

## Effect of 5-HTTLPR on connectivity and topological properties of resting state EEG networks

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Serotonin transporter is one of the most widely investigated genetic markers of individual variation in serotonergic function. The promoter region of the serotonin transporter gene (5-HTTLPR) contains long (L) and short (S) variants with the latter one having reduced transcriptional efficiency. S allele has been found to increase the risk of depression and other mental health problems, but some evidence suggests that S-allele carriers outperform subjects carrying the long allele in an array of cognitive tasks. Functional magnetic resonance imaging studies demonstrate a heightened amygdala response to negative emotional stimuli and diminished connectivity among key areas involved in emotion regulation in S allele carriers. However, evidence linking this polymorphism with individual variation in electrophysiological properties of resting state brain networks is still very limited. This study investigated the effect of 5-HTTLPR polymorphism on EEG current source density, connectivity, and topological properties of resting state networks. As compared to L homozygotes, S-allele carriers showed lower current source density and connectivity in most frequency bands in areas overlapping with the default mode and emotion regulation regions. The analysis of graph-theoretical measures showed that as compared to L homozygotes, S-allele carriers have less optimal topological properties of brain networks in theta, but more optimal in alpha band. This dissociation may reflect predisposition to emotional disorders, which is inherent to S allele carriers, and, on the other hand, their superior functioning in some cognitive domains.

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