

1 de Dezembro

08h30 | 10h00 – Sala 1

Retina Médica | Medical Retina

Moderadores | Chairs: Maria Luz Cachulo (CHUC), Diogo Cabral (HGO), Luis Mendonça (HB)

CO 3

EYS-ASSOCIATED RETINAL DEGENERATION: NATURAL HISTORY, GENETIC LANDSCAPE AND PHENOTYPIC SPECTRUM

Pedro Nuno Pereira¹, Ricardo Machado Soares², Ana Luísa Carvalho³, Sílvia Simão¹, Jorge Saraiva³, Joaquim Murta¹, Rufino Silva¹, João Pedro Marques¹

(¹Department of Ophthalmology, Centro Hospitalar e Universitário de Coimbra (CHUC), Coimbra, Portugal, ²Department of Ophthalmology, Centro Hospitalar de Vila Nova de Gaia Espinho (CHVNGE), Gaia, Portugal, ³Medical Genetics Unit, Centro Hospitalar e Universitário de Coimbra (CHUC), Coimbra, Portugal)

Introduction: Eyes shut homolog (*EYS*) is the third most frequently mutated gene in inherited retinal diseases (IRDs) worldwide, and the single most commonly mutated IRG gene in Portugal.^{1,2} Despite initially associated with a relatively homogeneous and slowly progressive form of retinitis pigmentosa (RP), recent studies have highlighted its genetic, phenotypic, and clinical heterogeneity.

Purpose: To describe the natural history, genetic landscape, and phenotypic spectrum of *EYS*-associated retinal degeneration (*EYS*-RD)

Methods: Single-center retrospective cohort study conducted at an inherited retinal degeneration (IRD) referral center in Portugal. Patients with biallelic *EYS* variants were invited to participate. Every patient underwent a cross-sectional examination comprising a comprehensive ophthalmologic examination including best-corrected visual acuity (BCVA), dilated slit-lamp anterior segment and fundus autofluorescence (UWF-FAF) imaging; and spectral domain-optical coherence tomography (SD-OCT). Additional information was collected from the patient file. Main outcome measures included clinical/demographic, genetic, and multimodal imaging data. BCVA variation during follow-up was used as an endpoint to describe *EYS*-RD natural history.

Results: Fifty-eight patients (59% males; mean age 52±14 years) from 48 Caucasian families of Portuguese ancestry were included. Twenty distinct *EYS* variants were identified, eight of which are novel. In 32.8% patients, onset of symptoms was in early adulthood (21-30 years). On UWF imaging, 75.0% patients (n=41) were graded as typical, while 25.0% were atypical. Overall, a negative correlation was found between age and BCVA ($r=-0.50$; $p<0.001$), with an average loss of 1.45 letters per year of follow-up. Higher BCVA and larger ellipsoid zone (EZ) widths were larger EZ widths, thus presenting an overall better prognosis.

Conclusions: This study expands the genetic spectrum of *EYS*-RD by reporting 8 novel variants. A high frequency of atypical phenotypes was identified. These patients have better BCVA and larger EZ widths, thus presenting an overall better prognosis.