Tero Hiekkalinna

List of Publications by Year
in descending order

30 papers	2,110 citations	331670 21 h-index	454955 30 g-index
30 all docs	30 docs citations	30 times ranked	4375 citing authors

#	Article	IF	Citations
1	NCOR2 is a novel candidate gene for migraine-epilepsy phenotype. Cephalalgia, 2022, 42, 631-644.	3.9	6
2	An epigenome-wide association study of metabolic syndrome and its components. Scientific Reports, 2020, 10, 20567.	3.3	27
3	Life is a simulation of life â€“ or is it?. Evolutionary Anthropology, 2017, 26, 151-156.	3.4	1
4	AMIGO-Kv2.1 Potassium Channel Complex is Associated With Schizophrenia-Related Phenotypes. Schizophrenia Bulletin, 2016, 42, sbv105.	4.3	25
5	A genome-wide screen for acrophobia susceptibility loci in a Finnish isolate. Scientific Reports, 2016, 6, 39345.	3.3	2
6	DataSHIELD: taking the analysis to the data, not the data to the analysis. International Journal of Epidemiology, 2014, 43, 1929-1944.	1.9	188
7	Identifying flavor preference subgroups. Genetic basis and related eating behavior traits. Appetite, 2014, 75, 1-10.	3.7	59
8	PSEUDOMARKER 2.0: efficient computation of likelihoods using NOMAD. BMC Bioinformatics, 2014, 15, 47.	2.6	26
9	The prevalence of metabolic syndrome and metabolically healthy obesity in Europe: a collaborative analysis of ten large cohort studies. BMC Endocrine Disorders, 2014, 14, 9.	2.2	440
10	On the statistical properties of family-based association tests in datasets containing both pedigrees and unrelated caseâ€“control samples. European Journal of Human Genetics, 2012, 20, 217-223.	2.8	11
11	Mortality Rate Increases Steeply With Nonadherence to Statin Therapy in Patients With Acute Coronary Syndrome. Clinical Cardiology, 2012, 35, E22-7.	1.8	29
12	On the Validity of the Likelihood Ratio Test and Consistency of Resulting Parameter Estimates in Joint Linkage and Linkage Disequilibrium Analysis under Improperly Specified Parametric Models. Annals of Human Genetics, 2012, 76, 63-73.	0.8	5
13	PSEUDOMARKER: A Powerful Program for Joint Linkage and/or Linkage Disequilibrium Analysis on Mixtures of Singletons and Related Individuals. Human Heredity, 2011, 71, 256-266.	0.8	39
14	Novel Susceptibility Locus at 22q11 for Diabetic Nephropathy in Type 1 Diabetes. PLoS ONE, 2011, 6, e24053.	2.5	12
15	Linkage analysis of schizophrenia controlling for population substructure. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2009, 150B, 827-835.	1.7	12
16	Genome-wide linkage screen for stature and body mass index in 3.032 families: evidence for sex- and population-specific genetic effects. European Journal of Human Genetics, 2009, 17, 258-266.	2.8	16
17	Combined Genome Scans for Body Stature in 6,602 European Twins: Evidence for Common Caucasian Loci. PLoS Genetics, 2007, 3, e97.	3.5	145
18	Families with the risk allele of DISC1 reveal a link between schizophrenia and another component of the same molecular pathway, NDE1. Human Molecular Genetics, 2007, 16, 453-462.	2.9	74

#	Article	IF	Citations
19	Genetic component of identification, intensity and pleasantness of odours: a Finnish family study. European Journal of Human Genetics, 2007, 15, 596-602.	2.8	32
20	An utter refutation of the â€˜Fundamental Theorem of the HapMapâ€™. European Journal of Human Genetics, 2006, 14, 426-437.	2.8	164
21	AUTOGSCAN: Powerful Tools for Automated Genome-Wide Linkage and Linkage Disequilibrium Analysis. Twin Research and Human Genetics, 2005, 8, 16-21.	0.6	25
22	AUTOGSCAN: Powerful Tools for Automated Genome-Wide Linkage and Linkage Disequilibrium Analysis. Twin Research and Human Genetics, 2005, 8, 16-21.	0.6	17
23	Search for cognitive trait components of schizophrenia reveals a locus for verbal learning and memory on 4q and for visual working memory on 2q. Human Molecular Genetics, 2004, 13, 1693-1702.	2.9	74
24	Locus for quantitative HDL-cholesterol on chromosome 10q in Finnish families with dyslipidemia. Journal of Lipid Research, 2004, 45, 1876-1884.	4.2	22
25	Evidence of susceptibility loci on 4q32 and 16p12 for bipolar disorder. Human Molecular Genetics, 2003, 12, 1907-1915.	2.9	70
26	A Susceptibility Locus for Migraine with Aura, on Chromosome 4q24. American Journal of Human Genetics, 2002, 70, 652-662.	6.2	146
27	Genome Scans Provide Evidence for Low-HDL-C Loci on Chromosomes 8q23, 16q24.1-24.2, and 20q13.11 in Finnish Families. American Journal of Human Genetics, 2002, 70, 1333-1340.	6.2	91
28	Quantitative-Trait-Locus Analysis of Body-Mass Index and of Stature, by Combined Analysis of Genome Scans of Five Finnish Study Groups. American Journal of Human Genetics, 2001, 69, 117-123.	6.2	111
29	Genome-wide scan of predisposing loci for increased diastolic blood pressure in Finnish siblings. Journal of Hypertension, 2000, 18, 1579-1585.	0.5	104
30	Genomewide Scan for Familial Combined Hyperlipidemia Genes in Finnish Families, Suggesting Multiple Susceptibility Loci Influencing Triglyceride, Cholesterol, and Apolipoprotein B Levels. American Journal of Human Genetics, 1999, 64, 1453-1463.	6.2	137