



# Corrigendum

# Identification of specific genes involved in schizophrenia aetiology – what difference does it make? – CORRIGENDUM

**David Curtis** 

## Keywords

Schizophrenia; genes; exome; NMDA receptor; SETD1A.

## Copyright and usage

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The author would like to update the citation of a pre-print in the references to the full citation. The full citation of reference 1 should read:

Singh T, on behalf of the Schizophrenia Exome Meta-analysis (SCHEMA) Consortium. Rare coding variants in ten genes confer substantial risk for schizophrenia. *Nature* 2022; 604: 509–516. https://doi.org/10.1038/s41586-022-04556-w

The author and Publisher apologise for this error.

### References

Curtis D. Identification of specific genes involved in schizophrenia aetiology – what difference does it make? *The British Journal of Psychiatry* 2022; 1–3. doi: 10.1192/bip.2021.153

Singh T, on behalf of the Schizophrenia Exome Meta-analysis (SCHEMA) Consortium. Rare coding variants in ten genes confer substantial risk for schizophrenia. *Nature* 2022; **604**: 509–516. https://doi.org/10.1038/s41586-022-04556-w