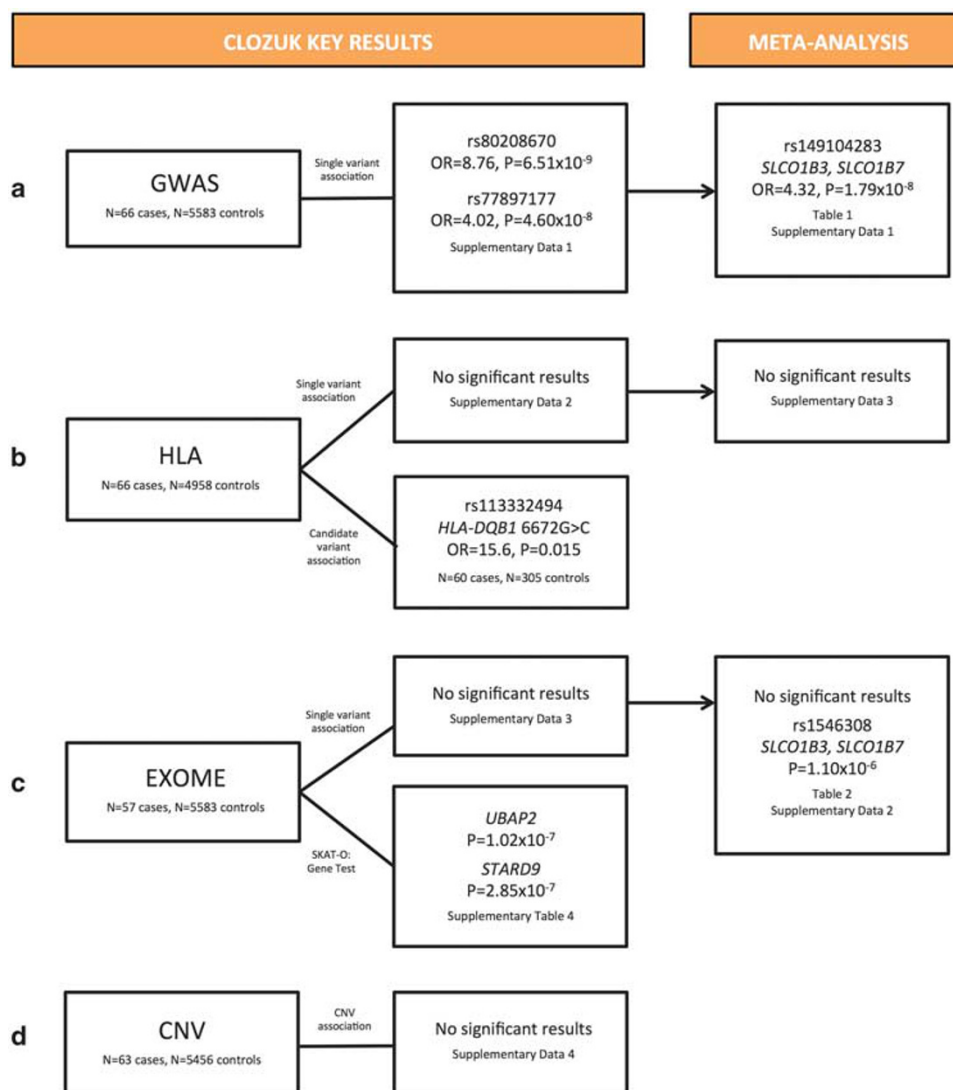


CORRIGENDUM

Genome-wide common and rare variant analysis provides novel insights into clozapine-associated neutropenia

SE Legge, ML Hamshere, S Ripke, AF Pardini, JI Goldstein, E Rees, AL Richards, G Leonenko, LF Jorskog, Clozapine-Induced Agranulocytosis Consortium, KD Chambert, DA Collier, G Genovese, I Giegling, P Holmans, A Jonasdottir, G Kirov, SA McCarroll, JH MacCabe, K Mantripragada, JL Moran, BM Neale, H Stefansson, D Rujescu, MJ Daly, PF Sullivan, MJ Owen, MC O'Donovan and JTR Walters

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Correction to: *Molecular Psychiatry* (2017) 22: 1502-1508; advance online publication, 12 July 2017; doi: 10.1038/mp.2016.97

In the first paragraph of the Results section and Figure 1, the authors incorrectly referred to the finding of SNP rs77897117. The correct SNP is rs77897177.

The corrected figure appears in previous page.



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