 30 and 60 min after application of 10^9 FITC-labeled bacteria. At the end of the experiment, each animal received 5% FITC-labeled dextran 150,000 and 0.1% rhodamine 6G i.a. for the quantitative assessment of microhemodynamic parameters, macromolecular leakage, and leukocyteendothelial cell interaction.

Enzyme-linked immunosorbent assay (ELISA)

To quantify levels of anti-gC1qR antibodies in the hamster serum, an ELISA was used as described previously (Peerschke et al., 2006). In brief, soluble, recombinant gC1qR was immobilized on 96-well plates. After blocking the wells, standard concentrations of antigC1qR antibodies (0–10 μg/mL) were prepared in naïve hamster serum diluted $1/10$ with 0.3M NaCl and then added to the wells. Test hamster serum was diluted 1/10 in 0.3M NaCl

and added to separate wells. Bound antibody was detected spectrophotometrically (OD405nm), using alkaline phosphatase-conjugated goat anti-mouse immunoglobulin G (IgG; Sigma) and p-NPP substrate. The reactivity of test plasmas was compared to that of standard anti-gC1qR antibody solutions. Each serum sample was evaluated in duplicate for data reproducibility. Results were expressed in μg/mL serum.

In vitro adherence assay

Mock or TNF-α-pretreated EA.hy endothelial cells (Edgell et al., 1983) were detached by trypsin treatment and subsequently exposed to FITC-labeled bacteria. Samples were harvested at the indicated times and attachment of bacteria was analyzed by using flowcytometry (FACS Calibur and CellQuest Pro Software; Becton Dickinson, Heidelberg, Germany). Experiments were performed with MAb 74.5.2 (500μg/mL in serum-free medium $+10\%$ FCS), respective controls (serum free medium $+10\%$ FCS) and with as well as without TNF-α stimulation. The results were confirmed in five independent experiments.

Statistics

Differences between the groups were tested separately at each time point by the unpaired Student's t-test and Mann-Whitney U test. To test for time effects within individual groups, multivariate ANOVA for repeated measures was applied. This was followed by the paired Student's t-test, including correction of the α -error according to Bonferroni probabilities for repeated measurements (SigmaStat; Jandel Corporation, San Rafael, CA, USA). All values are expressed as means \pm SEM. Statistical significance was accepted for a value of p <0.05.

Results

Serum concentration of MAb 74.5.2

In contrast to control animals (gC1qR MAb titer: 0 ± 0 μg/mL), significantly high sustainable concentrations of MAb 74.5.2 were measured 48h following i.p. administration of 100mg/kg MAb 74.5.2 (gC1qR MAb titer: 109±23 μg/mL).

Effects of gC1qR blockade during TNF-α-induced inflammation

TNF-α-induced inflammation resulted in the adherence of many late exponential phase *staphylococci* to the endothelium of postcapillary and collecting venules in dorsal skinfold chambers of control animals (Figs. 2A and C). Highest numbers of adherent bacteria (~850mm−²) could be observed 5 min after *S. aureus* administration. This was followed by a progressive decline in bacterial adherence to ~510mm⁻² at time point 60 min (Fig. 2C). In contrast, blockade of gC1qR by application of MAb 74.5.2 effectively reduced *S. aureus* adherence (Figs. 2B and C). Throughout the entire experiment, antibody-treated animals presented with significantly decreased numbers of adherent bacteria of \sim 260–480mm⁻² when compared to controls (Fig. 2C; 5min: $p = 0.038$; 10min: $p = 0.011$; 20min: $p = 0.011$; 30min: $p = 0.002$; 60min: $p = 0.021$).

The analysis of microhemodynamic parameters, i.e. microvessel diameter, centerline velocity, volumetric blood flow and wall-shear rate, did not show any significant differences between antibody-treated animals and respective controls (Tab. 1). Furthermore, leukocyteendothelial cell interaction remained unaffected under administration of the gC1qR antibody. In both groups, animals exhibited numbers of rolling and adherent leukocytes of ~6–7min⁻¹ and ~800mm⁻² (Figs. 3A and B). However, treatment with MAb 74.5.2 resulted in a significantly decreased macromolecular leakage (0.40 ± 0.03) when compared to controls $(0.54 \pm 0.02; p = 0.002)$ (Fig. 3C).

Effects of gC1qR blockade under baseline conditions

To determine whether the observed reduction of *S. aureus* adherence by gC1qR blockade was based on the interference with TNF-α-induced pathways, we further examined the effects of gC1qR blockade in dorsal skinfold chambers, which were not exposed to $TNF-\alpha$. As expected, bacterial adherence in postcapillary and collecting venules was distinctly reduced under these baseline conditions when compared to TNF-α-treated tissue (Figs. 4A and C). Numbers of adherent bacteria ranged between ~200–335mm−² in vehicle-treated animals throughout the observation period of 60 min. Nonetheless, treatment with the antigC1qR antibody again significantly decreased *S. aureus* adherence. In fact, antibody-treated animals presented with numbers of adherent bacteria of only \sim 70–135mm⁻², according to a ~60% reduction when compared to controls (Figs. 4B and C; 5min: $p = 0.052$; 10min: $p =$ 0.052; 20min: $p = 0.018$; 30min: $p = 0.009$; 60min: $p = 0.030$). These findings clearly indicate that bacterial adherence to the microvascular endothelium is directly affected by gC1qR blockade, independently from TNF-α-induced pathways. According to our first set of experiments, analysis of microhemodynamic parameters and leukocyte-endothelial cell interaction did not show any significant differences between antibody-treated and control animals (Tab. 1, Figs. 3A and B). Moreover, macromolecular leakage remained unaffected under treatment with MAb 74.5.2 (Fig. 3C).

Effects of gC1qR blockade on S. aureus adherence to human endothelial cells in vitro

In line with our *in vivo* experiments, we could also demonstrate *in vitro* the tendency that gC1qR blockade reduced the adherence of *S. aureus* to human endothelial cells, which had been exposed to TNF- α (Fig. 5A; 5min: p = 0.761; 10min: p = 0.217; 20min: p = 0.171; 30min: $p = 0.220$; 60min: $p = 0.686$). Moreover, bacterial adherence was also decreased, when cells were not pretreated with TNF- α (Fig. 5B; 5min: p = 0.529; 10min: p = 0.204; 20min: p = 0.180; 30min: p = 0.363; 60min: p = 0.486). However, the results of the *in vitro* experiments were not proven to be statistically significant.

Discussion

The present study demonstrates that blockade of gC1qR by administration of MAb 74.5.2 significantly reduces the adherence of *S. aureus* to the venular endothelium under *in vivo* conditions. For this purpose, we used the dorsal skinfold chamber model in combination with the technique of intravital fluorescence microscopy. This approach is unique as it allows for the real-time observation of staphylococcal interaction with the microvascular endothelium.

In a first set of experiments we studied the effects of MAb 74.5.2 administration on the adherence of *S.aureus* to the venular endothelium activated via TNF-α. TNF-α exerts a variety of different inflammatory effects on the endothelium (Bradley, 2008; Pober et al., 1998). Accordingly, we have previously shown that TNF-α markedly increases the number of vessel wall-adhering *staphylococci* (Laschke et al., 2005a). This was also the case in the present study, however, only under *in vivo* but not under *in vitro* conditions, indicating that the observed TNF-α-stimulated bacterial adherence may involve mechanisms additional to the expression of endothelial binding molecules on *in situ* or cultured endothelial cells. In fact, the endovascular system is characterized by various humoral and cellular factors contained in whole blood, many of which are also prone to TNF-α-mediated activation. These include binding of fibrinogen to endothelial cells or activation of circulating cells such as leukocytes (Vaday et al., 2001), which in turn may affect the binding of bacteria to the vessel wall.

In the present study, we now demonstrate that the TNF-α-induced adherence of *S. aureus* to the venular endothelium is significantly inhibited by blockade of gC1qR. On the one hand, this effect may be explained by the blockade of $TNF-\alpha$ -induced endothelial adherence of activated gC1qR-expressing platelets or leukocytes, which can serve as binding sites for free flowing bacteria in the microcirculation (Peerschke and Ghebrehiwet, 2007). However, in unactivated tissue we observed comparable results. Therefore, we suggest that administration of MAb 74.5.2 also directly affects the gC1qR-dependent binding of *S. aureus* to the microvascular endothelium. In line with this assumption, we could demonstrate in our *in vitro* assay excluding interfering platelets and leukocytes that *S. aureus* binding to human endothelial cells was inhibited by gC1qR-blockade albeit to a reduced extent when compared to our *in vivo* findings. Thereby, it was not surprising that in our *in vitro* and *in vivo* experiments not all adhesion of *staphylococcus* was inhibited by the anti-gC1qR antibody. In fact, there are various surface adhesins and ligands promoting the adhesion of *S. aureus* to the vascular endothelium (Moreillon et al., 2002). Particularly the interaction of vWF with its *staphylococcal* receptors protein A (Hartleib et al., 2000) or with the vWF binding protein (Bjerketorp et al., 2002) may mediate this interaction. Accordingly, it could not be expected that the MAb directed against gC1qR would completely abrogate this interaction, yet our study proves it to be a relevant and novel interaction partner in this system.

gC1qR is recognized by protein A on the surface of *S. aureus* (Nguyen et al., 2000). Although gC1qR interacts with protein A at or near the IgG binding site (Nguyen et al., 2000), we believe that the observed effects of our MAb 74.5.2 administration are based on specific interactions of the MAb rather than on unspecific IgG binding to protein A, because physiological concentrations of circulating IgG are high and application of low amounts of IgG as performed in this study is unlikely to have a significant effect.

As it was previously demonstrated that soluble gC1qR binds to *S. aureus*, it is conceivable that *S. aureus* decorated with gC1qR might interact with additional ligand molecules recognizing gC1qR. Candidate ligands include the fibrinogen D domain, which is a known binding partner for gC1qR. In fact, since fibrinogen/fibrin is an important component of endovascular lesions (Archer, 1998; Moreillon and Que, 2004; Peerschke et al., 2006), gC1qR-decoration may enhance *S. aureus'* ability to interact with sites of local fibrinogen/ fibrin exposure, e.g. sites of subendothelial exposure. Moreover, gC1qR has been shown to interact with various components of the human contact system, also known as the kallikreinkinin cascade or intrinsic pathway of coagulation such as factor XII or high molecularweight kininogen (HMW-kininogen) (Ghebrehiwet and Peerschke, 1998). The surface of *S. aureus* activates this contact system, resulting in the release of short-lived bradykinin (Mattsson et al., 2001) and a subsequent pro-inflammatory response mediated by conversion of bradykinin to the bradykinin 1 receptor (B1R) agonist desArg(9)bradykinin, which leads to a further up-regulation of B1R, and thereby ensues a chronic inflammatory response (Bengtson et al., 2006). Considering our findings, demonstrating a reduced interaction of *S. aureus* with the vessel wall upon blockade of gC1qR, it may be speculated that this interaction may also interfere with the *S. aureus*-triggered local contact system activation, resulting in a decreased inflammatory leukocyte and endothelial cell activation.

For these reasons, we also assessed the leukocyte-endothelial cell interaction in venules of the dorsal skinfold chamber. We found that exposure of the tissue to $TNF-\alpha$ resulted in a strong inflammation with significantly increased numbers of adherent leukocytes when compared to PBS-treated tissue. Meanwhile, it is well known that the interaction of leukocytes with the endothelium is regulated by specific ligand-receptor interactions. On activation, leukocytes roll along the venular endothelium mediated by the selectin family of adhesion molecules (Klintman et al., 2004; Laschke et al., 2008; Springer, 1994). This

leukocyte rolling is a major prerequisite for the subsequent firm adhesion of leukocytes, which is mediated by integrins expressed on leukocytes, in particular lymphocyte function antigen-1 (LFA-1) and macrophage antigen-1 (MAC-1), which interact with molecules of the immunoglobulin supergene family expressed on endothelial cells, such as intercellular adhesion molecule-1 (ICAM-1) and vascular cell adhesion molecule-1 (VCAM-1) (Carlos and Harlan, 1994; Springer, 1994). Interestingly, we found that blockade of gC1qR did not affect this well coordinated process, indicating that gC1qR blockade may selectively inhibit the adherence of *S. aureus* to the endothelium without interfering with the interaction of leukocytes and endothelial cells.

Furthermore, we could observe in our study a significantly decreased macromolecular leakage in TNF-α-exposed tissue of animals, which were treated with anti-gC1qR antibodies. Because we did not detect any effects of gC1qR blockade on leukocyteendothelial cell interaction, we propose that this observation may be due to a direct inhibition of the gC1qR-mediated activation of the kallikrein-kinin cascade and, thus, the formation of bradykinin (Joseph et al., 1999). In fact, bradykinin has been shown to be a potent direct inducer of vascular permeability during infection (Matsumoto et al., 1986; Maeda et al., 1999). Accordingly, blockade of gC1qR may also prevent transvascular spread of bacteria.

Taken together, we could demonstrate in the present study that gC1qR blockade effectively inhibits the adherence of *S. aureus* to the microvascular endothelium. In light of the fact that gC1qR plays a pivotal role in the pathogenesis of *S. aureus*-associated endovascular disease and that the resistance of *S. aureus* to antimicrobial therapy is increasing (Chastre, 2008), immunological blockade of gC1qR represents a promising novel therapeutic strategy in the treatment of *S. aureus*-associated endovascular diseases.

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Research highlights

- **•** Intravital fluorescence microscopy for the analysis of bacteria-endothelial cell interaction
- **•** Blockade of gC1qR/p33 by administration of monoclonal antibodies
- **•** gC1qR/p33 blockade inhibits adherence of *S. aureus* to the microvascular endothelium
- **•** gC1qR/p33 blockade does not affect leukocyte-endothelial cell interaction

Figure 1.

A: Syrian golden hamster equipped with a dorsal skinfold chamber (chamber weight \sim 4g). Within the observation window (arrow) all segments of the microcirculation including arterioles, capillaries and venules of the striated skin muscle and subcutaneous tissue can be analyzed using trans- and epi-illumination microscopy. B: Overview of the observation window. C: Intravital fluorescence microscopic image of the micro-angioarchitecture of the dorsal skinfold chamber displaying a representative venule (arrowheads) and arteriole (arrows). Note that the arteriole exhibits a straight-lined wall structure and much less bifurcations and branches than the venule. Blue light epi-illumination with intravascular plasma contrast enhancement by 5% FITC-labeled dextran 150,000 i.v.. Scale bars: A = 3cm, $B = 1.7$ mm, $C = 400$ μ m.

Figure 2.

A, B: Intravital fluorescence microscopy of FITC-labeled *Staphylococcus aureus* in postcapillary and collecting venules of the dorsal skinfold chamber (5 min after intra-arterial injection of bacteria) after local TNF-α-exposure. Note that blockade of gC1qR by application of MAb 74.5.2 (B) results in a significant reduction of bacterial adherence when compared to controls (A). Blue light epi-illumination; scale bars: 100μm. C: Adherent bacteria (mm−²) in venules of TNF-α-exposed dorsal skin-fold chambers of Syrian golden hamsters, which were treated with MAb 74.5.2 (black bars; $n = 7$) or vehicle (control, white bars; $n = 7$), as assessed by intravital fluorescence microscopy and computer-assisted image analysis. Measurements were performed at 5, 10, 20, 30, and 60 min after intra-arterial injection of the bacteria. Values are given as means \pm SEM; *p<0.05 vs. control.

Figure 3.

Rolling leukocytes (min−¹) (A), adherent leukocytes (mm−²) (B) and macro-molecular leakage (E_1/E_2) (C) in venules of TNF- α - or PBS-exposed dorsal skinfold chambers of Syrian golden hamsters, which were treated with MAb 74.5.2 (MAb, black bars; TNF- α : n = 7, PBS: $n = 6$) or vehicle (Con, white bars; TNF- α : $n = 7$, PBS: $n = 5$), as assessed by intravital fluorescence microscopy and computer-assisted image analysis. Values are given as means \pm SEM; *p<0.05 vs. control.

Figure 4.

A, B: Intravital fluorescence microscopy of FITC-labeled *Staphylococcus aureus* in postcapillary and collecting venules of the dorsal skinfold chamber (5 min after intra-arterial injection of bacteria) under physiological conditions. Note that blockade of gC1qR by application of MAb 74.5.2 (B) results in a significant reduction of bacterial adherence when compared to controls (A). Blue light epi-illumination; scale bars: 100μm. C: Adherent bacteria (mm−²) in venules of PBS-exposed dorsal skinfold chambers of Syrian golden hamsters, which were treated with MAb 74.5.2 (black bars; $n = 6$) or vehicle (control, white bars; $n = 5$), as assessed by intravital fluorescence microscopy and computer-assisted image analysis. Measurements were performed at 5, 10, 20, 30, and 60min after intra-arterial injection of the bacteria. Values are given as means \pm SEM; *p<0.05 vs. control.

Figure 5.

In vitro adherence of *Staphylococcus aureus* to endothelial cells. FITC-labeled bacteria were incubated with endothelial cells and adherence of *Staphylococcus aureus* to endothelial cells was measured using FACS analysis in the presence of MAb 74.5.2 (black bars; $n = 5$) and in respective controls (serum free medium $+ 10\%$ FCS; white bars; n = 5). The experiments were performed with (A) and without TNF- α treatment (B). Values are given as means \pm SEM.

Table 1

Diameter, centerline velocity, volumetric blood flow and wall shear rate of postcapillary and collecting venules within the dorsal skinfold chamber of antibody-treated (MAb) and vehicle-treated (Con) Syrian golden hamsters. Chambers were pretreated with TNF- α (MAb: n = 7; Con: n = 7) or PBS (MAb: n = 6; Con: n = 5). Parameters were assessed by intravital fluorescence microscopy and computer-assisted image analysis. Values are given as means \pm SEM; data did not significantly differ between the groups.

