Andrew J Pakstis

List of Publications by Year
in descending order

40 papers	2,656 citations	394286 19 h-index	345118 36 g-index
40 all docs	40 docs citations	40 times ranked	2331 citing authors

#	Article	IF	Citations
1	The world-wide distribution of allele frequencies at the human dopamine D4 receptor locus. Human Genetics, 1996, 98, 91-101.	1.8	429
2	Evidence against linkage of schizophrenia to markers on chromosome 5 in a northern Swedish pedigree. Nature, 1988, 336, 167-170.	13.7	405
3	Progress toward an efficient panel of SNPs for ancestry inference. Forensic Science International: Genetics, 2014, 10, 23-32.	1.6	211
4	A global survey of haplotype frequencies and linkage disequilibrium at the DRD2 locus. Human Genetics, 1998, 103, 211-227.	1.8	197
5	SNPs for a universal individual identification panel. Human Genetics, 2010, 127, 315-324.	1.8	194
6	Current sequencing technology makes microhaplotypes a powerful new type of genetic marker for forensics. Forensic Science International: Genetics, 2014, 12, 215-224.	1.6	182
7	Evaluating 130 microhaplotypes across a global set of 83 populations. Forensic Science International: Genetics, 2017, 29, 29-37.	1.6	117
8	Copy Number Variation in Obsessive-Compulsive Disorder and Tourette Syndrome: A Cross-Disorder Study. Journal of the American Academy of Child and Adolescent Psychiatry, 2014, 53, 910-919.	0.3	111
9	Candidate SNPs for a universal individual identification panel. Human Genetics, 2007, 121, 305-317.	1.8	96
10	Genome scan for linkage to Gilles de la Tourette syndrome. , 1999, 88, 437-445.		82
11	DRD2 Haplotypes Containing the TaqI A1 Allele: Implications for Alcoholism Research. Alcoholism: Clinical and Experimental Research, 1996, 20, 697-705.	1.4	63
12	A panel of 74 AISNPs: Improved ancestry inference within Eastern Asia. Forensic Science International: Genetics, 2016, 23, 101-110.	1.6	63
13	Ancestry inference of 96 population samples using microhaplotypes. International Journal of Legal Medicine, 2018, 132, 703-711.	1.2	48
14	Mini-haplotypes as lineage informative SNPs and ancestry inference SNPs. European Journal of Human Genetics, 2012, 20, 1148-1154.	1.4	45
15	Validation of novel forensic DNA markers using multiplex microhaplotype sequencing. Forensic Science International: Genetics, 2020, 47, 102275.	1.6	42
16	52 additional reference population samples for the 55 AISNP panel. Forensic Science International: Genetics, 2015, 19, 269-271.	1.6	41
17	Improving ancestry distinctions among Southwest Asian populations. Forensic Science International: Genetics, 2018, 35, 14-20.	1.6	40
18	Selecting microhaplotypes optimized for different purposes. Electrophoresis, 2018, 39, 2815-2823.	1.3	39

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19	Increasing the reference populations for the 55 AISNP panel: the need and benefits. International Journal of Legal Medicine, 2017, 131, 913-917.	1.2	38
20	Genetic variation in Tunisia in the context of human diversity worldwide. American Journal of Physical Anthropology, 2016, 161, 62-71.	2.1	29
21	Genetic relationships of European, Mediterranean, and SW Asian populations using a panel of 55 AISNPs. European Journal of Human Genetics, 2019, 27, 1885-1893.	1.4	22
22	The redesigned Forensic Research/Reference on Genetics-knowledge base, FROG-kb. Forensic Science International: Genetics, 2018, 33, 33-37.	1.6	21
23	Development of a map of chromosome 11p. Genetic Epidemiology, 1986, 3, 153-158.	0.6	17
24	Detection of a large CTG/CAG trinucleotide repeat expansion in a Danish schizophrenia kindred. , 1997, 74, 546-548.		15
25	Population relationships based on 170 ancestry SNPs from the combined Kidd and Seldin panels. Scientific Reports, 2019, 9, 18874.	1.6	15
26	The population genetics characteristics of a 90 locus panel of microhaplotypes. Human Genetics, 2021, 140, 1753-1773.	1.8	15
27	Mongolians in the Genetic Landscape of Central Asia: Exploring the Genetic Relations among Mongolians and Other World Populations. Human Biology, 2015, 87, 73.	0.4	14
28	ALFRED: An Allele Frequency Database for Microevolutionary Studies. Evolutionary Bioinformatics, 2005, 1, 117693430500100.	0.6	12
29	ALFRED: A WEB-ACCESSIBLE ALLELE FREQUENCY DATABASE. , 1999, , 639-50.		12
30	The distinctive geographic patterns of common pigmentation variants at the OCA2 gene. Scientific Reports, 2020, 10, 15433.	1.6	8
31	FrogAncestryCalc: A standalone batch likelihood computation tool for ancestry inference panels catalogued in FROG-kb. Forensic Science International: Genetics, 2020, 46, 102237.	1.6	8
32	Usefulness of COMT gene polymorphisms in North African populations. Gene, 2019, 696, 186-196.	1.0	7
33	Genetic relationships of Southwest Asian and Mediterranean populations. Forensic Science International: Genetics, 2021, 53, 102528.	1.6	7
34	North Asian population relationships in a global context. Scientific Reports, 2022, 12, 7214.	1.6	3
35	A more powerful method to evaluate p-values in GENEHUNTER. Genetic Epidemiology, 1999, 17, S415-S420.	0.6	2
36	Genetic diversity of the North African population revealed by the typing of SNPs in the DRD2/ANKK1 genomic region. Gene, 2021, 777, 145466.	1.0	2

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37	New Insight into the human genetic diversity in North African populations by genotyping of <scp>SNPs</scp> in <scp>DRD3</scp>, <scp>CSMD1</scp> and <scp>NRG1</scp> genes. Molecular Genetics & Genomic Medicine, 2022, 10, e1871.	0.6	2
38	Dinucleotide polymorphism at the DXS1178 locus is tightly linked to PGK1 at Xq13. Human Genetics, 1995, 95, 467-8.	1.8	1
39	STAT3 polymorphisms in North Africa and its implication in breast cancer. Molecular Genetics & Genomic Medicine, 2021, 9, e1744.	0.6	1
40	Genetic analysis workshop III: Sampling considerations and assumptions in gene mapping. Genetic Epidemiology, 1985, 2, 219-220.	0.6	0