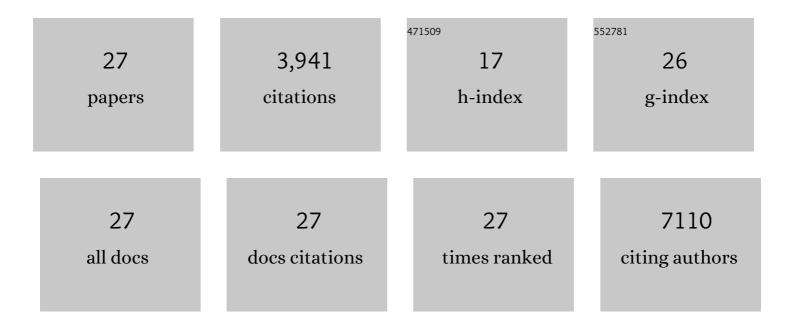
Ondrej Libiger

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Impacts of personal DNA ancestry testing. Journal of Community Genetics, 2021, 12, 37-52.	1.2	12
2	Anxiety is related to indices of cortical maturation in typically developing children and adolescents. Brain Structure and Function, 2016, 221, 3013-3025.	2.3	43
3	Dyslexia and language impairment associated genetic markers influence cortical thickness and white matter in typically developing children. Brain Imaging and Behavior, 2016, 10, 272-282.	2.1	27
4	The Pediatric Imaging, Neurocognition, and Genetics (PING) Data Repository. Neurolmage, 2016, 124, 1149-1154.	4.2	251
5	Partial Least Squares Regression Can Aid in Detecting Differential Abundance of Multiple Features in Sets of Metagenomic Samples. Frontiers in Genetics, 2015, 6, 350.	2.3	3
6	Family income, parental education and brain structure in children and adolescents. Nature Neuroscience, 2015, 18, 773-778.	14.8	979
7	Chip-based direct genotyping of coding variants in genome wide association studies: Utility, issues and prospects. Gene, 2014, 540, 104-109.	2.2	10
8	Correlation analysis of genetic admixture and social identification with body mass index in a Native American Community. American Journal of Human Biology, 2014, 26, 347-360.	1.6	13
9	The NIH Toolbox Cognition Battery: Results from a large normative developmental sample (PING) Neuropsychology, 2014, 28, 1-10.	1.3	163
10	Admixture and Clinical Phenotypic Variation. Human Heredity, 2014, 77, 73-86.	0.8	20
11	Patterns of population epigenomic diversity. Nature, 2013, 495, 193-198.	27.8	543
12	Complex Patterns of Genomic Admixture within Southern Africa. PLoS Genetics, 2013, 9, e1003309.	3.5	94
13	A Method for Inferring an Individual's Genetic Ancestry and Degree of Admixture Associated with Six Major Continental Populations. Frontiers in Genetics, 2013, 3, 322.	2.3	33
14	Genomic Risk Models Improve Prediction of Longitudinal Lipid Levels in Children and Young Adults. Frontiers in Genetics, 2013, 4, 86.	2.3	6
15	Long-term influence of normal variation in neonatal characteristics on human brain development. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 20089-20094.	7.1	158
16	Multimodal imaging of the self-regulating developing brain. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 19620-19625.	7.1	192
17	Clinical Implications of Human Population Differences in Genome-Wide Rates of Functional Genotypes. Frontiers in Genetics, 2012, 3, 211.	2.3	29
18	Transgenerational Epigenetic Instability Is a Source of Novel Methylation Variants. Science, 2011, 334, 369-373.	12.6	635

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#	Article	IF	CITATIONS
19	AN APPLICATION AND EMPIRICAL COMPARISON OF STATISTICAL ANALYSIS METHODS FOR ASSOCIATING RARE VARIANTS TO A COMPLEX PHENOTYPE. , 2010, , 76-87.		10
20	Statistical analysis strategies for association studies involving rare variants. Nature Reviews Genetics, 2010, 11, 773-785.	16.3	426
21	Comparison of Genetic Distance Measures Using Human SNP Genotype Data. Human Biology, 2009, 81, 389-406.	0.2	29
22	Accommodating Linkage Disequilibrium in Genetic-Association Analyses via Ridge Regression. American Journal of Human Genetics, 2008, 82, 375-385.	6.2	115
23	Generalized Analysis of Molecular Variance. PLoS Genetics, 2007, 3, e51.	3.5	75
24	Detecting genetic variation in microarray expression data. Genome Research, 2007, 17, 1228-1235.	5.5	10
25	DNA variation and brain region-specific expression profiles exhibit different relationships between inbred mouse strains: implications for eQTL mapping studies. Genome Biology, 2007, 8, R25.	9.6	57
26	A simulation-based analysis of chromosome segment sharing among a group of arbitrarily related individuals. European Journal of Human Genetics, 2007, 15, 1260-1268.	2.8	6
27	Simulation-based homozygosity mapping with the GAW14 COGA dataset on alcoholism. BMC Genetics, 2005, 6, S35.	2.7	2