To analyze the glycosylation of anti-β2GP1, we investigated purified IgG from healthy children, patients with APS, and asymptomatic adult carriers of antiphospholipid antibodies. We observed that in the sera of healthy children and of patients with APS, IgG3 and IgG2 were predominant, respectively. The potentially protective anti-β2GP1-IgM was lower in the sera of healthy children. Although anti-β2GP1-associated C1q did not differ between children and patients with antiphospholipid syndrome, the associated C3c was significantly higher in the sera of healthy children. This indicates a more efficient clearance of anti-β2GP1 immune complexes in the healthy children. This clearance is not accompanied by inflammation or coagulatory events. It is likely that the most important pathogenic factor of the anti-β2GP1-IgG is related to the different glycosylation observed in healthy and diseased individuals. We detected a significantly higher sialylation of anti-β2GP1-IgG isolated from the sera of healthy children and asymptomatic adults when compared with that of patients with clinically apparent antiphospholipid syndrome. Low sialylated IgG reportedly ameliorates inflammation and inflammation promotes hyposialylation. Thus, both reactions create a vicious circle that precipitates the pathology of the antiphospholipid syndrome including thrombus-formation. We conclude that the increased sialylation of anti-β2GP1-IgG of sera of healthy individuals limits their pathogenicity.

1. Introduction

The antiphospholipid syndrome (APS) is an autoimmune disease affecting the coagulation system. Its diagnosis requires at least one clinical criterion and one serological criterion as defined during conferences held in 1998 and in 2006 [1, 2]. The serological criteria include the detection of so-called antiphospholipid antibodies (aPL) which comprise the lupus anticoagulant (LA), anticardiolipin autoantibodies (aCL), and anti-β2-glycoprotein I autoantibodies (anti-β2GP1) [1].

Thrombosis as one major symptom in APS is mainly associated with anti-β2GP1 [3]. β2GP1 is a plasma protein
composed of five domains [4] and it is known to be an inhibitor of the contact activation of the intrinsic coagulation pathway [5]. Additionally, aPL can cause a prothrombotic state in individuals by activating platelets and endothelial cells resulting in a higher expression of tissue factor or in the upregulation of proinflammatory cytokines and cell adhesion molecules [6, 7].

Patients with APS can be classified into two groups depending on the occurrence of further diseases: (I) patients merely suffering from APS (primary antiphospholipid syndrome; PAPS) and (II) those with an underlying systemic lupus erythematosus (SLE). The latter condition is referred to as secondary antiphospholipid syndrome (SAPS) [8].

Up to now, a variety of hypotheses about possible triggers for the induction of anti-β2GP1 have been proposed. Published studies suggest that their production can be provoked during infectious diseases like parvovirus B19, HIV, CMV, or HCV [3, 9–11]. Certain molecular structures of these infectious agents may resemble structures of β2GP1 and thus cause the induction of cross-reacting antibodies [7, 12–16]. Another one claims that nutritional exposure to β2GP1 could also trigger the production of transient and harmless anti-β2GP1 in children [4, 17–19].

Interestingly, anti-β2GP1 can be detected in a quite large proportion of the healthy population who do neither suffer from APS nor another autoimmune disease [3, 18, 20]. As complication-free periods of patients with APS occur although the aPL are persistently detectable in their circulation [15], it seems to be likely that the mere presence of these antibodies is not sufficient for the development of clinical active APS complications.

For this reason, a two-hit hypothesis has been proposed to provide a putative explanation for this observation. (I) the production of anti-β2GP1 represents an initial hit that increases the risk for thrombotic events and (II) infectious agents can then serve as the second hit provoking the typical manifestations of APS by activating toll-like receptors or complement [9, 15, 21, 22]. This two-hit hypothesis surely provides a good first model for the etiopathogenesis of APS. Yet it is not completely satisfying and does not explain why some people harbouring aPL remain healthy. Some study results indicate that this could be grounded in structural differences of anti-β2GP1 found in patients with APS and healthy children [4, 23].

Comparing obviously harmless aPL from healthy children with putatively pathogenic aPL from adults with APS, we suggested a difference in the epitope specificity as differentiating factor. Anti-β2GP1 from the sera of the children specifically bound to the domain IV/V of β2GP1, whereas anti-β2GP1 from adults with APS preferentially targeted domain I [24]. These results were consistent with those described in other studies [19, 25]. Besides distinct epitope binding specificities, a different composition of the spectrum of the anti-β2GP1 IgG subclasses between patients with APS and healthy carriers of this antibody has also been described [19, 23, 26, 27].

The differences in glycosylation of antibodies which are accompanied by crucial modifications of their effector functions [28] provide another promising approach in performing structural analyses of anti-β2GP1. IgG molecules possess a biantennary oligosaccharide attached to the asparagine at position 297 of the Fc portion (Asn-297). Various glycoforms containing distinct sugar moieties have been described [28, 29]. Interestingly, it has been observed that IgG molecules whose glycans terminate with sialic acid residues show a decreased affinity to activating Fcy-receptors (FcyRs) while simultaneously provoking an upregulation of the inhibitory FcyRIIB [29–31]. These effects lead to a rather anti-inflammatory activity and might consequently be able to prevent potentially harmful immune responses.

In particular in children, aPL can readily be detected although they do not show any clinical signs of APS. In a study carried out by Avˇcin et al., the prevalence of anti-β2GP1 in 61 healthy children was 6.6% [18]. Based on this observation we intended to find out why these circulating autoantibodies do not cause harm in their hosts. We determined the IgG-subclass distribution in the samples from healthy children and patients with APS and examined the binding of the complement factors C1q and C3c to anti-β2GP1 immune complexes, since an important involvement of the complement system in the pathogenesis of APS has been reported [21, 22, 32]. Finally we analysed the sialylation status of anti-β2GP1-IgG isolated from healthy children, asymptomatic adult carriers, and patients with APS.

Our results revealed that, compared to adults with APS, the healthy children showed a higher content of IgG3 at the cost of IgG2 of the anti-β2GP1. Furthermore, anti-β2GP1-IgG of healthy children showed a higher sialylation and displayed more bound C3c. Hyposialylated IgG and inflammation mutually influence the pathogenicity of the antiphospholipid autoantibodies and form a vicious pathogenic circle.

2. Materials and Methods

2.1. Patients. For the investigation of anti-β2GP1-IgG, its subclasses, C1q- and C3c-binding and anti-β2GP1-IgM, we studied sera of 46 adult patients (24 patients with PAPS and 22 with SAPS) and 16 healthy children harbouring anti-β2GP1 without any clinical manifestations associated with APS from our partner hospital in Brescia, Italy. As a control group, we examined sera of 17 age and sex matched adult normal healthy donors (NHD) without aPL. Inclusion criteria for the children were the presence of anti-β2GP1-IgG, no infectious diseases in the course of three months before blood taking, and no chronic pathology. Additionally, it was excluded that their parents suffered from any autoimmune disease. At the time of blood taking, the mean age of the children was about 9 months (median age: 8.5 months; range: 3 months and 5 days–19 months and 10 days).

Since sugar residues attached to molecules can be damaged by repeated thawing and freezing, we received new sera from our partner hospital in Brescia, Italy, to analyse the sialylation of anti-β2GP1-IgG. We obtained sera from 12 patients with PAPS and 10 patients with SAPS, 15 patients with SLE and aPL but without clinical evidence of APS and sera from 6 asymptomatic carriers of aPL. Furthermore, we obtained sera from 16 healthy children showing no symptoms of APS although harbouring anti-β2GP1-IgG in
their circulation (mean age: 9.5 months; median age: 10 months; range: 3 months and 5 days–19 months and 10 days).

2.2. Blood Drawing. Heparinised venous blood was drawn by venipuncture and processed within two hours. Serum was obtained by centrifugation at 1000 g for 15 minutes of clotted blood, then aliquoted, frozen, and stored at −20°C until use for investigation.

Informed consent was obtained from all blood donors and the children’s parents, respectively, and the study was approved by the ethics committees of the University Hospital Erlangen and the Spedali Civili Brescia.

2.3. Isolation of β2GPI. For β2GPI-isolation from diluted sera of normal healthy donors (in 0.15 M NaCl), perchloric acid was slowly added until the mixture got a creamy aspect. After centrifugation at 10,000 g, the supernatant was neutralized with NaOH and dialyzed against a buffer of 0.02 M Tris and 0.03 M NaCl, pH 8. Then it was given into columns containing HiTrap heparin and contaminants of the supernatant were eluted by the addition of 0.02 M Tris and 0.35 M NaCl, pH 8 was injected into the columns and the fractions were been pooled and dialyzed against a solution containing 0.05 M acetate buffer, 0.05 M NaCl, pH 4.8. This material was filtrated with 0.2 μ membranes and injected into resin columns equilibrated with 0.05 M acetate buffer, 0.05 M NaCl, pH 4.8. After washing with 0.05 M acetate buffer, 0.65 M NaCl, pH 5.2, the fractions were pooled and dialyzed against PBS, pH 7.2. To verify the purity of isolated β2GPI, a sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) has been performed.

2.4. Purification of Anti-β2GPI-IgG. For the analysis of the sialylation of anti-β2GPI-IgG, we isolated IgG antibodies from the sera to exclude possible interactions of the detection lectin with anti-β2GPI-IgM in the samples. Purification of anti-β2GPI-IgG was performed as follows: 20 μL of serum was added to 50 μL magnetic Dynabeads Protein G (Nvex by life technologies, Carlsbad, CA, USA) and incubated for 45 minutes at room temperature. After washing with 1x TBS, glycine-HCl, pH 2.6, was added for 5 minutes to elute the anti-β2GPI-IgG. The supernatant, now containing the IgG antibodies, was neutralized with Tris-HCl, pH 8.8, and all samples were centrifuged using 100 K Amicon Ultra-centrifugal filter devices (Millipore, Tullagreen, Ireland) to transfer them into 300 μL TBS-buffer.

2.5. Serum ELISAs: Anti-β2GPI-IgG, -IgG1, -IgG2, -IgG3, -IgG4, and -IgM and C3c and Clq Bound to Anti-β2GPI. To quantify the amount of anti-β2GPI-IgG and its four subclasses, of C3c and Clq associated with anti-β2GPI and of anti-β2GPI-IgM in the sera of patients with PAPS and SAPS, NHD, and healthy children, we performed enzyme-linked immunosorbent assays (ELISA). In detail, the surface of F96 NUNC maxisorp microtiter plates (Thermo Scientific, Roskilde, Denmark) was coated with 50 μL of a 2 μg/mL solution of β2GPI in coating buffer (100 mM Na₂CO₃, 100 mM NaHCO₃, pH 9.6) at 4°C overnight. After washing the plates three times with washing buffer (WB; 0.05% Tween 20, 0.1 M CaCl₂, 0.1 M MgCl₂ in TBS buffer (150 mM NaCl, 50 mM TrisHCl, pH = 7.4)), the wells of the coated plates were blocked with 100 μL/well of a 2% BSA solution for two hours at 37°C. After three further washing steps with WB, 50 μL/well of diluted serum (1:200 in WB) was added and the plates were incubated at 37°C for two further hours. Then, the plates were washed three times with WB again.

Development of the plates based on an enzymatic reaction of horseradish peroxidase (HRP) with TMB as chromogenic substrate. Therefore, the appropriate HRP-conjugated detection-antibody was added as a next step (Table 1). After incubation at 37°C for another hour and three washing steps with WB, the plates were developed by the addition of 50 μL/well of freshly prepared TMB-solution (1 mL of 0.01 g 3,3′,5,5′-tetramethylbenzidine (Sigma-Aldrich, Steinheim, Germany) dissolved in 10 mL dimethyl sulfoxide, mixed with 9 mL of 0.1 M Na₂HPO₄, 0.05 M citric acid, pH 4.5–5.5, and 2 μL of H₂O₂). The reaction was stopped by adding 50 μL/well of 25% sulfuric acid (H₂SO₄). Finally, optical density (OD) was measured at 450 nm and 650 nm as correction factor using an infinite F200 pro ELISA-reader (Tecan).

After subtraction of the corresponding blank values of the plates, experimental values were expressed as optical density or as mean optical density of each serum when ELISA tests were performed in monoplicates or duplicates, respectively.

2.6. IgG-ELISA: SNA. In order to examine the sialylation of anti-β2GPI-IgG, we slightly modified the previously described ELISA protocol. Before coating, the β2GPI antigen was treated with 1% of periodic acid (PIA, Sigma-Aldrich,

---

**Table 1: Antibodies employed for each ELISA.**

<table>
<thead>
<tr>
<th>ELISA</th>
<th>Detection-antibody</th>
<th>Dilution</th>
</tr>
</thead>
<tbody>
<tr>
<td>Anti-β2GPI-IgG</td>
<td>HRP-conjugated rabbit F(ab’) 2 anti-human-IgG*</td>
<td>1:20,000</td>
</tr>
<tr>
<td>Anti-β2GPI-IgG subclasses</td>
<td>HRP-conjugated mouse anti-human IgG1, 2, 3 and 4 (Clones HP6001, HP6002, HP6050 &amp; HP6025)*</td>
<td>1:20,000</td>
</tr>
<tr>
<td>Anti-β2GPI-IgM</td>
<td>HRP-conjugated goat anti-human-IgM*</td>
<td>1:20,000</td>
</tr>
<tr>
<td>Anti-β2GPI-bound C3c</td>
<td>HRP-conjugated sheep anti-C3c**</td>
<td>1:500</td>
</tr>
<tr>
<td>Anti-β2GPI-bound Clq</td>
<td>HRP-conjugated sheep anti-C1q**</td>
<td>1:500</td>
</tr>
</tbody>
</table>

*:* purchased from Southern Biotech, Birmingham, Alabama, USA.

**HP6002, HP6050 & HP6025**

HRP = horseradish peroxidase.
Steinheim, Germany) for 48 hours at 43°C to remove its sugar residues to exclude unspecific binding of Sambucus nigra lectin (SNA) to naive β2GP1. After that, we centrifuged the β2GP1-solution with 10 K Amicon Ultra-centrifugal filter devices (Millipore, Tullagreen, Ireland) to transfer the treated β2GP1 into coating buffer (100 mM Na2CO3, 100 mM NaHCO3, pH 9.6).

Then, the plates were coated with 50 μL/well of a 2 μg/mL solution of the PIA-treated β2GP1 at 4°C overnight and subsequently blocked for two hours at 37°C with 100 μL/well of PIA-treated (0.5%) 3% gelatine (Sigma-Aldrich, Steinheim, Germany) in TBST buffer with 1 mM CaCl2 and 1 mM MgCl2. The subsequent steps were performed as described in the previous ELISA protocol with two exceptions. Instead of serum, we added 50 μL/well of purified IgG-samples to exclude possible binding of SNA to anti-β2GP1-IgM. And the development of the plate required the addition of a second detection reagent (Table 2), which was incubated for one hour at room temperature and demanded three preliminary washing steps. We calculated the SNA/anti-β2GP1-IgG ratio from the OD values we obtained in the SNA ELISA and a simultaneously conducted anti-β2GP1-IgG ELISA.

2.7. Statistical Analysis. Differences between the cohorts were assessed using the one-way analysis of variance (ANOVA) test. In the case of multiple testing, the significance values were adjusted by the Newman-Keuls posttest correction.

3. Results

3.1. Analysis of Anti-β2GP1-IgG. We measured the OD values of total anti-β2GP1-IgG in the sera of 24 patients with PAPS and 22 patients with SLE/APS, 16 healthy children, and 17 NHD as controls. As shown in Figure 1, the lowest IgG values were to be observed in the cohort of healthy donors (308 mOD), whereas the values of the healthy children (970 mOD) even exceeded that of the patients with APS (PAPS: 686 mOD, p = 0.010; SLE/APS: 546 mOD, p < 0.001). All differences between NHD and the other cohorts as well as between the children and the other cohorts were significant. Patients with PAPS and SLE/APS did not differ significantly.

We calculated the 97.5% quantile of the anti-β2GP1-IgG OD values obtained from the NHD cohort and set this value as cut-off (738 mOD). Sera exceeding this value were considered seropositive for anti-β2GP1-IgG. Applying this cut-off value, we obtained 31 seropositive samples (12/24 PAPS, 6/22 SAPS, 13/16 healthy children), which were further analysed in our study (Supplementary Figure 1, in Supplementary Material available online at http://dx.doi.org/10.1155/2015/638129).

3.2. Analysis of the IgG-Subclasses of Anti-β2GP1. Analysing the absolute OD values of the anti-β2GP1-IgG subclasses, we observed that the children showed significantly higher levels of the anti-β2GP1-IgG1 (69 mOD) and -IgG3 subclasses (212 mOD) compared to NHD, PAPS, and SAPS (Figures 2(a) and 2(c); Table 3(a)). The differences between healthy children and NHD regarding these two subclasses showed a higher level of significance than the comparison between healthy children and patients with APS. However, healthy children did not show significant differences for anti-β2GP1-IgG2 and -IgG4 when compared to NHD, PAPS, and SAPS (Figures 2(b) and 2(d); Table 3(b)).

To summarize, patients with APS showed higher OD values than NHD in IgG1, IgG2, and IgG3 than NHD (Figures 2(a), 2(b), and 2(c); Table 3(a)). As in the anti-β2GP1-IgG ELISA, we observed no significant difference between the patients with PAPS and SAPS.

These results were consistent with the analysis of the number of seropositive samples, as shown in Table 4. To determine seropositivity, we calculated the mean OD and the standard deviation (SD) of the NHD group for the different IgG subclasses. IgG values exceeding the NHD’s mean OD...
Figure 2: mOD values of anti-β2GP1-IgG1 (a), anti-β2GP1-IgG2 (b), anti-β2GP1-IgG3 (c), and anti-β2GP1-IgG4 (d) in NHD, PAPS, SLE/APS, and children. The horizontal lines represent the cut-off for seropositivity of the corresponding anti-β2GP IgG subclass (> mean mOD_NHD + 2 × SD_NHD); NHD = normal healthy donor; PAPS = primary antiphospholipid syndrome; SLE/APS = antiphospholipid syndrome with systemic lupus erythematosus; OD = optical density. Median OD values of the anti-β2GP1-IgG subclasses in the study cohorts (e). Regarding the anti-β2GP1-IgG subclass-distribution within the cohorts, children showed a significantly higher contribution of IgG3 than IgG1 and IgG2. Patients with antiphospholipid syndrome (APS) showed a higher content of anti-β2GP1-IgG2 than -IgG1 and -IgG3. NHD showed a higher contribution of anti-β2GP1-IgG2 than -IgG1, -IgG3, and -IgG4. The columns represent the median OD values of IgG1, IgG2, IgG3, and IgG4 in the cohorts. OD = optical density; NHD = normal healthy donor. Percentage distribution of anti-β2GP1-IgG subclasses in children and patients with APS (f). Children harbour significantly less anti-β2GP1-IgG2 than patients with APS but more anti-β2GP1-IgG3. APS = antiphospholipid syndrome.
Figure 3: OD values of C3c (a) bound to anti-β2GP1 and OD values of anti-β2GP1-IgM (b) in sera of patients with APS and healthy children. Anti-β2GP1 of healthy children displayed significantly more bound C3c than that of patients with APS ($p = 0.00007$). This suggests a more efficient clearance of β2GP1-immune complexes in the healthy children. Anti-β2GP1-IgM OD values of patients with APS were significantly higher than those of healthy children. Horizontal lines represent the median OD value of the cohort. APS = antiphospholipid syndrome; OD = optical density.

Table 3: (a) OD values of the anti-β2GP1-IgG subclasses in the investigated cohorts. (b) Comparison ($p$ values) of the OD values of the anti-β2GP1-IgG subclasses.

(a)

<table>
<thead>
<tr>
<th></th>
<th>IgG1 [mOD]</th>
<th>IgG2 [mOD]</th>
<th>IgG3 [mOD]</th>
<th>IgG4 [mOD]</th>
</tr>
</thead>
<tbody>
<tr>
<td>NHD</td>
<td>24</td>
<td>64</td>
<td>27</td>
<td>67</td>
</tr>
<tr>
<td>Children</td>
<td>69</td>
<td>77</td>
<td>212</td>
<td>210</td>
</tr>
<tr>
<td>PAPS</td>
<td>30</td>
<td>85</td>
<td>45</td>
<td>72</td>
</tr>
<tr>
<td>SAPS</td>
<td>28</td>
<td>74</td>
<td>55</td>
<td>173</td>
</tr>
<tr>
<td>APS</td>
<td>29</td>
<td>81</td>
<td>48</td>
<td>106</td>
</tr>
</tbody>
</table>

(b)

<table>
<thead>
<tr>
<th></th>
<th>IgG1</th>
<th>IgG2</th>
<th>IgG3</th>
<th>IgG4</th>
</tr>
</thead>
<tbody>
<tr>
<td>Children versus NHD</td>
<td>&lt;0.0001</td>
<td>n.s.</td>
<td>&lt;0.0001</td>
<td>n.s.</td>
</tr>
<tr>
<td>Children versus PAPS</td>
<td>&lt;0.0001</td>
<td>n.s.</td>
<td>&lt;0.001</td>
<td>n.s.</td>
</tr>
<tr>
<td>Children versus SAPS</td>
<td>&lt;0.001</td>
<td>n.s.</td>
<td>&lt;0.001</td>
<td>n.s.</td>
</tr>
<tr>
<td>APS versus NHD</td>
<td>n.s.</td>
<td>n.s.</td>
<td>n.s.</td>
<td>n.s.</td>
</tr>
<tr>
<td>PAPS versus SAPS</td>
<td>n.s.</td>
<td>n.s.</td>
<td>n.s.</td>
<td>n.s.</td>
</tr>
</tbody>
</table>

OD = optical density 450 nm; PAPS = primary antiphospholipid syndrome; SAPS = secondary antiphospholipid syndrome; APS = PAPS and SAPS; NHD = normal healthy donors; n.s. = not significant.

plus two times the SD of the considered subclass were defined as seropositive (Supplementary Figure 1). Applying this cutoff, we observed that the healthy children had high levels of the IgG1 and IgG3 subclasses, with 13 (100%) and 8 (62%) seropositive samples, respectively. In contrast, only 7 (39%) and 0 (0%) sera of patients with APS were seropositive for IgG1 and IgG3, respectively. APS showed seropositivity of anti-β2GP1-IgG2 in 7 cases (39%), whereas the cohort of healthy children contained just 2 seropositive sera (15%). Just few samples were seropositive for anti-β2GP1-IgG4 in each cohort, representing a range from 8% (PAPS) to 17% (SAPS).

To determine the subclass distribution within the cohorts, we calculated the contribution of the mean OD values of each individual subclass to the sum.

As shown in Figures 2(e) and 2(f), we saw that for the healthy children IgG3 was the most abundant subclass with a contribution of 40% (IgG3 > IgG1 ($p < 0.001$) and IgG3 > IgG2 ($p < 0.01$)). In contrast, IgG2 dominated the subclass profile for NHD (46%; IgG2 > IgG1, IgG2 > IgG3, and IgG2 > IgG4 ($p < 0.0001$)) and patients with APS (38%; IgG2 > IgG1 ($p < 0.0001$), IgG2 > IgG3 ($p = 0.001$)). The contribution of IgG1 ranged from 12% (SAPS) to 17% (NHD) and that of IgG4 from 21% (PAPS) to 35% (SAPS). Comparing the contribution of the individual subclasses between the study cohorts, we observed that children showed significantly less IgG2 than patients with APS and NHD ($p < 0.001$; $p < 0.0001$) and a higher IgG3 content than NHD and patients with APS ($p < 0.0001$; $p < 0.001$). No significant difference between the cohorts was to be seen for the contributions of IgG1 and IgG4.

3.3. Analysis of IgM, C1q, and C3c Associated to Anti-β2GP1-IgG. Next we analysed potential β2GP1-binding proteins in the sera of healthy children and adults with APS. The signals for C1q were low and did not differ significantly between the cohorts.

However, the ELISAs detecting C3c bound to anti-β2GP1 and anti-β2GP1-IgM showed highly significant differences between healthy children and adults with APS. Whereas C3c-anti-β2GP1 adducts were increased in healthy children ($p < 0.0001$; Figure 3(a)), anti-β2GP1-IgM was decreased ($p < 0.001$; Figure 3(b)).
Table 4: Number (frequency) of seropositive samples of the anti-β2GP1-IgG subclasses.

<table>
<thead>
<tr>
<th></th>
<th>IgG1 (&gt;2,9 mOD)</th>
<th>IgG2 (&gt;7,9 mOD)</th>
<th>IgG3 (&gt;8,5 mOD)</th>
<th>IgG4 (&gt;31,9 mOD)</th>
</tr>
</thead>
<tbody>
<tr>
<td>NHD (n = 17)</td>
<td>0 (0%)</td>
<td>0 (0%)</td>
<td>1 (6%)</td>
<td>1 (6%)</td>
</tr>
<tr>
<td>Children (n = 13)</td>
<td>13 (100%)</td>
<td>2 (15%)</td>
<td>8 (62%)</td>
<td>2 (15%)</td>
</tr>
<tr>
<td>PAPS (n = 12)</td>
<td>6 (50%)</td>
<td>5 (42%)</td>
<td>0 (0%)</td>
<td>1 (8%)</td>
</tr>
<tr>
<td>SAPS (n = 6)</td>
<td>1 (17%)</td>
<td>2 (33%)</td>
<td>0 (0%)</td>
<td>1 (17%)</td>
</tr>
<tr>
<td>APS (n = 18)</td>
<td>7 (39%)</td>
<td>7 (39%)</td>
<td>0 (0%)</td>
<td>2 (11%)</td>
</tr>
</tbody>
</table>

OD = optical density 450 nm; PAPS = primary antiphospholipid syndrome; SAPS = secondary antiphospholipid syndrome; APS = PAPS and SAPS; NHD = normal healthy donors. The cut-off for seropositivity was calculated from the mean OD of the corresponding anti-β2GP1-IgG subclass of the NHD cohort and its standard deviation (SD) according to the formula: mean OD[\text{NHD}] + 2 × SD[\text{NHD}]. The respective cut-off OD values are indicated in parentheses below the anti-β2GP1-IgG subclasses.

Table 5: Cohorts studied with the lectin ELISA.

<table>
<thead>
<tr>
<th>Cohort</th>
<th>Abbreviation</th>
<th>Number of samples</th>
</tr>
</thead>
<tbody>
<tr>
<td>Patients with primary APS</td>
<td>PAPS</td>
<td>12</td>
</tr>
<tr>
<td>Patients with APS and SLE as underlying disease</td>
<td>SAPS</td>
<td>10</td>
</tr>
<tr>
<td>Patients with SLE without symptoms of APS harbouring circulating aPL</td>
<td>SLE + aPL</td>
<td>15</td>
</tr>
<tr>
<td>Asymptomatic carriers of aPL</td>
<td>aaPL</td>
<td>6</td>
</tr>
<tr>
<td>Healthy children with circulating anti-β2GP1</td>
<td>—</td>
<td>16</td>
</tr>
</tbody>
</table>

APS = antiphospholipid syndrome; aPL = antiphospholipid antibodies; SLE = systemic lupus erythematosus.

3.4. Analysis of the Sialylation of Anti-β2GPI. We analysed the glycosylation of anti-β2GPI-IgG, focusing on its sialylation since highly sialylated Fc-parts of IgG are known to cause rather anti-inflammatory responses [30]. We measured the binding of the sialic acid specific lectin SNA to the anti-β2GPI-IgG. As shown in Table 5, we employed five cohorts of individuals: 12 patients with PAPS, 10 patients with SAPS (SLE as primary disease), 15 patients with SLE who did not show clinical sign of APS but with aPL in their sera (SLE + aPL), 6 asymptomatic adult carriers of aPL (aaPL), and 16 healthy children, harbouring aPL.

Analysing the SNA/IgG-ratios (Figure 4), we identified the highest sialylation level in the sera of the healthy children. The significances were rather high and decreased from PAPS (p < 0.0001) to SAPS (p < 0.0001), to SLE + aPL (p < 0.0001), and to aaPL (p < 0.001).

4. Discussion

We analysed two distinct sets of sera. One was used to investigate anti-β2GPI-IgG, its subclasses, and its associated proteins C1q and C3c. For the analysis of the sialylation of anti-β2GPI-IgG, we had to collect fresh sera from patients with APS and the healthy children since the stability of the antibody-glycosylation is unknown. In order to avoid its possible decay by processes like enzymatic degradation, this period has been kept as short as possible and was followed by immediate investigation of the glycosylation.

The antiphospholipid syndrome is characterized by clinical manifestations such as arterial and venous thrombosis or pregnancy-related complications and the detection of specific autoantibodies (lupus anticoagulant, anti-β2GPI- and anticardiolipin-autoantibodies) [1]. In particular anti-β2GPIs are considered responsible for provoking the pathological complications [3]. We first measured the total amount of anti-β2GPI-IgG in the sera of 16 healthy children without clinical manifestations of APS and 46 adult patients with APS to identify individual cohorts seropositive for anti-β2GPI. 17 adult normal healthy donors served as negative control.

The samples of the patients with APS showed higher levels of anti-β2GPI-IgG than those of NHD. We also observed a higher concentration of anti-β2GPI-IgG in the sera of healthy children compared to NHD. This is due to the fact that the detection of these antibodies has been the inclusion criterion for the healthy children examined in this study. Interestingly, the children showed even higher OD values for the putatively pathogenic anti-β2GPI-IgG than the patients with APS. This finding was surprising since the children were clinically healthy and did not show any signs of APS such as thrombosis or thrombocytopenia, complications in which the existence of aPL, especially anti-β2GPI, is made responsible for [7, 33].

High levels of circulating aPL in 61 healthy children, when compared to healthy adults, have already been described by Avčin and colleagues [18]. The anti-β2GPI-IgG was reportedly higher in preschool children with a mean age of 5 years than in adolescents with a mean age of 13.5 years [18]. The strongly elevated circulating anti-β2GPI-IgG in our cohort of healthy children may be explained by the fact that they had an even lower age (mean age was nine months). One possible explanation for this result is the induction of anti-β2GPI by viral and bacterial infections [10, 11] or nutritional exposure to β2GPI [17]. As the immune system in children is still to evolve, they are more susceptible to...
infections than adults and their immune system has to cope with a plethora of viral and bacterial pathogens for the first time in their life. This scenario is prone to result in the stimulation of autoantibody-production. Molecular mimicry of certain pathogens can lead to the production of antibodies which then cross-react with β2GP1 [7, 12–15]. When infants diversify their dietary habits and start to consume products like milk and meat, they can get into contact with nutritional β2GP1. No significant differences were found between the remaining cohorts. Horizontal lines indicate the mean SNA/IgG ratio of the cohort. SNA = Sambucus nigra agglutinin; APS = antiphospholipid syndrome; PAPS = patients with primary APS; SAPS = patients with APS and SLE as underlying disease; SLE + aPL = patients with SLE without symptoms of APS harbouring circulating aPL; aPL = antiphospholipid antibodies; SLE = systemic lupus erythematosus.

**Figure 4:** SNA/anti-β2GP1-IgG ratio in PAPS, SAPS, SLE + aPL, aaPL, and children. Healthy children show a significantly higher SNA/anti-β2GP1-IgG ratio than PAPS, SAPS, SLE + aPL, and aaPL (**p < 0.0001; *** p < 0.00001**). This suggests higher sialylation of the oligosaccharides attached to Fc fragments of the children’s anti-β2GP1. No significant differences were found between the remaining cohorts. Horizontal lines indicate the mean SNA/IgG ratio of the cohort. SNA = Sambucus nigra agglutinin; APS = antiphospholipid syndrome; PAPS = patients with primary APS; SAPS = patients with APS and SLE as underlying disease; SLE + aPL = patients with SLE without symptoms of APS harbouring circulating aPL; aPL = asymptomatic carriers of aPL; aPL = antiphospholipid antibodies; SLE = systemic lupus erythematosus. Tested serum of the healthy children surpassed the threshold set for seropositivity (>mean OD_{NHD} + 2 × SD_{NHD}) and 8 samples (62%) did so in the IgG3-subclass. In the sera of patients with APS, just 7 (39%) and no samples were seropositive for IgG1 or IgG3. In the sera of patients with APS compared to NHD, we observed higher OD levels of anti-β2GP1 in all subclasses, except for IgG4. This result is consistent with the findings of Guerin et al. who reported elevated levels in all anti-β2GP1-IgG subclasses, even including IgG4, in patients with APS compared to a control group of healthy adults [23].

It might be that the elevated absolute OD values of the anti-β2GP1-IgG subclasses are the logical consequence of the generally higher level of anti-β2GP1-IgG in the patients with APS and healthy children who were selected for further testing. Therefore, we also analysed the subclass distribution of this antibody. We first analysed the anti-β2GP1-IgG subclass distribution within the individual cohorts and observed that in the healthy children the IgG3 antibody was most abundant with a contribution of 40% to the total reactivity. This is consistent with the finding that all sera of the healthy children were seropositive in this subclass and that their anti-β2GP1-IgG3 OD values were the highest of all cohorts. This indicates a predominance of this subclass in the healthy children. The remaining anti-β2GP1-IgG subclasses in the sera of the children ranged from 15% (IgG1) over 20% (IgG2) to 25% (IgG4) but the differences between their proportions were not statistically significant. In contrast, in the sera of the patients with APS, the anti-β2GP1-IgG2 subclass had the highest contribution with 38%.

The finding of an elevated proportion of the IgG2 subclass in the patients with APS confirms former published data which also describe a skewing towards anti-β2GP1-IgG2 in APS [19, 23, 26, 27]. The authors detected an even higher contribution of anti-β2GP1-IgG2 (70.2% [26] and 87% [27], resp.). In contrast to our results, the IgG1 subclass was more prominent in the published studies described above. Importantly, in the cohort of the healthy infants IgG2 was relatively low. Therefore one may argue that anti-β2GP1-IgG2 is not just a marker of the disease but is also involved in the pathogenic action of the aPL autoantibodies. When we compared the anti-β2GP1-IgG subclass distribution between the three cohorts, we noticed a significant difference in the distribution concerning the children on one hand and NHD and patients with APS on the other. Healthy children showed a higher proportion of anti-β2GP1-IgG3 and a lower one of anti-β2GP1-IgG2. In NHD and patients with APS, it was the other way round. IgG1 and IgG4 did not significantly differ between healthy children and patients with APS or NHD. The predominance of anti-β2GP1-IgG3 in the healthy children is a hint to an immune response elicited by a proteinaceous antigen [26, 27, 34]. This supports the hypothesis that the harmless anti-β2GP1 in children are infection-induced with pathogen-derived proteins serving as trigger for their production. However, it is still elusive if the IgG3-dominated anti-β2GP1 response, found in the sera of healthy children, contributes to its low pathogenic potential. Alternatively, it can just be ascribed to the proceeding evolution of the infant’s immune system typically associated with an elevated
proportion of IgG antibodies [34, 35]. During childhood, the proportion of IgG continuously decreases to adult ranges usually reached at the age of 10–12 years. However, the above average elevated proportion of the anti-\(\beta_2\)GP1-IgG (40%) in the sera of the healthy children may thus not just be due to the infant immune system. In particular when bearing in mind that normal adult levels of the IgG3 subclass are just around 6% [34, 36].

IgG3 is the IgG-subclass exerting the most effective activation of the complement cascade through the classical pathway [34]. Since complement activation is associated with the pathogenesis of APS [21, 32, 37], the high proportion of complement-activating anti-\(\beta_2\)GP1-IgG3 in the children seems to contradict a possible protective effect against APS of these kinds of antibodies. However, the role of complement in the etiopathogenesis of autoimmune diseases is manifold as the complement components C1q, C2, and C4 contribute to the clearance of postapoptotic, secondary necrotic cells and their deficiency is, therefore, a risk factor for autoimmunity and autoinflammation [38]. In a similar manner, studies in mouse in vivo models of various autoimmune diseases demonstrated that IgG mediated tissue inflammation was blocked in mice deficient in activating FcyRs, although the complement component C3 was still abundantly deposited in the tissue [39–41]. Thus, the contribution of complement deposits in tissue with respect to tissue inflammation remains to be established.

To obtain new insights into the involvement of the complement system in the pathogenesis of APS, we analysed the C1q- and C3c-binding to anti-\(\beta_2\)GP1. We observed that anti-\(\beta_2\)GP1 in the sera of the healthy children and in patients with APS similarly bound C1q, the first molecule of the classical pathway of complement activation [42]. Performing anti-\(\beta_2\)GP1-IgM-ELISAs, we detected significantly higher IgM values in the patients with APS. One would expect a higher C1q-binding of the IgM positive sera, since this immunoglobulin binds and activates C1q more strongly than IgG. However, the children mainly harbour IgG3 autoantibodies, the most potent subclass for the activation of the classical complement pathway [34]. This may compensate for the lower IgM level in their sera and thus be responsible for the similar C1q-binding of both cohorts. This suggests that preconditions for complement activation via the classical pathway are comparable in both cohorts. We conclude that classical complement activation has a minor impact on the differential pathogenesis of aPL in healthy children and patients with APS.

What then is the culprit that induces their pathogenic potential?

Next we quantified the contribution of C3c to the immune complexes bound to \(\beta_2\)GP1. C3c is a domain in the C3b molecule and one of its cleavage products created after cleavage by factor I of C3b [42, 43]. C3c not just represents a marker of complement activation but also serves as an opsonin, improving the clearance of bound targets. We observed significantly more C3c bound to anti-\(\beta_2\)GP1 in the sera of the healthy children than in those of patients with APS. This reflects a better opsonization by C3c of the potentially harmful anti-\(\beta_2\)GP1 immune complexes, anti-\(\beta_2\)GP1-opsonized cells, or cellular fragments in the sera of the healthy children. Consequently, the anti-\(\beta_2\)GP1 complexes can be cleared more efficiently by the powerful complement system of the healthy children. In patients with chronic inflammatory autoimmune diseases, complement is usually low, which may result from null alleles [44] or from complement consumption [38].

But why do these processes not cause inflammation and aberrant coagulation in the children? Apart from anti-\(\beta_2\)GP1-IgG subclass distribution and distinct ability to activate the complement system, it is likely that other properties of the anti-\(\beta_2\)GP1 in the sera of children contribute to their low pathogenicity. We hypothesized that an increased sialylation of the anti-\(\beta_2\)GP1-IgG in healthy individuals renders them harmless for their hosts. IgG is usually N-glycosylated at the CH2-domain of its Fc-part. This N-glycan crucially modifies the effector functions the IgG molecule is endowed with [45–47]. This biantennary, core-fucosylated oligosaccharide carries a variable amount of sugar monomers attached to both arms [28, 48]. IgG antibodies terminating with sialic acids on one or both antennae have been shown to behave anti-inflammatorily after infusion of IVIG [29, 31]. The signal exerted by the terminal sialic acid is dominant since the fraction of sialylated IgG in healthy plasma is just 4 to 14% [28, 46, 49].

In mice, these effects are mainly ascribed to decreased binding of sialylated IgG to activating Fcy-receptors and an upregulation of the inhibitory FcyRIIB [29, 31]. We observed that anti-\(\beta_2\)GP1-IgG isolated from sera of healthy children was significantly more sialylated than that from symptomatic patients with APS. One may predict that these autoantibodies, in complex with \(\beta_2\)GP1, show low affinity for activating FcyRs and, therefore, do not provoke proinflammatory and procoagulant effector functions. FcyRIIA is an example for an activating receptor. It is located on a plethora of cells. On the surfaces of platelets, it mediates activation by immune complexes. The activation is accompanied by platelet aggregation and the release of mediators such as the prothrombotic Thromboxane A2 [50]. Consecutively, this cascade can result in thrombus formation [51, 52]. Sialylated anti-\(\beta_2\)GP1-IgG is unable to bind to and to activate platelets via FcyRIIA.

Further mechanisms involved in thrombus formation in APS comprise the interaction of anti-\(\beta_2\)GP1 with various pattern recognition receptors [53] and its interference with regulatory functions of molecules involved in the coagulation process (protein C, prothrombin, or tissue factor [52, 54]). Direct binding to \(\beta_2\)GP1 on endothelial cells (EC) shifts their phenotype towards coagulation and inflammation [54]. Considering that Fcγ-R-binding is involved in many of these mechanisms, it seems likely that a high sialylation of anti-\(\beta_2\)GP1-IgG leads to a low pathogenicity of these autoantibodies and does not cause APS. Therefore, we conclude that the higher sialylation of anti-\(\beta_2\)GP1-IgG in the sera of the healthy children plays a major role in protecting them from typical complications of APS. This hypothesis is further supported by the observation that antiphospholipid antibodies from sera of asymptomatic adult carriers are more sialylated than those from patients with APS. A high
sialylation status of antiphospholipid antibodies seems to prevent the development of symptoms of APS also in these individuals.

We conclude that a plethora of factors contributes to the low pathogenicity of anti-β2GPI-IgG in the sera of healthy children and asymptomatic adults. We observed a distinct spectrum of IgG subclasses compared to patients with APS and a better binding of C3c indicating a more efficient clearance of potentially harmful anti-β2GPI. The hypersialylation of the anti-β2GPI-IgG is prone to ameliorate their inflammatory and procoagulatory effects in healthy carriers, whereas hyposialylation and inflammation mutually precipitate the antiphospholipid syndrome.

Disclosure

The present work was performed in fulfillment of the requirements for obtaining the degree “Dr. med.” for Christoph Fickentscher.

Conflict of Interests

The authors declare that there is no conflict of interests regarding the publication of this paper.

Authors’ Contribution

Laura Andreoli and Martin Herrmann contributed equally to senior authorship.

Acknowledgments

This project was supported by the German Research Foundation (training research programme of the collaborative research center 643), by the Emerging Fields Initiative (EFI) of the FAU Erlangen-Nuremberg, and by the K. und R. Wucherpfennig-Stiftung.

References


