

Largest study of its kind reveals that many psychiatric disorders arise from common genes

December 12 2019



Credit: CC0 Public Domain

Many distinct psychiatric diseases share a common genetic structure, according to new research by scientists at Massachusetts General Hospital (MGH) and the Psychiatric Genomics Consortium, an international team of investigators. Psychiatric disorders affect more

than 25 percent of the population in a given year. In the largest-ever study of its kind, published in the journal *Cell*, researchers identified more than 100 genetic variants that affect the risk for more than one mental health condition.

A gene is made up of segments of DNA; an alteration in the DNA sequence produces a [gene variant](#), which can increase or decrease the risk for disease. Many individual gene variants that affect the risk for specific psychiatric [disorders](#) have been identified. However, [genes](#) are often pleiotropic, meaning they produce multiple effects in the body.

Identifying gene variants that influence the risk for more than one psychiatric disorder is an important step toward improving the diagnosis and treatment of these conditions, says the study's senior author, Jordan W. Smoller, MD, ScD, director of MGH's Psychiatric and Neurodevelopmental Genetics Unit and a professor of Psychiatry at Harvard Medical School (HMS). "Understanding how specific genetic variations may contribute to a broad spectrum of illnesses can tell us something about the degree to which these disorders may have a shared biology," says Smoller.

To identify these multi-purpose gene variants, the researchers used a technique called genome-wide association to analyze genetic data from 494,162 healthy control subjects and 232,964 people diagnosed with at least one of eight common psychiatric disorders. The analysis identified 109 gene variants that affect the risk for more than one psychiatric disorder.

Certain disorders shared many variants, allowing the researchers to divide the conditions into three groups of genetically-related conditions: disorders characterized by compulsive behaviors (anorexia nervosa, [obsessive-compulsive disorder](#) and, to a lesser extent, Tourette syndrome); mood and psychotic disorders (bipolar disorder, major

depression and schizophrenia); and early-onset neurodevelopmental disorders (autism spectrum disorder, ADHD and Tourette syndrome). The researchers also found evidence that genes associated with multiple disorders show increased expression beginning in the second trimester of pregnancy and appear to play an important role in [brain development](#).

Knowing which gene variants increase the odds for developing multiple psychiatric disorders provides new clues about the biological pathways that contribute to mental illness, says computational geneticist Phil H. Lee, Ph.D., of the Center for Genomic Medicine at MGH and HMS, lead author of the study. "And learning how disorders are related at a biological level may inform how we classify and diagnose mental health conditions," says Lee.

Moreover, the study identified several gene variants that had especially widespread influence on the risk for a number of psychiatric disorders, which could prove to be an important finding for the prevention and care of [psychiatric disorders](#). "To the extent that these genes may have broad effects," says Smoller, "they could be potential targets for developing new treatments that might benefit multiple conditions."

More information: *Cell* (2019). [DOI: 10.1016/j.cell.2019.11.020](https://doi.org/10.1016/j.cell.2019.11.020)

Provided by Massachusetts General Hospital

Citation: Largest study of its kind reveals that many psychiatric disorders arise from common genes (2019, December 12) retrieved 21 May 2023 from <https://medicalxpress.com/news/2019-12-largest-kind-reveals-psychiatric-disorders.html>

This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.