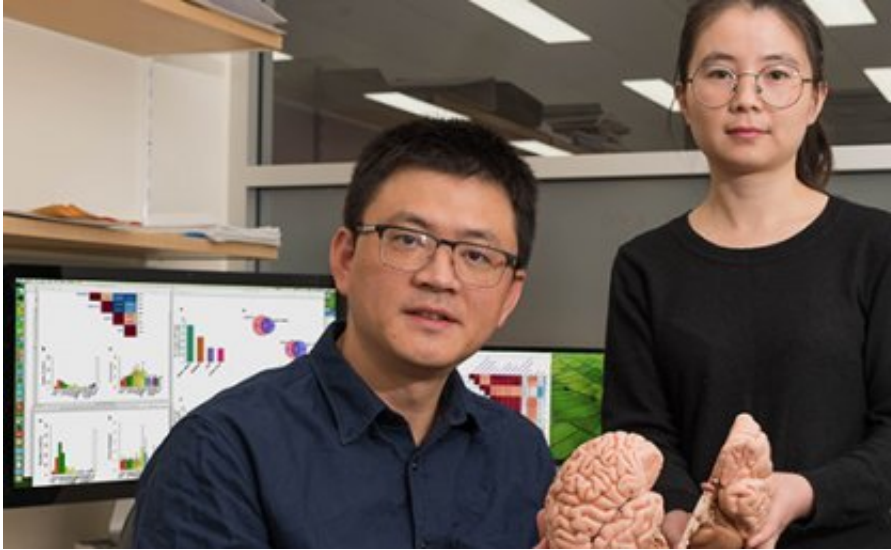


Brain secrets that flow in our blood

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Professor Jian Yang and Dr Ting Qi. Credit: University of Queensland

Our blood can be used to uncover genetic secrets inside the brain, according to University of Queensland research.

A team led by Professor Jian Yang from UQ's Institute for Molecular Bioscience and Queensland Brain Institute, and Dr Ting Qi from the Institute for Molecular Biosciences, discovered blood samples can be used to identify gene targets for brain-related traits like IQ and diseases such as schizophrenia.

Professor Yang said this discovery would open up new avenues of research and improve scientists' ability to unravel the complexity of

brain disease.

"Identifying gene targets for brain diseases is an essential part of understanding [disease risk](#) and developing treatment and prevention strategies," he said.

"But the effects of individual genes on brain traits are often very small, meaning we require large sample numbers to find the gene targets, and only a few hundred brain samples are available in most databases.

"We wondered whether we could overcome this issue by using blood samples, which number in tens of thousands, to identify gene targets for brain-related complex traits and diseases."

Professor Yang, Dr Qi and the team found the local genetic controls of [gene expression](#) in the brain are very similar to those in blood.

"This discovery provides strong justification for using genetic effects on gene expression estimated from [blood samples](#) as a proxy for those in the brain to boost the power of our studies to identify the genes underlying brain traits and disorders," Dr Qi said.

"Using [blood](#) data enabled us to find 97 genes associated with IQ, schizophrenia, years of education and cigarette smoking, compared to 61 using [brain](#) data alone."

Professor Yang last year won the Frank Fenner Prize for Life Scientist of the Year at the Prime Minister's Prizes for Science, and this year has made several significant discoveries, including using genetic data to identify modifiable disease risk factors, pinpointing the [genes](#) and epigenetic factors that are linked to specific diseases, natural selection and evolution.

The study is published in international scientific journal *Nature Communications*.

More information: Identifying gene targets for brain-related traits using transcriptomic and methylomic data from blood, *Nature Communications* (2018). [DOI: 10.1038/s41467-018-04558-1](https://doi.org/10.1038/s41467-018-04558-1)

Provided by University of Queensland

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