

A Kong

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

2,973
citations

686830

13
h-index

940134

16
g-index

16
all docs

16
docs citations

16
times ranked

3558
citing authors

#	ARTICLE	IF	CITATIONS
1	Allele-Sharing Models: LOD Scores and Accurate Linkage Tests. American Journal of Human Genetics, 1997, 61, 1179-1188.	2.6	931
2	Allegro, a new computer program for multipoint linkage analysis. Nature Genetics, 2000, 25, 12-13.	9.4	737
3	Loci on chromosomes 2 (NIDDM1) and 15 interact to increase susceptibility to diabetes in Mexican Americans. Nature Genetics, 1999, 21, 213-215.	9.4	374
4	Familial Aggregation of Parkinson's Disease in Iceland. New England Journal of Medicine, 2000, 343, 1765-1770.	13.9	271
5	A Genome-Wide Scan Reveals a Maternal Susceptibility Locus for Pre-Eclampsia on Chromosome 2p13. Human Molecular Genetics, 1999, 8, 1799-1805.	1.4	196
6	A genome-wide scan for preeclampsia in the Netherlands. European Journal of Human Genetics, 2001, 9, 758-764.	1.4	140
7	Graphical templates for model registration. IEEE Transactions on Pattern Analysis and Machine Intelligence, 1996, 18, 225-236.	9.7	104
8	Allelic Frequencies and Patterns of Single-nucleotide Polymorphisms in Candidate Genes for Asthma and Atopy in Iceland. American Journal of Respiratory and Critical Care Medicine, 2001, 164, 2036-2044.	2.5	85
9	CDKN2A mutations and melanoma risk in the Icelandic population. Journal of Medical Genetics, 2008, 45, 284-289.	1.5	30
10	The role of linkage studies for common diseases. Current Opinion in Genetics and Development, 2001, 11, 264-267.	1.5	26
11	Asymptotic theory for gene mapping.. Proceedings of the National Academy of Sciences of the United States of America, 1994, 91, 9705-9709.	3.3	22
12	Sequential imputation and multipoint linkage analysis. Genetic Epidemiology, 1993, 10, 483-488.	0.6	18
13	Efficient methods for computing linkage likelihoods of recessive diseases in inbred pedigrees. Genetic Epidemiology, 1991, 8, 81-103.	0.6	16
14	Importance sampling. I. Computing multimodel p values in linkage analysis. American Journal of Human Genetics, 1992, 51, 1413-29.	2.6	13
15	Linkage analysis with adjustment for covariates: A method combining peeling with Gibbs sampling. Cytogenetic and Genome Research, 1992, 59, 208-210.	0.6	6
16	On a Randomization Procedure. American Journal of Human Genetics, 2000, 67, 1352-1355.	2.6	4