

CORRIGENDUM: Autozygosity reveals recessive mutations and novel mechanisms in dominant genes: implications in variant interpretation

Dorota Monies PhD, Sateesh Maddirevula PhD, Wesam Kurdi MD, Mohammed H. Alanazy MD, Hisham Alkhalidi MD, Mohammed Al-Owain MD, Raashda A. Sulaiman MD, Eissa Faqeih MD, Ewa Goljan, MPharm, Niema Ibrahim BSc, Firdous Abdulwahab BSc, Mais Hashem BSc, Mohamed Abouelhoda PhD, Ranad Shaheen PhD, Stefan T. Arold PhD and Fowzan S. Alkuraya MD

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In the published version of this article, the legends for the Supplementary Figures were missing. The legends appear below. The authors regret the error.

Supplementary Figure S1. Pedigrees of families reported in this study. Note the consanguineous nature of all pedigrees. The genotypes for the identified recessive variants are shown in red under each symbol.

Supplementary Figure S2. Homology model of the GABRD molecule, established based on the 50% identical PDB entry 4cof (QMEAN score –2.03). The five chains that form the pentameric channel are color-coded. Thr292, located in the helical transmembrane channel domain, is shown in green. Left: side view. Right: view into the channel, orientation toward the cytoplasm.