

Most genetic risk for autism resides with common varia

Nature Genetics

46, 881-885

DOI: [10.1038/ng.3039](https://doi.org/10.1038/ng.3039)

Citation Report

#	ARTICLE	IF	CITATIONS
2	Somatic Mosaicism in the Human Genome. <i>Genes</i> , 2014, 5, 1064-1094.	1.0	122
3	Navigating the new road in psychiatry. <i>Science-Business EXchange</i> , 2014, 7, 913-913.	0.0	0
4	The Perfect Storm: Preterm Birth, Neurodevelopmental Mechanisms, and Autism Causation. <i>Perspectives in Biology and Medicine</i> , 2014, 57, 470-481.	0.3	11
5	Editorial: Translational genetics of child psychopathology: a distant dream?. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2014, 55, 1065-1067.	3.1	4
6	Common microRNAs Target Established ASD Genes. <i>Frontiers in Neurology</i> , 2014, 5, 205.	1.1	15
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8	Synaptic, transcriptional and chromatin genes disrupted in autism. <i>Nature</i> , 2014, 515, 209-215.	13.7	2,254
9	Common genetic variants linked with large percentage of autism risk. <i>American Journal of Medical Genetics, Part A</i> , 2014, 164, vii-viii.	0.7	0
10	Opposite risk patterns for autism and schizophrenia are associated with normal variation in birth size: phenotypic support for hypothesized diametric gene-dosage effects. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2014, 281, 20140604.	1.2	31
11	Etiology of Autism Spectrum Disorder: A Genomics Perspective. <i>Current Psychiatry Reports</i> , 2014, 16, 501.	2.1	12
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18	Network assisted analysis to reveal the genetic basis of autism. <i>Annals of Applied Statistics</i> , 2015, 9, 1571-1600.	0.5	43
19	Association of genetic variants of GRIN2B with autism. <i>Scientific Reports</i> , 2015, 5, 8296.	1.6	39

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20	Kernel Approach for Modeling Interaction Effects in Genetic Association Studies of Complex Quantitative Traits. <i>Genetic Epidemiology</i> , 2015, 39, 366-375.	0.6	12
21	Rising interdisciplinary collaborations refine our understanding of autisms and give hope to more personalized solutions. <i>Personalized Medicine</i> , 2015, 12, 359-369.	0.8	1
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90	A Quantitative Electrophysiological Biomarker of Duplication 15q11.2-q13.1 Syndrome. <i>PLoS ONE</i> , 2016, 11, e0167179.	1.1	54
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