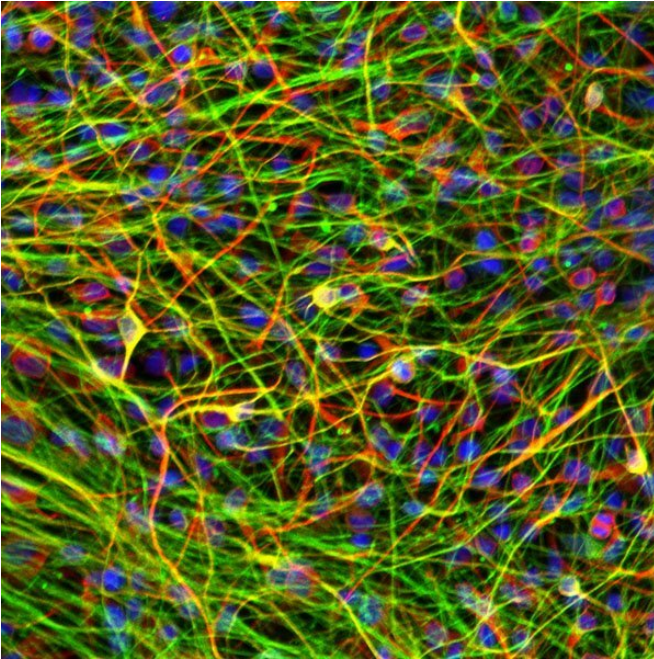


Gene pathway linked to schizophrenia identified through stem cell engineering

21 December 2020, by Deborah Mann Lake



These are neurons derived from induced pluripotent stem cells from a schizophrenia patient. Credit: Photo by Laura Stertz, PhD/UTHealth

Using human-induced pluripotent stem cells engineered from a single family's blood samples, a gene signaling pathway linked to a higher risk for developing schizophrenia was discovered by scientists at The University of Texas Health Science Center at Houston (UTHealth). The research was published in a recent issue of *Neuropsychopharmacology*.

The signaling [pathway](#) researchers pinpointed is called phosphoinositide 3-kinase/glycogen synthase kinase 3 (PI3K/GSK3). Among the differentially expressed genes along the pathway was one called serum-glucocorticoid kinase 1 (SGK1), an inhibitor of GSK3 beta, which has been associated with [schizophrenia](#).

"We believe this has direct implications for the treatment of patients," said senior author Consuelo Walss-Bass, Ph.D., professor in the Louis A. Faillace, MD, Department of Psychiatry and Behavioral Sciences at McGovern Medical School at UTHealth. "There is a new antipsychotic that just received approval from the Food and Drug Administration that directly targets the pathway we identified as dysregulated in neurons from the patients, and several other antipsychotics also target this pathway. This could help pinpoint who may respond better to treatments."

Researchers led by Walss-Bass and first author, postdoctoral research fellow Laura Stertz, Ph.D., used [blood samples](#) from adult members of a large family with multiple individuals affected by schizophrenia. Through human-induced pluripotent stem cell (hiPSC) technology, the [blood cells](#) were reprogrammed into stem cells, which were then directed to become brain neurons. Those neurons could be studied in a virtual biopsy and compared to neurons engineered from individuals who did not have schizophrenia, but came from the same family from a homogenous population in the Central Valley of Costa Rica.

"Mental health research has lagged behind because we don't know what is happening biologically. We are diagnosing people based on what they are telling us," Walss-Bass said. "Even postmortem, the [brain tissue](#) in mental health disorders looks perfectly fine. In Alzheimer's disease, you can see a difference compared to controls. But not in psychiatric disorders. Now by studying virtual brain biopsies, we can tell what is happening biologically."

Among the differentially expressed genes the researchers saw in the virtual biopsies were five that have previously been identified as schizophrenia candidate genes by [genome-wide association studies](#).

Among the genes associated with the PI3K/GSK3 pathway was SGK1, which inhibits GSK3 activity. Those alterations are linked to whether a person has a higher risk of developing schizophrenia.

"We were able to find significant, meaningful differences with a small control group," Walss-Bass said. "Neurons of patients with schizophrenia had alterations in the signaling pathway. This research may help to understand how or why some antipsychotics targeting GSK3 work and also to develop other target-specific medications."

Walss-Bass said identifying patients with specific biological pathway markers could identify them as the best candidates for medications, creating the personal pharmacology that is needed to treat psychiatric disorders.

More information: Laura Stertz et al. Convergent genomic and pharmacological evidence of PI3K/GSK3 signaling alterations in neurons from schizophrenia patients, *Neuropsychopharmacology* (2020). DOI: [10.1038/s41386-020-00924-0](https://doi.org/10.1038/s41386-020-00924-0)

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