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ASSOCIATION STUDY BETWEEN NTRK2 GENE POLYMORPHISMS AND SCHIZOPHRENIA IN CHINESE HAN PEOPLE

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Objective: To explore the relationships between Neurotrophic Tyrosine Kinase Receptor Type 2 (NTRK2) gene polymorphisms and schizophrenia in Chinese Han population.

Methods: The NTRK2SNPs rs1187272 were genotyped by applying polymerase chain reaction-restriction fragment length polymorphism (PCR-RLFP) technique in 100 patients with schizophrenia and 119 unrelated healthy controls, then the comparison of the alleles and genotypes frequencies of the NTRK2 gene between two groups was performed.

Results: There were no significant difference in alleles (χ^2 =1.78, df=1, P>0.05) and genotypes (χ^2 =1.01, df=2, P>0.05) distribution of NTRK2 gene between patients and controls. There were also no significant difference in alleles (χ^2 =2.70, df=3, P>0.05) and genotypes (χ^2 =4.72, df=6, P>0.05) distribution among male and female patients and controls. There were significant difference in alleles (χ^2 =5.53, df=1, P=0.02) distribution between different first-onset age patients groups, There were also significant difference in alleles (χ^2 =4.38, df=1, P=0.04) distribution between schizophrenia with earlier first-onset age (≤25 years old) and control subjects.

Conclusion: The NTRK2 gene was associated with schizophrenia with earlier first-onset age, and it may be one of common susceptible genes for schizophrenia in Chinese Han population.