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healthy relatives (n=137). A case-control association analysis was performed in 993 patients and 977 controls. We identified a novel missense mutation, which we genotyped in the PEX6-1 region in a large replication sample of 1298 patients and 1257 controls. The deletion-1 allele was over-represented among the schizophrenia patients compared to the controls ( $P=0.002$ ), with an allele-specific risk ratio of 2.3 (95% CI: 1.3-4.1). In the case-control association analysis, the deletion-1 allele was significantly associated with schizophrenia ( $P=0.0023$ ). The association remained significant after correction for multiple testing. We found no evidence for an association 82157476af

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