Quantifying the Impact of Rare and Ultra-rare Coding V
Spectrum

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Citation Report

#	Article	IF	Citations
1	Functional architecture of low-frequency variants highlights strength of negative selection across coding and non-coding annotations. Nature Genetics, 2018, 50, 1600-1607.	9.4	132
3	Rare-variant collapsing analyses for complex traits: guidelines and applications. Nature Reviews Genetics, 2019, 20, 747-759.	7.7	147
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6	The Relevance of Variants With Unknown Significance for Autism Spectrum Disorder Considering the Genotypeâ€“Phenotype Interrelationship. Frontiers in Psychiatry, 2019, 10, 409.	1.3	8
7	Spatial genome exploration in the context of cognitive and neurological disease. Current Opinion in Neurobiology, 2019, 59, 112-119.	2.0	12
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17	Targeted exon sequencing in deceased schizophrenia patients in Denmark. International Journal of Legal Medicine, 2020, 134, 135-147.	1.2	2
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19	Functional characterization of rare NRXN1 variants identified in autism spectrum disorders and schizophrenia. Journal of Neurodevelopmental Disorders, 2020, 12, 25.	1.5	17

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20	Insufficient Evidence for â€œAutism-Specificâ€•Genes. American Journal of Human Genetics, 2020, 106, 587-595.	2.6	110
21	The mutational constraint spectrum quantified from variation in 141,456 humans. Nature, 2020, 581, 434-443.	13.7	6,140
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23	Genetic Variation across Phenotypic Severity of Autism. Trends in Genetics, 2020, 36, 228-231.	2.9	21
24	Genetics of schizophrenia in the South African Xhosa. Science, 2020, 367, 569-573.	6.0	93
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