OXYTOCIN PATHWAY POLYMORPHISMS AND SOCIAL FUNCTIONING IN SCHIZOPHRENIA

Vera Mikhailova

I.M. Sechenov First Moscow State Medical University, Moscow, Russian Federation

Correspondence to: voviand.vm@gmail.com

Cite as: Mikhailova, V. (2020) OXYTOCIN PATHWAY POLYMORPHISMS AND SOCIAL FUNCTIONING IN SCHIZOPHRENIA. The Physician vol 6; issue 3; epub 21 Aug 2020; DOI: https://doi.org/10.38192/1.6.3.firmst20.ab08

Article Information
Published: 21 Aug 2020
Open Access – Creative Commons Licence CC-BY-ND-4.0

Introduction

The social adaptation for any schizophrenia patient and their family is the main factor allowing to judge whether therapy is working or not. That makes the improvement of patients’ social functioning the main goal for an attending physician, the pharmaceutical industry specialists and also for biomedical researchers.

To obtain a better knowledge of the pathogenesis of poor social performance in schizophrenia patients under treatment, it seems necessary to take into consideration the disturbance of the oxytocin pathway.

Neurohormone oxytocin is known to play an important role in social behaviour. The latest neuroimaging studies have proven OXTR (oxytocin receptor) expressing cells to be largely involved in social cognition tasks. Moreover, such polymorphisms as CD38 (rs3796863), OXTR (rs53576), OXTR (rs4686302) have been observed to be associated with the pathogenesis of several diseases such as autism, OCD and bipolar disorder.

Methods

Social functioning was assessed with the Personal and the Social Performance (PSP) scale in 732 patients (women 343), aged 36.5 ± 12.3 years, with an ICD-10 diagnosis of schizophrenia or schizoaffective psychosis. Firstly, genotyping was performed with HRM (High Resolution Melt) method using DNA samples extracted from patients’ venous blood samples by organic solvents and amplified by PCR.

Secondly, all patients were stratified by the level of social functioning into two groups: “moderate and good” and “marked and poor”. Allele frequencies were aligned with the Hardy-Weinberg equation (p>0,05). Genotypes were calculated in each group of social functioning.

Results

CC allele CD38 (rs3796863) tended to be more frequent in “poor” performance group within a socially useful activity PSP scale. OXTR rs53576 GG allele was considerably more frequent in “poor” performance
The Physician

OXYTOCIN PATHWAY POLYMORPHISMS AND SOCIAL FUNCTIONING IN SCHIZOPHRENIA

Conclusion

It is, therefore, the first-time oxytocin pathway polymorphisms are found to be associated with social functioning in schizophrenia patients, with CD38 rs3796863 CC and OXTR rs53576 GG being risk allele.

Conflict of Interest

No conflict of interest declared