Elizabeth Atkinson

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	27.8	929
2	International meta-analysis of PTSD genome-wide association studies identifies sex- and ancestry-specific genetic risk loci. Nature Communications, 2019, 10, 4558.	12.8	363
3	An Unexpectedly Complex Architecture for Skin Pigmentation in Africans. Cell, 2017, 171, 1340-1353.e14.	28.9	134
4	Tractor uses local ancestry to enable the inclusion of admixed individuals in GWAS and to boost power. Nature Genetics, 2021, 53, 195-204.	21.4	125
5	No Evidence for Recent Selection at FOXP2 among Diverse Human Populations. Cell, 2018, 174, 1424-1435.e15.	28.9	99
6	Localization of adaptive variants in human genomes using averaged one-dependence estimation. Nature Communications, 2018, 9, 703.	12.8	83
7	Population and Evolutionary Genomics of <i>Amblyomma americanum</i> , an Expanding Arthropod Disease Vector. Genome Biology and Evolution, 2016, 8, 1351-1360.	2.5	70
8	The impact of non-additive genetic associations on age-related complex diseases. Nature Communications, 2021, 12, 2436.	12.8	55
9	Low-coverage sequencing cost-effectively detects known and novel variation in underrepresented populations. American Journal of Human Genetics, 2021, 108, 656-668.	6.2	49
10	Cortical Folding of the Primate Brain: An Interdisciplinary Examination of the Genetic Architecture, Modularity, and Evolvability of a Significant Neurological Trait in Pedigreed Baboons (Genus <i>Papio</i>). Genetics, 2015, 200, 651-665.	2.9	48
11	High rate of extrapair paternity in a human population demonstrates diversity in human reproductive strategies. Science Advances, 2020, 6, eaay6195.	10.3	41
12	Molecular genetic overlap between posttraumatic stress disorder and sleep phenotypes. Sleep, 2020, 43, .	1.1	32
13	Characterizing the effect of background selection on the polygenicity of brain-related traits. Genomics, 2021, 113, 111-119.	2.9	24
14	Evolutionary and developmental implications of asymmetric brain folding in a large primate pedigree. Evolution; International Journal of Organic Evolution, 2016, 70, 707-715.	2.3	9
15	Inferring archaic introgression from hominin genetic data. Evolutionary Anthropology, 2021, 30, 199-220.	3.4	9
16	Cross-ancestry genomic research: time to close the gap. Neuropsychopharmacology, 2022, 47, 1737-1738.	5.4	7
17	Neuropsychiatric Genetics of Psychosis in the Mexican Population: A Genome-Wide Association Study Protocol for Schizophrenia, Schizoaffective, and Bipolar Disorder Patients and Controls. Complex Psychiatry, 2021, 7, 60-70.	0.9	6
18	The ethics and logistics of field-based genetic paternity studies. Evolutionary Human Sciences, 2020, 2,	1.7	3

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19	Reply to: On powerful GWAS in admixed populations. Nature Genetics, 2021, 53, 1634-1635.	21.4	2
20	153. A Novel Framework for Well-Calibrated Analysis of Complex Traits in Admixed Individuals. Biological Psychiatry, 2019, 85, S63-S64.	1.3	0
21	Gene Discovery in Admixed Cohorts With Tractor. Biological Psychiatry, 2021, 89, S70.	1.3	0
22	<i>The Mysterious World of the Human Genome</i> . By Frank Ryan. Amherst (New York): Prometheus Books. \$28.00. x + 341 p.; ill.; index. ISBN: 978-1-63388-152-5 (hc); 978-1-63388-153-2 (eb). 2016 Quarterly Review of Biology, 2017, 92, 480-480.	0.1	0
23	Overturning the hypothesis for how humans evolved language. TheScienceBreaker, 2019, 05, .	0.0	0