

## Commentary: Genetic testing in cases of pediatric cataract

Pediatric cataract is a common cause of childhood blindness. In developing countries like India, it is 10 times more prevalent than in developed countries, where the prevalence is 2.03–2.49 per 10,000 children.<sup>[1,2]</sup> Early diagnosis and timely intervention in pediatric cataract cases improve the visual outcomes significantly.<sup>[3,4]</sup> Identifying the etiology of cataracts in children is also very important for counseling and preventive

public programs. The most common etiology for congenital cataracts in India is *Toxoplasma gondii*, other agents, rubella, cytomegalovirus (CMV), and herpes simplex virus (HSV) (TORCH) infections (33.4%), followed by familial causes (18.3%) and developmental anomalies (10.1%).<sup>[5-7]</sup> Congenital cataract has a lot of phenotypic heterogeneity also, with various genetic and environmental causes to consider. Genetic causes are also responsible for one-fifth of cases of unilateral cataracts and majority of bilateral cataracts in children. Autosomal dominant inheritance is the most common pattern of inheritance, followed by autosomal recessive mode in such cases. The morphology of

a cataract can often guide us toward the etiology of the cataract. Apart from detailed history, appropriate investigations should be done in all congenital cataract cases, including complete blood examination, serum calcium and phosphorus, blood sugar levels, urine for reducing sugars, and TORCH titers. If all these investigations are inconclusive of etiology, genetic testing becomes essential. Some mutations may be associated with some peculiar morphologies. The case published in this issue<sup>[8]</sup> highlights how morphology can sometimes be misleading and how crucial genetic testing is in such cases of pediatric cataracts. The authors report two cases of membranous cataracts in siblings with homozygous missense mutation in exon 3 of glucosaminyl (N-acetyl) transferase (*GCNT2*) gene.

In particular, a membranous cataract is a congenital disorder in which the lens is absorbed and flattened with no or little fibers. A membranous cataract is usually diagnosed during late childhood, but can also be seen in early childhood. Patients with a familial nature of cataract associated with Hallermann–Streiff–Francois syndrome, Lowe’s syndrome, and few inheritable mutations reported like *LIM2* (autosomal dominant)<sup>[9]</sup> and *GCNT2* (autosomal recessive) mutation,<sup>[10,11]</sup> as well as patients with intrauterine TORCH infections can have membranous cataracts.

Mutations in the genes encoding lens proteins can directly cause cataracts. Although most of these are inherited as autosomal dominant, five loci have been linked to autosomal recessive inheritance. A recent addition to this literature is the *GCNT2* gene that highlights the association between autosomal recessive congenital cataract and the rare adult i blood group phenotype. Further, three different transcript forms, designated as *GCNT2A*, -B, and -C, have been described of the *GCNT2* gene. Mutations in exon 1 of *GCNT2B* selectively inactivate the transcript expressed in the lens, resulting in congenital cataracts without the i blood phenotype. The case reported here has mutations in exon 3, and hence both reticulocytes and lens epithelial cells are likely to be affected. Altered protein glycosylation is a common feature of cancers, and hence, *GCNT2* could be a marker of systemic cancers.<sup>[12]</sup>

Accurate pediatric cataract diagnosis is crucial for patients and their families as it facilitates genetic counseling. There is a lot of variability in the investigative pathways of pediatric cataract patients. Hereditary cases in our country might be underestimated because of inconsistent testing. Hence, an attempt should be made to investigate the etiology of congenital cataracts thoroughly with the efficient use of supportive genetic tests. Not only does it establish the etiology of cataract, but also it guides us to approach other associated and undiagnosed systemic conditions.

**Savleen Kaur, Jaspreet Sukhija, Kiran Kumari**

Advanced Eye Centre, Post Graduate Institute of Medical Education and Research, Chandigarh, India

Correspondence to: Dr. Savleen Kaur,  
Advanced Eye Centre (Department of Ophthalmology),  
Post Graduate Institute of Medical Education and Research,  
Chandigarh, India.  
E-mail: mailsavleen@gmail.com

## References

1. Yi J, Yun J, Li ZK, Xu CT, Pan BR. Epidemiology and molecular genetics of congenital cataracts. *Int J Ophthalmol* 2011;4:422-32.
2. Haargaard B, Wohlfahrt J, Fledelius HC, Rosenberg T, Melbye M. A nationwide Danish study of 1027 cases of congenital/infantile cataracts: Etiological and clinical classifications. *Ophthalmology* 2004;111:2292-8.
3. Yangzes S, Kaur S, Gupta PC, Sharma M, Jinagal J, Singh J, *et al.* Intraocular lens implantation in children with unilateral congenital cataract in the first 4 years of life. *Eur J Ophthalmol* 2019;29:304-8.
4. Sukhija J, Kaur S, Ram J. Outcome of primary intraocular lens implantation in infants: Complications and rates of additional surgery. *J Cataract Refract Surg* 2016;42:1060-5.
5. Eckstein M, Vijayalakshmi P, Killedar M, Gilbert C, Foster A. Aetiology of childhood cataract in south India. *Br J Ophthalmol* 1996;80:628-32.
6. Jain I, Pillay P, Gangwar D, Kaul V. Congenital cataract: Etiology and morphology. *J Pediatr Ophthalmol Strabismus* 1983;20:238-42.
7. Singh VM, Badakere A, Patil-Chhablani P, Kekunnaya R. Profile of congenital cataract in the first year of life from a tertiary care center in South India-A modern series. *Indian J Ophthalmol* 2021;69:932-6.
8. Ganatra S, Kekunnaya R, Sachdeva V. Bilateral congenital membranous cataracts due to Glucosaminyl (N-Acetyl) Transferase 2 (*GCNT2*) mutation: Life-saving genetic analysis. *Indian J Ophthalmol* 2022;70:2622-3.
9. Pei R, Liang PF, Ye W, Li J, Ma JY, Zhou J. A novel mutation of *LIM2* causes autosomal dominant membranous cataract in a Chinese family. *Int J Ophthalmol* 2020;13:1512-20.
10. Happ H, Weh E, Costakos D, Reis LM, Semina EV. Case report of homozygous deletion involving the first coding exons of *GCNT2* isoforms A and B and part of the upstream region of *TFAP2A* in congenital cataract. *BMC Med Genet* 2016;17:64.
11. Pras E, Raz J, Yahalom V, Frydman M, Garzozzi HJ, Pras E, *et al.* A nonsense mutation in the glucosaminyl (N-acetyl) transferase 2 gene (*GCNT2*): Association with autosomal recessive congenital cataracts. *Invest Ophthalmol Vis Sci* 2004;45:1940-5.
12. Sweeney JG, Liang J, Antonopoulos A, Giovannone N, Kang S, Mondala TS, *et al.* Loss of *GCNT2*/I-branched glycans enhances melanoma growth and survival. *Nat Commun* 2018;9:3368.

This is an open access journal, and articles are distributed under the terms of the Creative Commons Attribution-NonCommercial-ShareAlike 4.0 License, which allows others to remix, tweak, and build upon the work non-commercially, as long as appropriate credit is given and the new creations are licensed under the identical terms.

Access this article online	
Quick Response Code:	Website: www.ijo.in
	DOI: 10.4103/ijo.IJO_124_22

Cite this article as: Kaur S, Sukhija J, Kumari K. Commentary: Genetic testing in cases of pediatric cataract. *Indian J Ophthalmol* 2022;70:2623-4.