

Genetic analysis can improve depression therapy

January 12 2018



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The failure of drugs such as SSRIs, used to treat depression, can be a result of genetic variations in patients. Variations within the gene that encodes the CYP2C19 enzyme results in extreme differences in the

levels of escitalopram achieved in patients, according to a new study published in *The American Journal of Psychiatry*. Prescribing the dose of escitalopram based on a patient's specific genetic constitution would greatly improve therapeutic outcomes. The study was conducted at Karolinska Institutet in Sweden in association with researchers at Diakonhjemmet Hospital in Oslo, Norway.

Pharmaceutical treatment of depression commonly makes use of selective serotonin reuptake inhibitors (SSRIs) of which escitalopram is the most frequently administered clinically. However, escitalopram therapy is currently limited by the fact that some [patients](#) do not respond well to the [drug](#), while others develop adverse reactions requiring discontinuation of treatment.

In order to individualise drug therapy, researchers are attempting to establish genetic biomarkers that can predict an individual's response to drugs. In a recent study, it was discovered that variation in the gene encoding the enzyme responsible for escitalopram metabolism (CYP2C19) is very important in this respect. Individuals with a variant of the gene promoting increased enzyme expression had blood levels of escitalopram too low to impact the depression symptoms, whereas patients with a defective CYP2C19 gene reached drug levels which were too high. Overall, one third of the 2,087 study participants achieved escitalopram blood levels that were either too high or too low.

Interestingly, the researchers found that 30 per cent of the patients carrying gene variants causing excessive or inadequate enzyme levels switched to other drugs within one year, in contrast with only 10 to 12 per cent of patients carrying the common gene.

"Our study shows that genotyping of CYP2C19 could be of considerable clinical value in individualising doses of [escitalopram](#) so that a better all-round antidepressive effect could be achieved for the patients," says

Professor Magnus Ingelman-Sundberg at Karolinska Institutet's Department of Physiology and Pharmacology who led the study together with Professor Espen Molden. "Because CYP2C19 is involved in the metabolism of many different SSRIs, the finding is also applicable to other types of antidepressants."

More information: Marin M. Jukić et al. Impact of CYP2C19 Genotype on Escitalopram Exposure and Therapeutic Failure: A Retrospective Study Based on 2,087 Patients, *American Journal of Psychiatry* (2018). [DOI: 10.1176/appi.ajp.2017.17050550](https://doi.org/10.1176/appi.ajp.2017.17050550)

Provided by Karolinska Institutet

Citation: Genetic analysis can improve depression therapy (2018, January 12) retrieved 25 April 2024 from <https://medicalxpress.com/news/2018-01-genetic-analysis-depression-therapy.html>

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