Levels of total IgA and IgA subclasses in the serum of chronic rhinosinusitis patients

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Abstract
Background: Different inflammatory mechanisms take part in the immunopathogenesis of chronic rhinosinusitis (CRS). Immunoglobulin (Ig) A is the first-line defense in the airway tracts and other mucosal sites, but little is reported regarding its serum level in CRS patients. The purpose of current study is to determine the serum levels of total IgA, and its subclasses (IgA1, and IgA2) in CRS with nasal polyps (CRSwNP), CRS without nasal polyps (CRSsNP), and control groups.

Methods: In this case-control study the serum levels of total IgA and IgA subclasses were determined by Nephelometry and ELISA methods, respectively. The difference of the median concentrations was analyzed with the Kruskal-Wallis test. Collected data were analyzed using SPSS and presented by GraphPad Prism software.

Results: A total of 10 CRSwNP patient, 10 CRSsNP patients and 10 healthy controls participated in our study. The mean age of the groups were 38.2±12.6, 25.6±10.54, and 30.1±9.5, respectively. The concentrations of total IgA were 156(120-165), 165 (149-173), and 172 (152.8-184.3) mg/dl, respectively. The concentrations of IgA1 were 107 (77.9-169.9), 156.1(112.8-175.6), and 130.4 (118.8-175.2) mg/dl, respectively. The concentrations of IgA2 were 26.11 (18.41-38.11), 26.96 (15.48-38.39), and 23.2 (18.42-31.78) mg/dl, respectively. There was no significant difference in total IgA (p=0.120), IgA1 (p=0.397) and IgA2 (p=0.925) serum levels among three groups.

Conclusion: Our study showed there is no difference in total IgA and IgA subclasses in the serum of CRS patients in comparison to healthy controls.

Keywords: Chronic Rhinosinusitis, Nasal polyps, CRSsNP, CRSwNP, Serum IgA, Isotype switching

Introduction
Chronic Rhinosinusitis (CRS) indicates a group of diseases characterized by permanent inflammation of paranasal sinuses. CRS is diagnosed on the basis of clinical symptoms, CT scan and endoscopic examinations (1) and is divided into two main groups depending upon the presence of nasal polyps in sinonasal tracts, including CRS with nasal polyps (CRSwNP) or CRS without nasal polyps (CRSsNP) (2). Although the etiology of CRS still remains to be fully understood, usually the dominant local response in CRSsNP patients is neutrophil accumulation whereas hallmark of local response in CRSwNP is an eosinophilic pattern (3). Accumulation of inflammatory cells such as eosinophils, mast cells, neutrophils, macrophages, B and T lymphocytes, and plasma cells are

↑ What is “already known” in this topic:
Although IgE levels increase in the sera of patients with nasal polyps, and most studies have examined its rate and role in these patients, IgA has not been studied as the most abundant antibody after IgG in sera of patients with CRS.

→ What this article adds:
There is no significant difference in total IgA, IgA1, and IgA2 levels in serum of patients with CRS in comparison to healthy controls. Therefore, serum IgA is not a good marker for diagnosis and prognosis of CRS, consequently local investigation of these biomarkers may result in better interpretation.
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frequently seen in CRS and is directly associated with pathological consequences of the disease (4). However, humoral immune system components, especially immunoglobulins are involved in protection against infectious diseases (5, 6).

Immunoglobin (Ig) A is mainly found in mucosal tissues and fluids in secretory form and plays protective roles by neutralization of microbes and their toxic products. Plasma cells produce dimers of IgA in lamina propria, where they are transported to mucosal surfaces across epithelium barriers (7). Active translocation of IgA is carried out by polymeric immunoglobulin receptor (pIgR) and its extracellular portion, known as secretory component (SC) which protects IgA from proteolytic degradation. Human IgA has two subclasses named as IgA1 and IgA2 which are differently expressed in mucosal surfaces, and their expression is thought to be controlled by several cytokines such as TGF-β1, BAFF, and APRIL (8). Although both subclasses are nearly detected in equal levels in small intestine, IgA1 is largely produced in airway tracts, and IgA2 is profoundly expressed in colon (9). Previous studies confirm that IgE level is higher in the airway tracts, and IgA2 is profoundly expressed in colon (9). Previous studies confirm that IgE level is higher in the airway tracts, and IgA2 is profoundly expressed in colon (9).

In this regard, in the present study, we aimed to compare the serum levels of total IgA, IgA1, and IgA2 in CRS patients. In limited studies (9), Previous studies confirm that IgE level is higher in the airway tracts, and IgA2 is profoundly expressed in colon (9). Previous studies confirm that IgE level is higher in the airway tracts, and IgA2 is profoundly expressed in colon (9). Previous studies confirm that IgE level is higher in the airway tracts, and IgA2 is profoundly expressed in colon (9). Previous studies confirm that IgE level is higher in the airway tracts, and IgA2 is profoundly expressed in colon (9). Previous studies confirm that IgE level is higher in the airway tracts, and IgA2 is profoundly expressed in colon (9). Previous studies confirm that IgE level is higher in the airway tracts, and IgA2 is profoundly expressed in colon (9). Previous studies confirm that IgE level is higher in the airway tracts, and IgA2 is profoundly expressed in colon (9). Previous studies confirm that IgE level is higher in the airway tracts, and IgA2 is profoundly expressed in colon (9).

In this regard, in the present study, we aimed to compare the serum levels of total IgA, IgA1, and IgA2 in CRSwNP and CRSsNP patients and healthy controls.

Methods

Patient selection

This case-control study was carried out in ENT and Head and Neck Research Center and Department of Hazrat Rasoul Akram hospital (IUMS, Tehran, Iran), from January 2016 to September 2017. Ten patients with CRSsNP, 10 CRSwNP patients, and 10 healthy controls who were candidates for anatomical septal deviation or rhinoplasty were recruited in the study. The diagnosis of CRSwNP and CRSsNP was performed according to EPOS criteria (European Position Paper on Rhinosinusitis and Nasal Polyposis), based on clinical examination, history, nasal endoscopic and CT scan scores (1). Demographic and clinical characteristics of participants are summarized in Table 1. Healthy controls were supposed to undergo elective surgery while patients with CRS underwent surgery as a treatment protocol. Written consent was filled by all participants, and the study was approved by the ethical committee of Iran University of Medical Sciences. Before blood sampling, all patients were prohibited to use oral, nasal or systemic corticosteroids or even antibiotics for at least 4 weeks before surgery. Patients who had immunodeficiencies, cystic fibrosis, bronchiectasis, chronic obstructive pulmonary disease, diabetes mellitus, neoplasia, or fungal rhinosinusitis were not included in the study.

Measurement of serum total IgA, IgA1, and IgA2

In brief, after bleeding, serum samples were aliquoted and frozen at -80°C. Total IgA was measured by Nephelometry method using Minineph™ human IgA kit (Minineph, Birmingham, UK) according to manufacturer’s instruction. IgA subclasses including IgA1 and IgA2 were assessed by indirect sandwich ELISA (MyBiosource Company, San Diego, USA) according to the kit manual. The minimum detectable amounts of the ELISA kits were 0.03 and 0.0005 mg/dl, respectively.

Statistical Analysis

The differences of the median concentrations were analyzed by the Kruskal-Wallis test. Data were analyzed using SPSS version 21 and presented using Graph Pad Prism 6 software. P values less than 0.05 was considered significant.

Results

A total of 20 patients participated in our study from January 2016 to September 2017 for measuring the levels of total serum IgA, IgA1, and IgA2, including CRSwNP and CRSsNP patients as well as healthy controls. The mean age of the participants was 38.2±12.6 in CRSwNP patients, 25.6±10.54 in CRSsNP patients, and 30.1±9.5 in healthy controls. There were 10 patients in CRSwNP group of whom 6 were males (60%), and 4 were females (40%). There were 10 in CRSsNP group of whom 7 were males (70%), and 3 were females (30%). There were 10 in control group of whom 5 were males (50%), and 5 were females (50%) (Table 1).

Measurement of total IgA by Nephelometry revealed that the median levels (IQR) of total IgA were 156 (120-165), 165 (149-173), and 172 (152.8-184.3) mg/dl in CRSwNP, CRSsNP, and control groups, respectively (Fig. 1 and Table 2). There was no significant difference in the amount of total IgA among the studied groups. The concentration of IgA1 was measured by ELISA, and they were 107 (77.9-169.9), 156.1 (112.8-175.6), and 130.4 (107.9-169.9) mg/dl in CRSsNP, CRSwNP, and control groups, respectively (Table 2).

Table 1. Characteristics of control and patient groups

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>Controls</th>
<th>CRSsNP</th>
<th>CRSwNP</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Number</td>
<td>10</td>
<td>10</td>
<td>10</td>
<td>-</td>
</tr>
<tr>
<td>Age (y) ± SD</td>
<td>30.1 ± 9.5</td>
<td>25.6 ± 10.54</td>
<td>38.2 ± 12.6</td>
<td>0.019</td>
</tr>
<tr>
<td>CT score</td>
<td>-</td>
<td>7.1 ± 5.36</td>
<td>14.9 ± 4.8</td>
<td>0.003</td>
</tr>
<tr>
<td>Endoscopic score</td>
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<td>0.6 ± 0.7</td>
<td>2 ± 0.67</td>
<td>0.001</td>
</tr>
<tr>
<td>VAS score</td>
<td>-</td>
<td>5.6 ± 2.6</td>
<td>8.4 ± 2.5</td>
<td>0.001</td>
</tr>
</tbody>
</table>

Table 2. The concentrations of total IgA, IgA1, and IgA2 in the serum of studied groups

<table>
<thead>
<tr>
<th>Group</th>
<th>Control</th>
<th>CRSsNP</th>
<th>CRSwNP</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Serum IgA total (mg/dl)</td>
<td>156 (120-165)</td>
<td>165 (149-173)</td>
<td>172 (152.8-184.3)</td>
<td>0.120</td>
</tr>
<tr>
<td>Serum IgA1 (mg/dl)</td>
<td>107 (77.9-169.9)</td>
<td>156.1 (112.8-175.6)</td>
<td>130.4 (118.8-175.2)</td>
<td>0.397</td>
</tr>
<tr>
<td>Serum IgA2 (mg/dl)</td>
<td>26.11 (18.41-38.11)</td>
<td>26.96 (15.48-38.39)</td>
<td>23.2 (18.42-31.78)</td>
<td>0.925</td>
</tr>
</tbody>
</table>

* DATA are presented as median (interquartile range)
(118.8-175.2) mg/dl in CRSwNP, CRSsNP, and control groups sera, respectively (Fig. 2 and Table 2). There was no significant difference in the level of IgA1 in three groups. Similarly, the levels of IgA2 was also measured by ELISA, and its concentration was 26.11 (18.41-38.11), 26.96 (15.48-38.39), and 23.2 (18.42-31.78) mg/dl in CRSwNP, CRSsNP, and control groups, respectively (Fig. 3 and Table 2). There was no significant statistical difference in the level of IgA2 in the studied groups.

**Discussion**

Chronic Rhinosinusitis (CRS) is an inflammatory disease affecting paranasal sinuses (1). IgA is a neutralizing antibody which is predominantly found in mucosal surfaces (7). IgA production particularly occurs within nasal polyps, and its local production is crucial for the activation of eosinophils and mast cells, which in turn influence the inflammation of local mucosal tissues (13). We investigated the levels of IgA and its subclasses in sera of CRS patients in comparison to healthy controls. Results showed that there is no significant difference in the concentration of total IgA and IgA subclasses (IgA1 and IgA2) in the serum of CRSwNP and CRSsNP patients in comparison to healthy controls. Bachert et al. showed that the level of immunoglobulins, especially IgA is locally increased in CRSwNP patients (14). Schleimer et al. showed that local concentrations of total IgA were higher in nasal polyposis patients compared with CRSsNP and control groups (15) and in another study, this author and his colleagues indicated that total IgA levels were increased in tissue extracts from CRS patients but not in lavage fluids (10).

It has been repeatedly reported that IgA is elevated in tissue homogenates from patients with CRSwNP (11, 16); however, it remains to be clarified whether these findings are as a result of increased production of IgA in nasal mucosa or defect in transportation to luminal parts. Mortuaire et al. investigated total IgA levels in blood and nasal secretions of allergic and non-allergic patients. They found that there is no difference between the mentioned two groups. In fact, the atopic status has no effect on the levels of IgA in both serum and nasal secretions (17). However, an old study reported that local IgA production occurred in both allergic and non-allergic CRSwNP patients, resulting in degradation of cytotoxic proteins from eosinophils. In contrast to our study they counted eosinophil numbers in the nasal polyps, nevertheless they acclaimed no relationship between eosinophils and total IgA in serum and nasal secretions (18). In addition to allergic status, the microbiota composition of sinonasal tracts may play a pivotal role in either IgA production or IgA class switching. Van Zele et al. revealed that IgA, IgG, and IgE concentrations against Staphylococcus aureus enterotoxins (SAEs) were significantly increased in tissue extracts, but not in the serum of nasal polyps in comparison to healthy controls (11). Therefore, it is necessary to investigate the microbiome status to find specific IgA and IgA subclasses against super antigens and other microbial agents settled down in sinonasal tracts.

Human individuals have two distinct IgA subclasses. Although both subclasses are nearly detected in equal levels in small intestine, IgA1 is largely produced in airway tracts, and IgA2 is profoundly expressed in the colon (9). In this regard, we aimed to measure IgA1 and IgA2 levels in serum of studied groups. We observed no difference in the protein concentrations of IgA1 and IgA2 in three groups. Schleimer et al. determined IgA1 and IgA2 levels in the samples taken from tissue extracts and nasal lavage by ELISA. Although IgA1 levels were significantly higher in nasal polyps, they reported only a trend toward elevated IgA2 levels. Surprisingly, neither total nor IgA1 or IgA2 levels was elevated in the lavage fluid of CRSwNP patients. These findings further focus on normal mechanisms for export of antibodies to the lumen. So that any disruptions, either Poly-Ig receptor (plgR) dysfunction or insufficient plgR expression, might occur in CRSwNP patients and lead to accumulation of antibodies within nasal tissues (10). The main limitation of our present research was small sample population.

**Fig. 1.** The concentrations of total IgA in serum of control, CRSsNP, and CRSwNP groups.

**Fig. 2.** The concentrations of serum IgA1 in control, CRSsNP, and CRSwNP groups.

**Fig. 3.** The concentrations of serum IgA2 in control, CRSsNP, and CRSwNP groups.
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Therefore, it is suggested that more investigation is necessary with a larger sample size. Eventually, for future studies, we suggest to evaluate local total IgA and IgA subclasses. Furthermore, evaluation of the concentration of the cytokines involved in IgA production such as TGF-β, BAFF, and APRIL is highly recommended. Furthermore, it is valuable to investigate local IgA class switching mechanism by molecular methods such as real-time PCR and the number of IgA producing B cells and plasma cells by flowcytometry and Immunohistochemistry methods. Future studies on the concentrations and the pattern of local IgA and IgA subclasses, determining the mechanisms by which IgA class switching occurs and IgA is secreted at sinonasal mucosa, as well as measurement of the ratio of IgA1 to IgA2 subclasses may develop new insights to target IgA and IgA subclasses in the treatment of CRS patients.

Conclusion

Our study showed there is no significant difference in total IgA, IgA1, and IgA2 levels in serum of CRSwNP and CRSsNP patients in comparison to healthy controls. Hence, serum IgA is not a good marker for diagnosis and prognosis of CRS.

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Conflict of Interests

The authors declare that they have no competing interests.

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