

# Isolated and syndromic retinal degeneration associated with POC5 variants.

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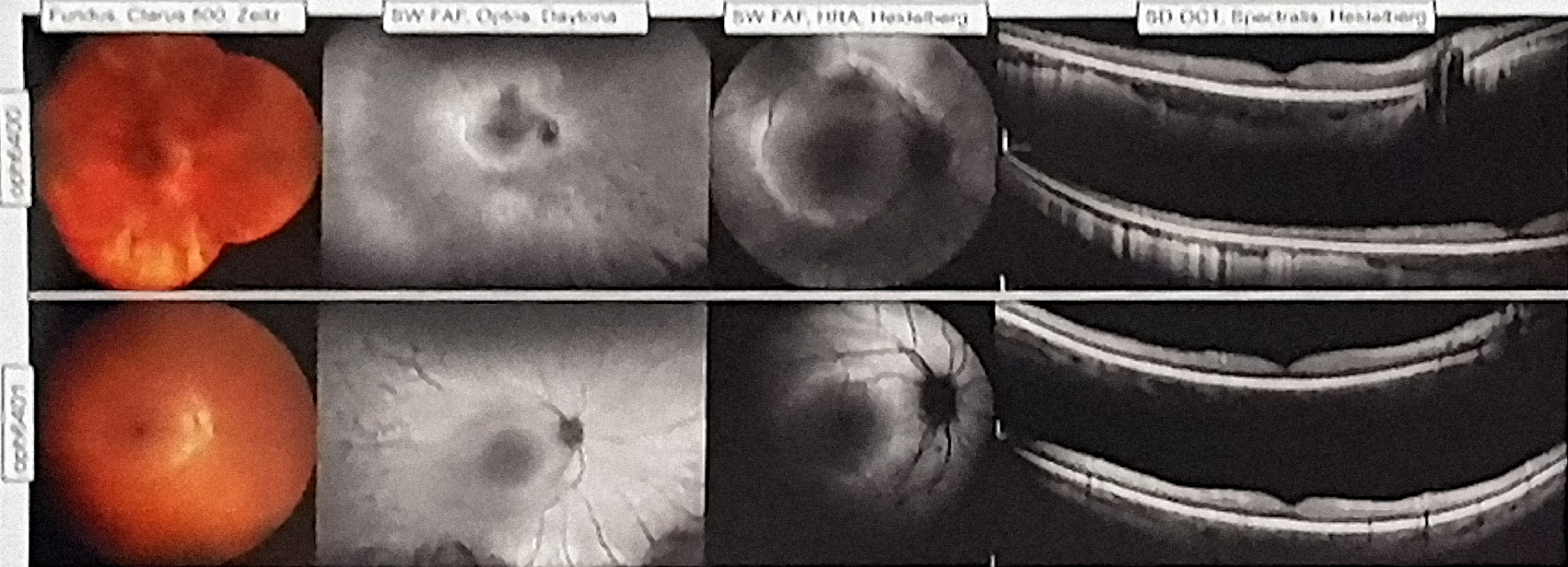
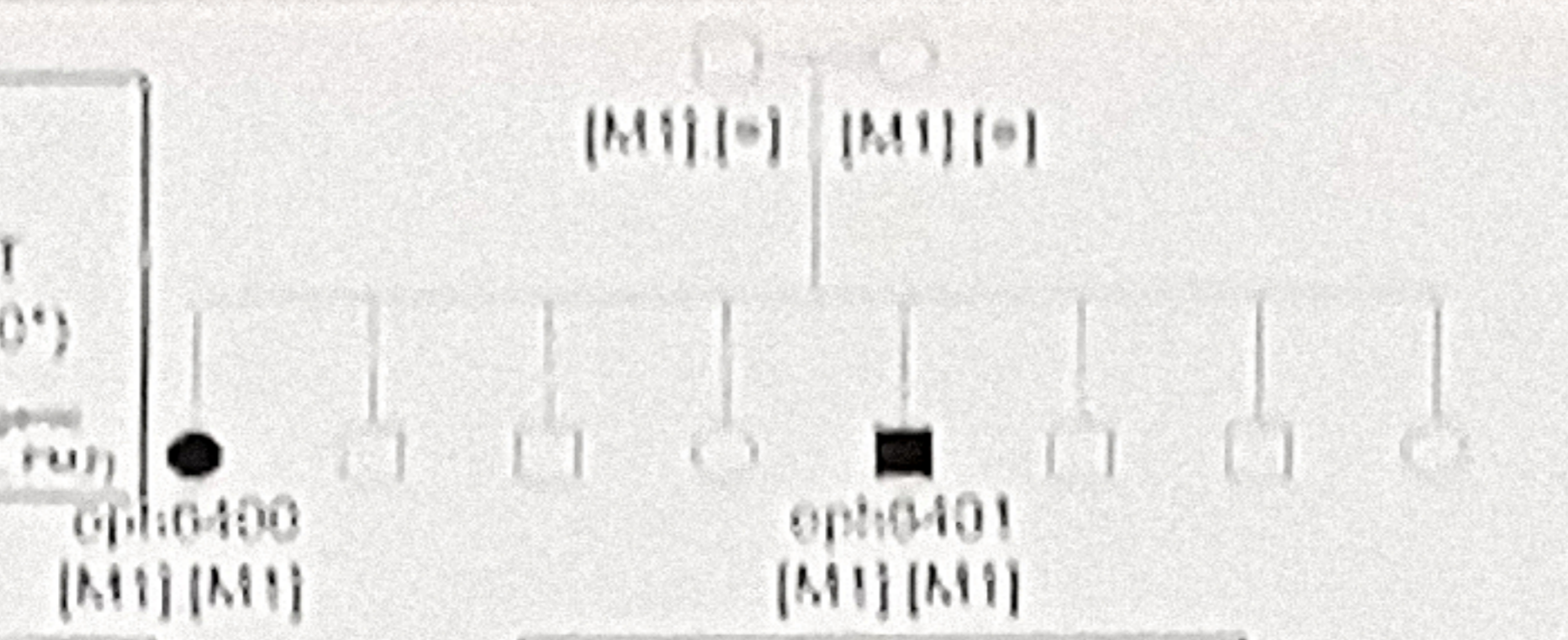
## Purpose:

Biallelic gene defect in *POC5* have been recently reported in association with syndromic rod-cone degeneration in one patient. Systemic findings included low weight, short stature, microcephaly and recurrent glomerulonephritis [1]. The purpose of this study was to report a detailed phenotypic description of French patients harboring *POC5* pathogenic variants.

### oph6400, 17 y.o., night blindness and progressive visual loss

	RO	LE
BCVA, Snellen	20/200	20/200
Refraction, D		
Anterior segment	Posterior subcapsular cataract	

M1:  
*POC5*  
c.568 C>T  
p.(Arg190\*)  
ACMG pathogenic  
(PVS1, PLS1, PM2)

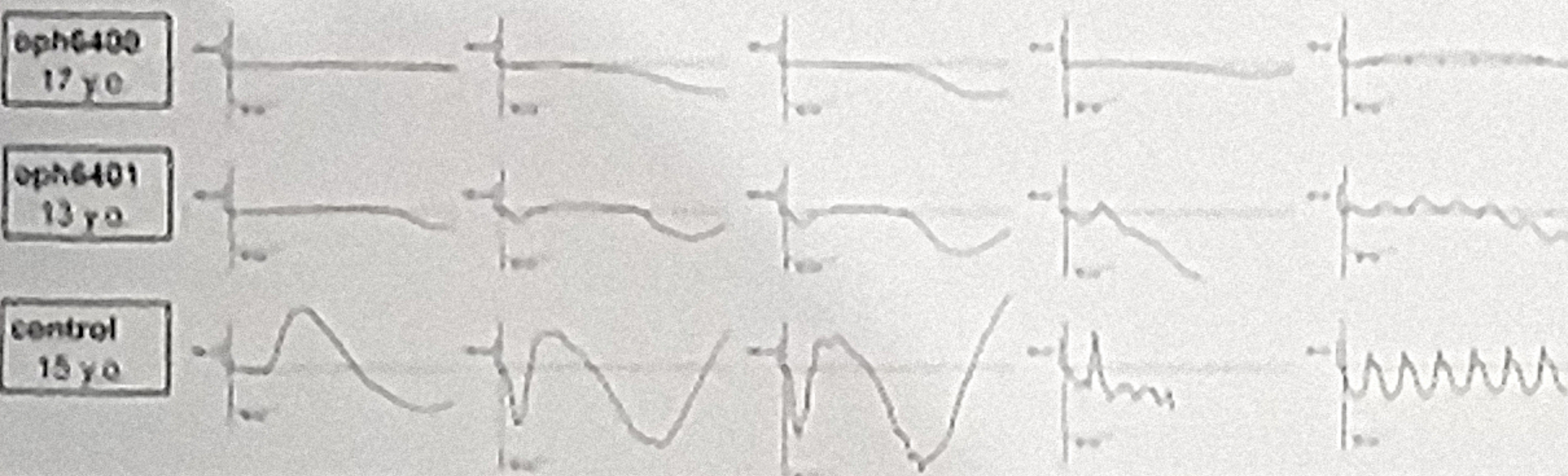


### oph6401, 13 y.o., convergent squint LE, no visual complaints

	RO	LE
BCVA, Snellen	20/125	20/125
Refraction, D	+8. -1.50/20°	+8.50/-1.80/170°
Anterior segment	Normal	

Full-field ERG, Mon Color, Metrovision®

DA 0.01 DA 3.0 DA 10.0 LA 3.0 LA 3.0 flicker



- No systemic involvement
- Low weight (-2DS both)
- Normal PC
- Normal kidney and liver function
- Regular schooling

oph6400 undetectable  
oph6401 rod-cone dystrophy,  
cone-dominated ERG under  
dark-adapted conditions

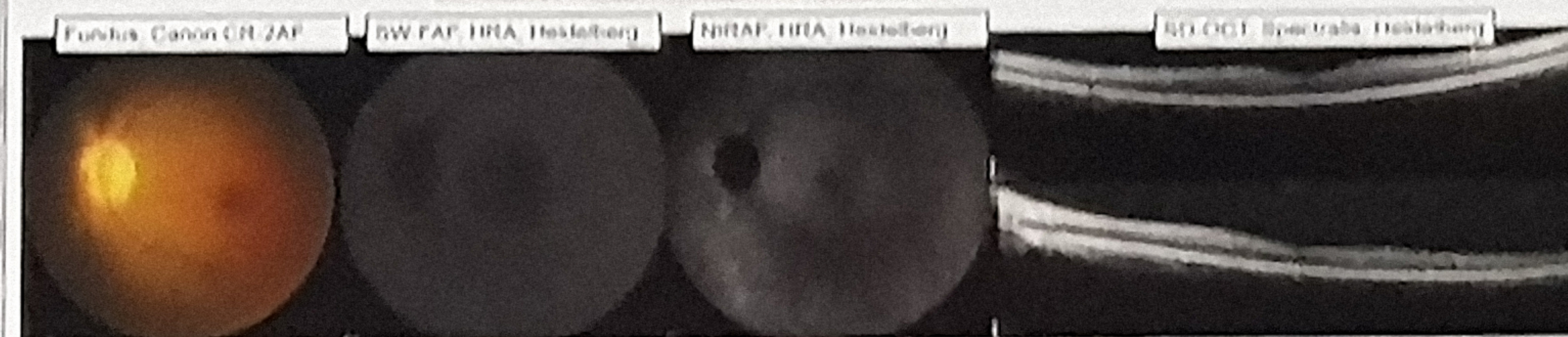
## Methods:

Patients harboring biallelic gene defects in *POC5* were selected from cohorts of Inherited Retinal Degeneration (IRD) patients followed at the Exploration de la Vision et Neuro-Ophthalmologie, CHU de Lille and at the National Reference Center for rare ocular diseases, REPERET, of Quinze-Vingts hospital, Paris. Best corrected visual acuity (BCVA), slit-lamp examination, static and kinetic perimetry, full field electroretinography (ffERG) and multimodal imaging including color photos, infrared reflectance (IRR), short-wave autofluorescence (SWAF) and optical coherence tomography (OCT) were done for all patients. Past medical history was also collected. *POC5* variants were identified applying customized IRD multigene NGS panels. Variant confirmation and familial segregation, when possible was performed by direct Sanger sequencing.

### F5404 CIC09410, 27 y.o., infantile nystagmus, low visual acuity since childhood, light sensitivity

	RO	LE
BCVA, Snellen	20/320	20/400
Refraction, D	-3.0	-2/-1170°
Anterior segment	Normal	

M2:  
*POC5*  
c.616C>T  
p.(Gln206\*)  
ACMG likely pathogenic (PVS1, PM2)



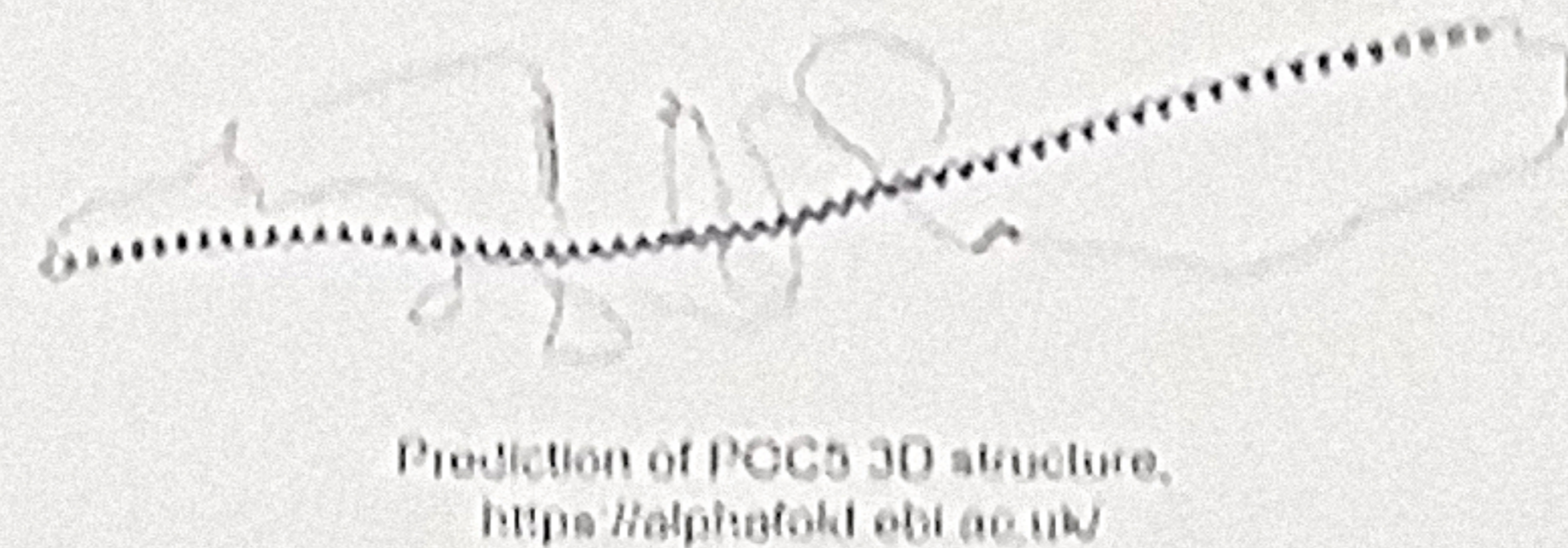
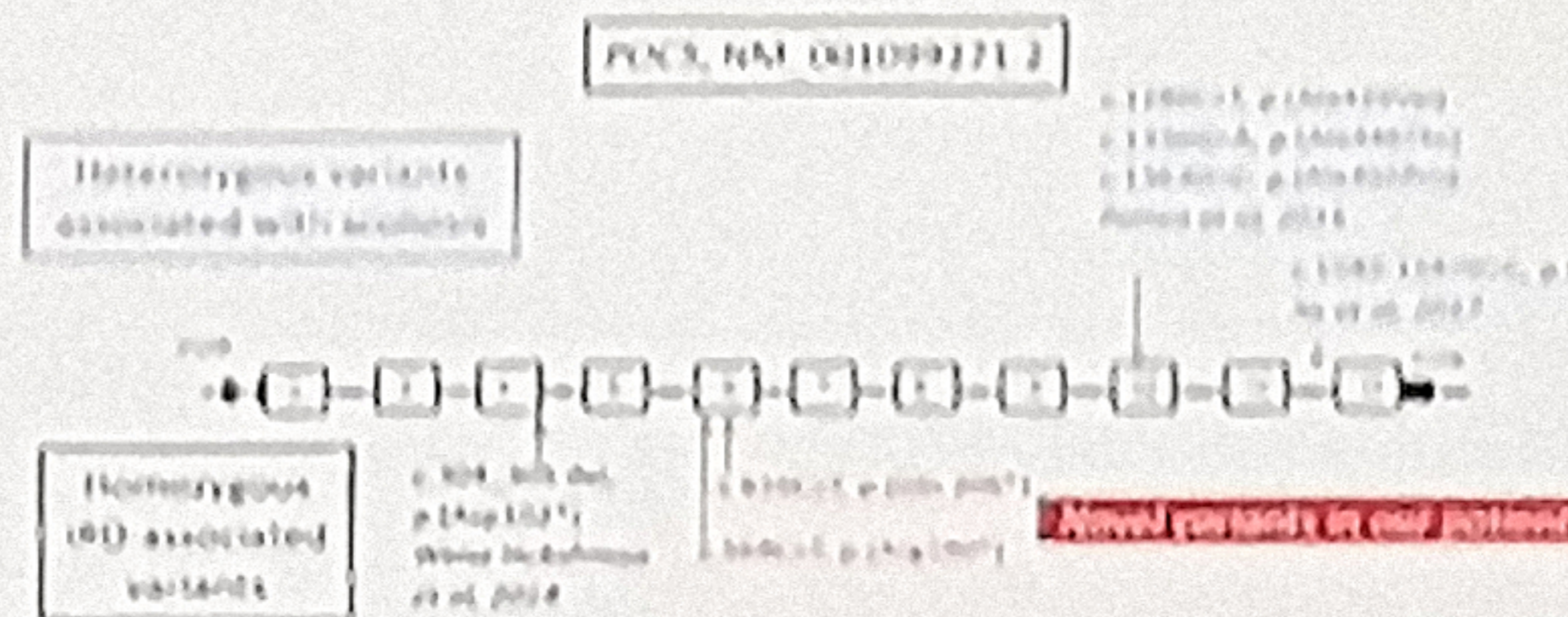
Full-field ERG:  
undetectable dark-  
and light-adapted  
responses

### Systemic features:

- Short stature (1m55) and low weight (46 kg), BMI 10
- Non-autoimmune insulin-requiring diabetes since 13 y.o., high insulin-resistance
- Chronic glomerulonephritis with persistent non-selective massive proteinuria (1,3 g/d) since 18 y.o.

- Cirrhosis (Child A5) with steatosis, hepato- and splenomegaly, mild portal hypertension
- Arterial hypertension
- Oligomenorrhea, polycystic ovary syndrome
- Psoriasis and acanthosis nigricans

## Genotype/phenotype correlations



Prediction of POC5 3D structure,  
<https://alphafold.ebi.ac.uk/>

### *POC5* (CSORF37)

- Ubiquitously expressed, but higher levels in retina, brain and reproductive tissues
- Protein associated with distal part of centrioles, required for centriole elongation and cell cycle progression [1]
- POC5* depleted morphant cells are stacked in S-phase and undergoes p53-mediated cell death [1]
- In photoreceptors, *POC5* is localized in the basal bodies of connecting cilium [2]
- In zebrafish *POC5* morphants outer photoreceptor segments were defective (decreased length) and the retinal function abnormal [2]
- Heterozygous missense variants have been reported in association with scoliosis, but data are still unclear as the Minor Allele Frequency of these variants is high [3]
- Patient reported by Weisz Huberman et al. presented a syndromic phenotype including a rod-cone retinal degeneration, chronic glomerular nephropathy, short stature and microcephaly, recurrent episodes of rhabdomyolysis
- In our patients oph6400 and oph6401 the phenotype was retina-restricted at the age of assessment
- In an older patient CIC09410 the clinical picture was in keeping with previously reported case (syndromic ciliopathy including growth, renal, liver, ovarian abnormalities and rod-cone retinal degeneration)
- These data expand the clinical spectrum of *POC5*-related ciliopathy

### References:

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2. Weisz Huberman M, Aronson A, van Hips E, et al. Whole-mount screening reveals POC5 as a novel gene associated with retinal rod-cone retinal degeneration. *Hum Mol Genet*. 2014 Feb 15;23(4):411-24.
3. Patten SA, Maggipinto KA, Bernard A, et al. Functional variants of POC5 identified in patients with scoliosis. *Am J Hum Genet*. 2015 Mar 2;97(3):414-8.

## Findings

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## Conclusions:

The clinical phenotype of patients harboring biallelic *POC5* defects could be both syndromic and retina restricted. Identification of *POC5* pathogenic variants in patients with inherited retinal degeneration should trigger the systemic work-up for morphological abnormalities, kidney and liver disease, endocrine dysfunction. Careful follow-up is mandatory, as the phenotype might broaden with time.