Axenfeld-Rieger anomaly and corneal endothelial dystrophy: a case series

Anomalia de Axenfeld-Rieger e distrofia corneana endotelial: uma série de casos

Mariana Borges Oliveira¹, Roberto dos Santos Mitraud², Riuitiro Yamane³

ABSTRACT

Report of a study of a family with a remarkable combination of endothelial corneal dystrophy, and anterior chamber dysgenesis classified as Axenfeld-Rieger anomaly. A 56 year-old female had blurred vision complaints. A slit lamp evaluation showed a bilateral posterior embryotoxon and guttata with corneal edema, Descemet membrane’s folds, characterizing Fuchs endothelial dystrophy. A 27 year-old daughter was asymptomatic. The biomicroscopic exam disclosed a bilateral nasal and temporal posterior embryotoxon and a central guttata. Gonioscopy showed iridocorneal strands with a prominent Schwalbe’s line. The intraocular pressure was normal. Her sister, a 25 year-old female presented very similar abnormalities along with her brother (22 years old). A 19 year-old female sister presented the iridocorneal angle alterations, without Fuchs endothelial dystrophy, until this moment. This paper presents a family with a central cornea guttata or Fuchs dystrophy associated with Axenfeld-Rieger anomaly. Two different inherited diseases appearing together in an entire family may suggest a single molecular and genetic etiology, with high penetrance.

Keywords: Eye abnormalities/genetics; Fuchs endothelial dystrophy; Anterior eye segment/abnormalities; Cornea/pathology; Syndrome; Case reports

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INTRODUCTION

Axenfeld-Rieger (AR) syndrome (MIM 180500) is an inherited autosomal dominant disorder with high penetrance and variable expressivity. It is included in a heterogeneous spectrum of overlapping phenotypes referred to as an anterior segment dysgenesis (ASD) and is characterized by posterior embryotoxon, iridocorneal strands, iris changes associated with systemic signs, like oligodontia, microdontia, cranial and facial defects. The AR anomaly presents only ocular changes. Glaucoma is associated with half of the cases in each subtype, with exception of the isolated posterior embryotoxon. There are other genetically related phenotypes that share many features with AR syndrome, like iridogoniodysgenesis (IGD) anomaly and syndrome as well as familial glaucoma iridogoniodyplasia. For this reason, some authors suggest that they might be considered part of the same group. Three genetic loci are linked to AR syndrome and related phenotypes, on chromosomes 4q25 (PITX2 gene), 6p25 (FOX-C1 gene), 13q14 (unrecognized gene) and recently, it was also linked to chromosome 11p13 (PAX6 gene). These genes are involved in the regulation of eye embryogenesis, and codify transcription factors expressed during this phase. Shields has postulated that both the ocular and systemic phenotypes result from the disruption of migration and/or the differentiation process of the neural crest-derived tissues, which include the corneal endothelium, the anterior chamber angle (trabecular meshwork, Schlemm’s canal) and the iris.

Fuchs endothelial dystrophy of the cornea (FECD, MIM 136800) is a relatively common and slowly progressive corneal dystrophy; adult-onset corneal degeneration begins with an axially (central or paracentral) arranged guttata. It becomes clinically evident when epithelial and stromal edemas produce painful bullae, which can lead to corneal swelling and severely reduced vision. The pathogenesis of Fuchs dystrophy is not entirely understood yet. There are clues suggesting a disturbance of regulation of apoptosis, while other studies suggest that the disturbance occurs on terminal differentiation of the neural crest-derived corneal endothelium. In this report, we describe a family (mother and four children) having a combination of endothelial corneal dystrophy with anterior segment dysgenesis classified as Axenfeld-Rieger anomaly.

Case Reports

The family was examined in order to find clinical signs of anterior chamber dysgenesis, cornea guttata or Fuchs endothelial dystrophy. The findings are listed on Table 1 and their heredogram is shown on Figure 1.

Figure 1: Heredogram exhibiting four of six members affected with both posterior embryotoxon and corneal central guttata or Fuchs dystrophy (black) and, one member affected only with anterior segment alterations (gray).

Case 1

A 56-year-old female (mother) had blurred vision complaints. Her best visual acuity was 20/25 in the right eye and 20/50 in the left eye. The biomicroscopic exam showed bilateral posterior embryotoxon (Figure 2) and guttata with corneal edema and Descemet membrane’s folds, characterizing Fuchs endothelial dystrophy. Gonioscopic view was not possible because of the corneal edema. The specular microscopic examination revealed that the endothelial cells were almost absent, and it was not possible to execute the cell count (Figure 3). The intraocular pressure (applanation tonometry) was 12 mmHg in the right eye and 14 mmHg in the left eye. The fundoscopy was normal and revealed no signs of glaucomatous optic nerve damage. The rest of the ocular examination was normal. This case was indicated to corneal transplant.

Case 2

A 27-year-old Caucasian female patient (daughter) had the best visual acuity equal to 20/20 in both eyes. Slit lamp evaluation disclosed a bilateral nasal and temporal posterior embryotoxon (Figure 4). No systemic signs were observed. The intraocular pressure (applanation tonometry) was 15 mmHg in the right eye and 16 mmHg in the left eye. Gonioscopy showed iridocorneal strands with a prominent Schwalbe’s line (Figure 5). Specular microscopic examination demonstrated bilateral dark bodies (guttata) and the endothelial cells showed pleomorphism and polymegathism (Figure 6). Central corneal thickness was 571 μm in the right eye and 587 μm in the left eye. The fundoscopy was normal and revealed no signs of glaucomatous optic nerve damage.
Table 1

<table>
<thead>
<tr>
<th>Case</th>
<th>Age</th>
<th>Posterior Embryotoxon</th>
<th>Cornea</th>
<th>Iris</th>
<th>Glaucoma</th>
</tr>
</thead>
<tbody>
<tr>
<td>Case 1 Mother</td>
<td>59</td>
<td>+</td>
<td>Fuchs' Dystrophy</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>Father</td>
<td>61</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>Case 2</td>
<td>27 female</td>
<td>+</td>
<td>Central Guttata</td>
<td>Mild Corectopy</td>
<td>-</td>
</tr>
<tr>
<td>Case 3</td>
<td>25 female</td>
<td>+</td>
<td>Central Guttata (mild)</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>Case 4</td>
<td>23 male</td>
<td>+</td>
<td>Central Guttata (mild)</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>Case 5</td>
<td>20 female</td>
<td>+</td>
<td>-</td>
<td>Mild Corectopy</td>
<td>-</td>
</tr>
</tbody>
</table>
12 mmHg in the left eye. Specular microscopic examination was normal as well as the rest of the ocular examination. The fundoscopy was normal and revealed no signs of glaucomatous optic nerve damage.

**DISCUSSION**

This is a remarkable case series in which a family, including two generations presented Fuchs endothelial dystrophy or central cornea guttata associated with Axenfeld-Rieger anomaly. The trait appeared to be autosomal dominant with high penetrance (100%), considering that all siblings inherited the abnormalities of the mother. Interestingly, Hwang et al. described a case in which mother and son had a combination of corectopy, congenital corneal opacities and cornea guttata (MIM 608484), with an overlap with Peters anomaly (characterized by central corneal opacities and absence of Descemet membrane), but it was not associated with posterior embryotoxon, polycoria or iris stromal hypoplasia.

There are two genes encoding transcription factors and three loci that are associated with the Axenfeld-Rieger malformation. The gene PITX2 (“pituitary homeobox transcription factor 2”, RIEG1) was cloned on chromosome 4p25 and regulates the expression of other genes during embryonic development. Several pathological mutations (deletions, translocations or spot mutations) in PITX2 have been described causing different phenotypes, including AR malformation, iridogoniodygenesis, iris hypoplasia and rarely Peters’ anomaly. A second gene associated with AR syndrome is FOXC1 (“Forkhead Box C1”), on chromosome 6p25. During the embryogenesis, FOXC1 is expressed in craniofacial structures, in periocular and pre-endothelial mesenchyma, sclera and cornea and further, in the post-fetal phase, in trabecular meshwork, iris and conjunctival epithelium. The mutations include duplications, insertions, deletions and missense types affecting the DNA ligation stability, activation of specific genes and transcriptional regulation. These mutations produce a very large spectrum of phenotypes of anterior segment dysgenesis, mild to severe form, including or not systemic manifestations.

Other locus for Rieger syndrome was mapped on 13q14 (RIEG2), but no gene was identified.

Besides the genes described previously, there are evidences of the involvement of PAX6 gene (“Paired Box
has been widely studied and discussed. It is necessary to investigate this hypothesis. Further studies are needed to explain the relationship among these overlaps.

In the Fuchs dystrophy, the mechanism of progressive dysfunction of corneal endothelium and the collagen deposition remains unknown, and it is not clear whether it is a primary or secondary effect. Several theories have been proposed to explain the etiology of FECD. Biswas et al. described missense mutations in COL8A2, a gene of the α2 chain of type VII collagen in association with FECD, suggesting that this molecule may be related with a disturbance in the terminal differentiation of the neural crest derived endothelial cells. There are studies suggesting that apoptosis plays an important role in endothelial cell degeneration. It is possible that a modification of the basement membrane composition (Descemet membrane) or the loss of contact between endothelial cell and Descemet membrane may cause apoptosis of these cells in Fuchs’ Dystrophy. Further studies are needed to investigate this hypothesis.

The mechanism of the ocular bad development has been widely studied and discussed. Often the factors that result in disturbance of neural crest may act at more than one phase of this process, producing anomalies that involve more than one tissue derived from neural crest.

This case series was the first description of this combination of two ocular abnormalities together (Fuchs dystrophy and Axenfeld-Rieger anomaly) and was reported on world ophthalmology congress in 2006. Later on, Kniestedt et al. showed very similar ocular findings presented in a five generation family with a PITX2 mutation and polymorphism.

**Conclusion**

Two different inherited diseases appearing together in an entire family may suggest a single molecular and genetic etiology, but additional studies are necessary to explain the relationship among these overlapping phenotypes and their inheritance. Perhaps, in future, it would be possible to develop alternative therapeutic tools for the treatment of Fuchs dystrophy and glaucoma, based on the knowledge of their etiology.

**References**


**Resumo**

Relato dos resultados do estudo de uma família com dystrofia corneana endotelial associada a uma disgenesia do segmento anterior, a anomalia de Axenfeld-Rieger. Paciente de 56 anos, feminina, queixava-se de visão borrada. O exame na lâmpada de fenda evidenciou embriotoxo posterior bilateral e guta na oclusão do ângulo iridocorneano. A pressão intra-ocular estava normal. Não foram observadas alterações sistêmicas. Sua irmã, 25 anos, sexo feminino, apresentava alterações muito similares, assim como seu irmão (22 anos). Outra irmã de 19 anos apresentava alterações do ângulo iridocorneano, porém sem distrofia de Fuchs, até o presente. Este trabalho relata os resultados de estudo de uma família com córnea guttata central ou distrofia de Fuchs associada à anomalia de Axenfeld-Rieger. Duas doenças hereditárias com expressão simultânea em toda a família sugerem uma etiologia molecular e genética única, com alta penetrância. **Descritores:** Anormalidades do olho/anormalidades; Distrofia endotelial de Fuchs; Segmento anterior do olho/anormalidades; Córnea/patologia; Síndrome; Relatos de casos

**Conclusão**

Duas diferentes doenças hereditárias apareceram juntas em uma família inteira, que sugerem uma única etiologia molecular e genética. No entanto, são necessários estudos adicionais para explicar a relação entre essas anomalias que envolvem mais de um tecido derivado do neural crest.

Este caso foi o primeiro relato dessa combinação de duas anomalias oculares, juntando Fuchs Distrofia e Anomalia de Axenfeld-Rieger, e foi relatado no Congresso Mundial de Oftalmologia em 2006. Mais tarde, Kniestedt et al. mostraram achados similares oculares em uma família de cinco gerações com a mutação e polimorfismo de PITX2.

**Referências**


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