

Why some moles become melanoma still a mystery

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Half of all melanomas develop from a naevus. Credit: University of Queensland

Testing for two gene mutations commonly associated with melanoma would be insufficient to determine whether a mole could turn cancerous, University of Queensland research has found.

UQ Diamantina Institute's Dr Mitchell Stark is among researchers investigating why melanomas develop from some naevi (moles).

"In Australia, about half of all melanomas develop from a naevus, but most moles will never progress to become a melanoma," Dr Stark said.



"We are trying to determine what causes some naevi to change so that we can better predict or more accurately detect those which could become dangerous.

"This would help avoid unnecessary excisions of those lesions unlikely to pose a risk."

Scientists from UQ's Dermatology Research Centre analysed samples from participants in the Brisbane Naevus Morphology Study, and discovered all had one of two key <u>mutations</u> associated with melanoma.

"We found that 85 per cent of samples had a mutation on the gene known as BRAF, and the remaining samples had a mutation on the NRAS gene," Dr Stark said.

"When either of these <u>genes</u> are mutated it activates the signalling pathway known as MAPK, which is commonly active in melanomas.

"Clearly our samples were not melanomas, so additional genomic events need to occur before a <u>mole</u> becomes malignant."

Dr Stark said further research was underway to determine other genetic changes that could trigger the development of melanoma from naevi.

Studies have consistently shown the number of naevi a person has is the strongest predictor of risk for <u>melanoma</u>.

Dr Stark said people with a high number of moles, and other risk characteristics such as fair skin or light coloured hair or eyes, should continue to see their treating dermatologist or skin cancer physician for routine skin examination.

The research was published in the British Journal of Dermatology.



More information: J. M. Tan et al. The BRAF and NRAS mutation prevalence in dermoscopic subtypes of acquired naevi reveals constitutive MAPK pathway activation, *British Journal of Dermatology* (2017). DOI: 10.1111/bjd.15809

Provided by University of Queensland

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