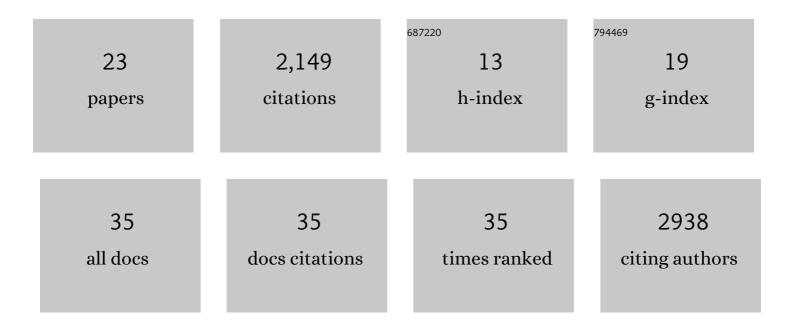
## Elizabeth Atkinson

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5415005/publications.pdf Version: 2024-02-01



FUZABETH ATKINSON

#	Article	IF	CITATIONS
1	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	13.7	929
2	Cross-ancestry genomic research: time to close the gap. Neuropsychopharmacology, 2022, 47, 1737-1738.	2.8	7
3	Characterizing the effect of background selection on the polygenicity of brain-related traits. Genomics, 2021, 113, 111-119.	1.3	24
4	The impact of non-additive genetic associations on age-related complex diseases. Nature Communications, 2021, 12, 2436.	5.8	55
5	Low-coverage sequencing cost-effectively detects known and novel variation in underrepresented populations. American Journal of Human Genetics, 2021, 108, 656-668.	2.6	49
6	Inferring archaic introgression from hominin genetic data. Evolutionary Anthropology, 2021, 30, 199-220.	1.7	9
7	Gene Discovery in Admixed Cohorts With Tractor. Biological Psychiatry, 2021, 89, S70.	0.7	Ο
8	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.	1.3	6
9	Tractor uses local ancestry to enable the inclusion of admixed individuals in GWAS and to boost power. Nature Genetics, 2021, 53, 195-204.	9.4	125
10	Reply to: On powerful GWAS in admixed populations. Nature Genetics, 2021, 53, 1634-1635.	9.4	2
11	Molecular genetic overlap between posttraumatic stress disorder and sleep phenotypes. Sleep, 2020, 43, .	0.6	32
12	The ethics and logistics of field-based genetic paternity studies. Evolutionary Human Sciences, 2020, 2,	0.9	3
13	High rate of extrapair paternity in a human population demonstrates diversity in human reproductive strategies. Science Advances, 2020, 6, eaay6195.	4.7	41
14	International meta-analysis of PTSD genome-wide association studies identifies sex- and ancestry-specific genetic risk loci. Nature Communications, 2019, 10, 4558.	5.8	363
15	153. A Novel Framework for Well-Calibrated Analysis of Complex Traits in Admixed Individuals. Biological Psychiatry, 2019, 85, S63-S64.	0.7	0
16	Overturning the hypothesis for how humans evolved language. TheScienceBreaker, 2019, 05, .	0.0	0
17	Localization of adaptive variants in human genomes using averaged one-dependence estimation. Nature Communications, 2018, 9, 703.	5.8	83
18	No Evidence for Recent Selection at FOXP2 among Diverse Human Populations. Cell, 2018, 174, 1424-1435.e15.	13.5	99

#	Article	IF	CITATIONS
19	An Unexpectedly Complex Architecture for Skin Pigmentation in Africans. Cell, 2017, 171, 1340-1353.e14.	13.5	134
20	<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.	0.0	0
21	Evolutionary and developmental implications of asymmetric brain folding in a large primate pedigree. Evolution; International Journal of Organic Evolution, 2016, 70, 707-715.	1.1	9
22	Population and Evolutionary Genomics of <i>Amblyomma americanum</i> , an Expanding Arthropod Disease Vector. Genome Biology and Evolution, 2016, 8, 1351-1360.	1.1	70
23	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.	1.2	48