

# Genetic mutations warn of skin cancer risk

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Researchers have discovered that mutations in a specific gene are responsible for a hereditary form of melanoma.

Every year in the UK, almost 12,000 people are diagnosed with melanoma, a form of skin cancer. About 1 in 20 people with melanoma have a strong family history of the disease. In these patients, pinpointing the genetic [mutations](#) that drive disease development allows dermatologists to identify people who should be part of melanoma surveillance programmes.

The team found that people with specific mutations in the POT1 gene were extremely likely to develop melanoma. These mutations deactivate the POT1 gene that protects the ends of our chromosomes from damage.

"Genomics is on the verge of transforming the healthcare system – this study highlights the potential clinical benefits that can be gained through genomic studies and offers potential strategies to improve patient care and disease management," says Dr David Adams, co-senior author from the Wellcome Trust Sanger Institute. "With this discovery we should be able to determine who in a family is at risk, and in turn, who should be regularly screened for early detection."

Known [genetic mutations](#) account for approximately 40 per cent of all occurrences of inherited forms of melanoma. The team set out to identify the hereditary mutations that account for the other ~60 per cent by sequencing part of the genome of 184 patients with hereditary

melanoma caused by unknown mutations.

They found that the inactivation of POT1 caused by these mutations leads to longer and potentially unprotected telomeres, regions at the end of our chromosomes that protect chromosomes from damage.

"This finding significantly increases our understanding of why some families have a high incidence of melanoma," says Professor Tim Bishop, Director of the Leeds Institute of Cancer and Pathology. "Since this gene has previously been identified as a target for the development of new drugs, in the future, it may be possible that [early detection](#) will facilitate better management of this disease."

The team also found that there were also cases of other cancer types in families with these hereditary mutations such as leukaemias and brain tumours. It seems that mutations that deactivate the POT1 gene may underlie other cancers, not just melanoma.

"Our research is making a real difference to understanding what causes melanoma and ultimately therefore how to prevent and treat melanoma and is a prime example of how genomics can transform public health," says Professor Julia Newton Bishop, co-senior author from the University of Leeds. "This study would not have been possible without the help and patience from the families that suffer from these devastating, inherited forms of melanoma."

The team are currently working on developing cells and mice with an inactive POT1 gene. These will be used to test potential drug therapies that alter telomere metabolism.

Dr Safia Danovi, Cancer Research UK's senior science communications officer, said: "This is a step forward for people with a strong family history of [melanoma](#), the most dangerous form of [skin cancer](#). But it's

important to remember that, for most of us, avoiding sunburn and sunbeds is the best way to reduce the risk of this disease."

**More information:** Carla Daniela Robles-Espinoza et al (2014). 'POT1 loss-of-function variants predispose to familial melanoma' Advanced online publication in *Nature Genetics* on 30 March 2014. [DOI: 10.1038/ng.2947](https://doi.org/10.1038/ng.2947)

Provided by Wellcome Trust Sanger Institute

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