

Identification of an Agrin Mutation that Causes Congenital Myasthenia and Affects Synapse Function

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An error during the printing process resulted in an incomplete version of Figure 1B in this paper. The correct panel (and legend) appears below. The online version was not affected. *AJHG* sincerely regrets the error.

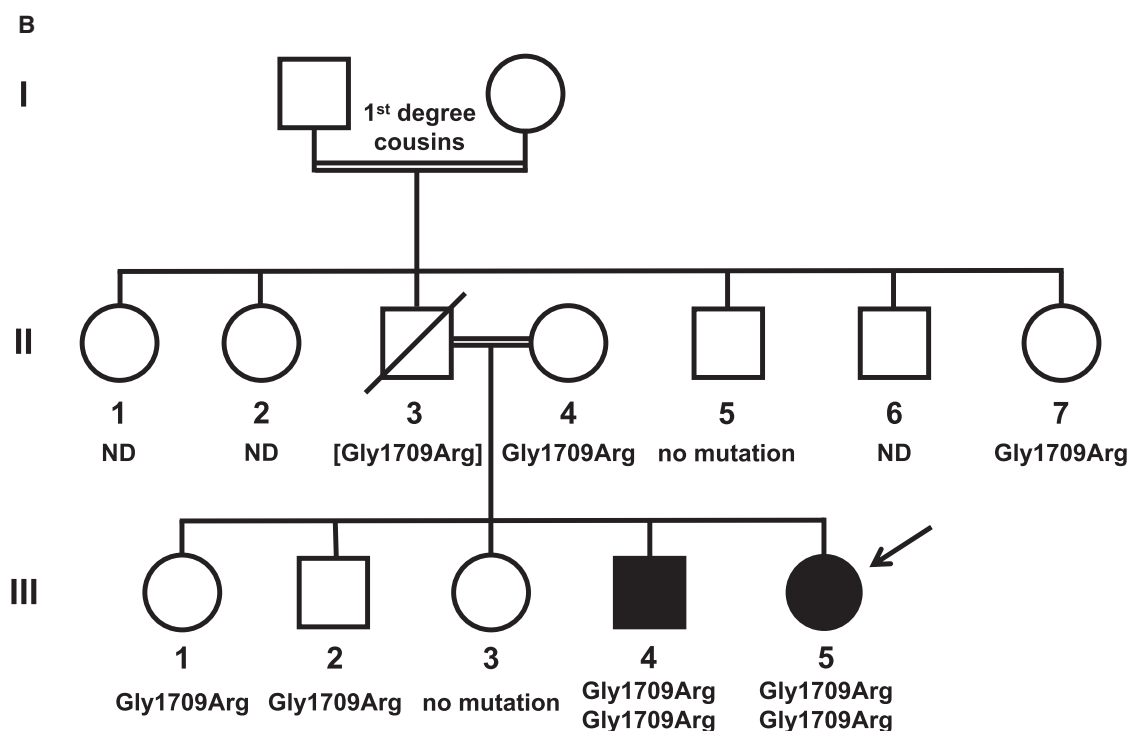


Figure 1. Identification of the Mutation c.5125G>C in *AGRN* Exon 29 and Hereditary Transmission: Corrected Panel B
(B) Family pedigree. The proband is indicated by an arrow. The expected nucleotide change p.Gly1709Arg transmitted in this consanguineous family is indicated below the symbols. ND: not determined.

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