

# Molecular characterization of retinomas and their role in retinoblastoma pathogenesis

**Hrant Ghazelian<sup>1</sup>**, Clement Hua<sup>1</sup>, Kelly Ferreira Pinto<sup>1</sup>, Francois Radvanyi<sup>1</sup>

<sup>1</sup>Institut Curie, CNRS, UMR144

## Abstract

**Retinoblastoma is a tumor arising in the developing retina** and is the most common intraocular malignancy afflicting children. Despite a good prognosis, current treatments induce side effects and still 30% of the patients have their eye removed surgically. Using multi-omics data, we demonstrated the existence of two distinct retinoblastoma subtypes. Subtype 1, of earlier onset, includes most of the heritable forms. It has few genetic alterations other than the initiating RB1 inactivation and corresponds to differentiated tumors expressing more mature cone markers. By contrast, subtype 2 tumors have frequent recurrent genetic alterations, including MYCN-amplification. They express markers of less differentiated cone together with neuronal/ganglion cell markers intra-tumor heterogeneity. Retinoma, also known as retinocytoma, is a benign, nonproliferative lesion of the retina. It can accompany retinoblastomas or occur alone in patients with germline *RB1* mutations. These tumors are poorly studied for multiple reasons.



A better understanding of retinoma will lead to a better understanding of the pathogenesis of these tumors and hopefully lead to more precise diagnostic tools, efficient therapies and thus avoiding invasive and toxic treatments.

## Background

### About retinoblastoma

- Rare childhood cancer of the developing retina
  - I per 18000 childbirths
  - Institut Curie: French referral center
- A curable cancer if diagnosed early...
  - In high-income countries: ≈100% survival
  - In low-income countries: 30% survival
- but invasive treatments
  - Enucleation ( $\approx 30\%$ )
  - Conservative chemotherapies: Side effects

About retinocytoma

Molecular alterations RB1 loss initiates most of retinoblastoma but it is not sufficient to make it malignant; additional molecular alterations are necessary.



## **Objectives**

1) Determine whether retinoma is the precursor to retinoblastoma or whether it is an abortive progression pathway distinct from that of retinoblastomas,

2) Understand why retinomas have halted their proliferation,

3) Verify that retinoblastomas associated with retinomas belong exclusively to subtype 2, as suggested by our preliminary data.

## **Methods**

- Benign counterpart of retinoblastoma : • seen in 2 % of people with a mutant *RB1* allele
  - seen in 15-20% of surgeries done for retinoblastoma
- Understudied tumor for multiple reasons :
  - very rare tumor (1 per 100000)
  - underdiagnosed
  - most tumors are not biopsied





Retinoblastoma subtypes are identified by the integration of three omics data (transcriptome, methylome and copy-number). A total of 102 retinoblastomas were studied, including 72 tumors with at least two of the three omic approaches. Two classifers were built (from transcriptome or methylation of 9 CpGs) to classify the other samples. These subtypes have distinct clinical and molecular features.

Distinct clinical and molecular features: Subtype 1 < 18 months</p>

- Metastatic potential +
- Intra-tumoral heterogeneity -
- Expression of cone marker, Stemness -
- Genomic instability +
- Mutation -

Subtype 2

- > 18 months
- Metastatic potential +++
- Intra-tumoral heterogeneity +++
- Neuronal/ganglions marker, Stemness +++
- Genomic instability ++



## Why is it important?



Understand the origins of the differences between the two subtypes? Between tumor subpopulations? Comparison the normal?

 $\rightarrow$  A better understanding of a rare tumor, that albeit benign, is frequently associated with a retinoblastoma and/or is a precursor of RB.

 $\succ$  Study the underlying genetic modifications of retinoma and retinoblastoma and deduce the genetic alterations necessary for the progression of retinoma into retinoblastoma

 $\succ$  Eventually help discover therapeutic targets that enable the reprogramming and/or induction of differentiation of Retinoblastoma to retinoma



Acknowledgment

Institut de la Vision : Olivier Goureau Institut Curie : Sandra Majo, Sabine Druillennec, **Celio Pouponnot** 

Pathology : Arnaud Gauthier, Paul Freneaux Pediatric oncology : Yassine Bouchoucha, Isabelle Aerts, Francois Doz **Ophtalmology : Alexandre Matet, Denis Malaise, Livia** 

Lumbroso-Le Rouic, Nathalie Cassoux

#### **Genetics : Jessica Le Gall, Lisa Golmard**

#### References

Aibar S et al. Nat Methods. 2017 PMID: 28991892 Dimaras H et al. Hum Mol Genet. 2008 PMID: 18211953 Eagle RC Jr. Arch Pathol Lab Med. 2009 PMID: 19653710 Gallie BL et al. Br J Cancer. 1982 PMID: 7073943 Liu J et al. Nat Commun. 2021 PMID: 34552068 Pfaff E et al. Acta Neuropathol. 2020 PMID: 31768671 Rushlow et al. Lancet Oncol. 2013 PMID: 23498719 Sampieri K et al. Acta Oncol. 2008 PMID: 18785023 Sampieri K et al. Cancer Sci. 2009 PMID: 19183342

#### Contact

#### Hrant Ghazelian, M2 student hrant.ghazelian@curie.fr

Molecular Oncology team Institut Curie, Paris (France)