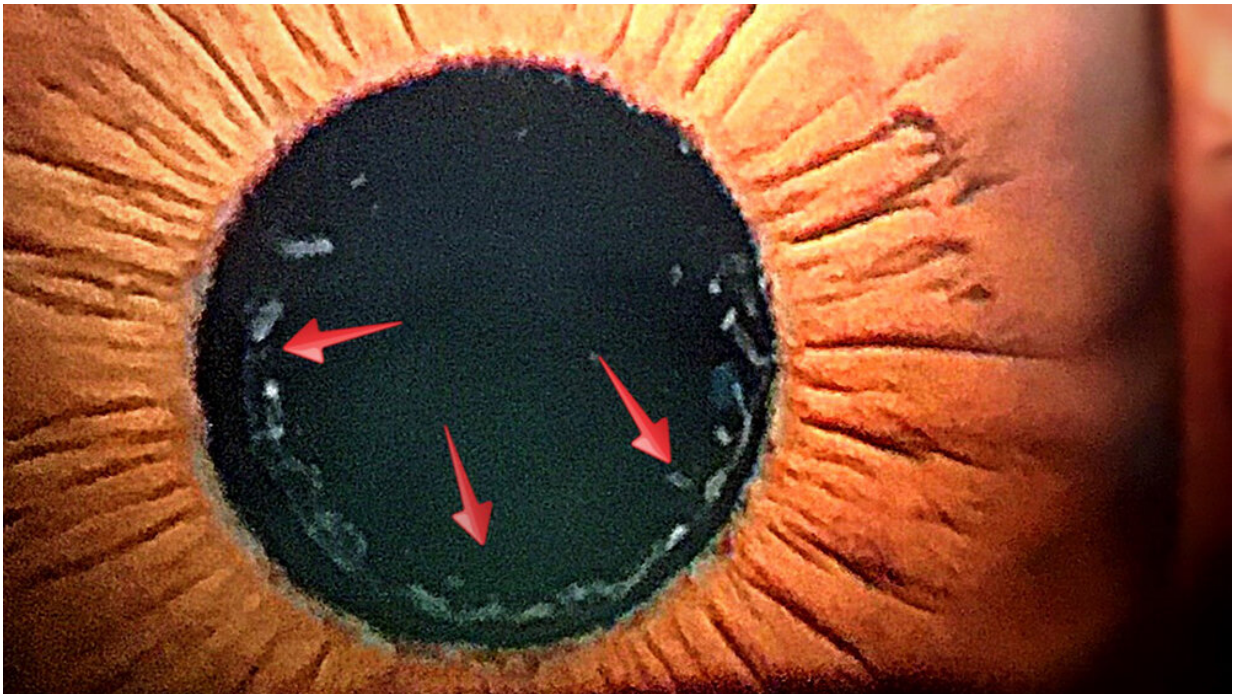


Scientists identify genetic mutation associated with exfoliative syndrome

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Eye of a patient with exfoliation syndrome. The typical white exfoliation material deposits on the surface of the lens are visible (red arrows) with the aid of a slit lamp at 16x magnification under white light. Exfoliation material may not be visible on unaided exam. Credit: Agency for Science, Technology and Research (A*STAR), Singapore

A team of researchers from the Agency for Science, Technology and Research's (A*STAR) Genome Institute of Singapore (GIS) and

Bioprocessing Technology Institute (BTI), as well as Singapore Eye Research Institute (SERI), have identified a genetic mutation (functionally defective CYP39A1 gene) associated with exfoliation syndrome, the most common cause of glaucoma. The findings could pave the way for future research on the cause of exfoliation syndrome and potential cures. Their research was published in *Journal of the American Medical Association (JAMA)* on 24 February 2021.

Exfoliation syndrome is a systemic disorder characterized by abnormal protein material that progressively accumulates in the front of the eye. This disorder is the most common cause of glaucoma, and a major cause of irreversible blindness.

In this study, the scientists sequenced all protein encoding genes of more than 20,000 participants from 14 countries across Asia, Europe, and Africa, including more than 1,200 Singaporeans. They observed that people with [exfoliation](#) syndrome are twice as likely to carry damaging mutations in the gene encoding for the CYP39A1 protein, an enzyme which plays an important role in the processing of cholesterol. Further extended analyses suggest that defective CYP39A1 function is strongly associated with increased risk of exfoliation syndrome.

Although exfoliation syndrome is the most common cause of glaucoma, its origin is shrouded in mystery because it is not known where the abnormal protein deposits (exfoliative material) originate, and how the disease comes about. Answers to these questions could provide approaches to design and develop an effective treatment. The current findings point to the important role of cholesterol processing in the exfoliation [syndrome](#) disease process. As cholesterol is found abundantly in all cells, disruption to how cholesterol is processed due to defective CYP39A1 activity could adversely impact their normal functions. In particular, this study discovered that epithelial cells in the front of the eye responsible for filtering the [blood supply](#) to produce the clear fluid

known as aqueous humor that bathes and nourishes other cells in the eye, were most affected by the CYP39A1 gene mutation. Disruption to the gene function can compromise the filtering function of [epithelial cells](#) and lead to leakage of exfoliative material from the blood into the eye.

Prof Patrick Tan, Executive Director of GIS, said, "This is a ground-breaking study that could facilitate future research efforts aimed at restoring defective CYP39A1 function and inhibiting the formation of exfoliation material in the eye as treatments for [exfoliation syndrome](#) and glaucoma." Prof Aung Tin, Director of SERI and Deputy Medical Director of SNEC, said, "This is a major eye disease, affecting over 70 million people worldwide, which causes a lot of visual morbidity and blindness, not only from glaucoma but also due to complications related to cataract surgery. This study was notable for involving many centers from many different countries around the world, but led from Singapore. The study findings are very exciting as we found a new pathway for the disease which opens up possibilities for new treatments."

Prof David Friedman, the Albert and Diane Kaneb Chair in Ophthalmology at Harvard University and Director of the Glaucoma Service at the Massachusetts Eye and Ear Infirmary, Boston, commented, "Very exciting work. The researchers have identified rare gene variants that results in disrupted cholesterol homeostasis and transport that will open the door to novel therapeutics. Having studied over 20,000 individuals, the study demonstrates the power of studying rare variants to detect disease-causing [genes](#) in complex conditions." Prof Friedman was not involved in the study.

More information: undefined undefined et al. Association of Rare CYP39A1 Variants With Exfoliation Syndrome Involving the Anterior Chamber of the Eye, *JAMA* (2021). [DOI: 10.1001/jama.2021.0507](https://doi.org/10.1001/jama.2021.0507)

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