Xiao-Qing Liu

List of Publications by Year
in descending order

25 papers	6,055 citations	361413 20 h-index	580821 25 g-index
25 all docs	25 docs citations	25 times ranked	8707 citing authors

#	Article	IF	Citations
1	Identity-by-descent mapping for diastolic blood pressure in unrelated Mexican Americans. BMC Proceedings, 2016, 10, 263-267.	1.6	4
2	The Effects of Anemia and Blood Transfusion on Patients With Stage III-IV Ovarian Cancer. International Journal of Gynecological Cancer, 2013, 23, 1569-1576.	2.5	19
3	Measurement equivalence of the autism symptom phenotype in children and youth. Journal of Child Psychology and Psychiatry and Allied Disciplines, 2013, 54, 1346-1355.	5.2	10
4	Individual common variants exert weak effects on the risk for autism spectrum disorders. Human Molecular Genetics, 2012, 21, 4781-4792.	2.9	334
5	ABCG2 null alleles define the Jr(aâ^') blood group phenotype. Nature Genetics, 2012, 44, 131-132.	21.4	77
6	Sex differences in repetitive stereotyped behaviors in autism: Implications for genetic liability. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2012, 159B, 5-12.	1.7	154
7	Identification of Genetic Loci Underlying the Phenotypic Constructs of Autism Spectrum Disorders. Journal of the American Academy of Child and Adolescent Psychiatry, 2011, 50, 687-696.e13.	0.5	26
8	Postpartum Septic Sacroiliitis Misdiagnosed as Sciatic Neuropathy. American Journal of the Medical Sciences, 2010, 339, 292-295.	1.1	26
9	Functional impact of global rare copy number variation in autism spectrum disorders. Nature, 2010, 466, 368-372.	27.8	1,803
10	A genome-wide scan for common alleles affecting risk for autism. Human Molecular Genetics, 2010, 19, 4072-4082.	2.9	538
11	Genetic Variation of DKK3 May Modify Renal Disease Severity in ADPKD. Journal of the American Society of Nephrology: JASN, 2010, 21, 1510-1520.	6.1	59
12	A genome-wide linkage and association scan reveals novel loci for autism. Nature, 2009, 461, 802-808.	27.8	570
13	Genome-wide Linkage Analyses of Quantitative and Categorical Autism Subphenotypes. Biological Psychiatry, 2008, 64, 561-570.	1.3	80
14	IL5RA and TNFRSF6B Gene Variants Are Associated With Sporadic IgA Nephropathy. Journal of the American Society of Nephrology: JASN, 2008, 19, 1025-1033.	6.1	30
15	Multiple Superoxide Dismutase 1/Splicing Factor Serine Alanine 15 Variants Are Associated With the Development and Progression of Diabetic Nephropathy. Diabetes, 2008, 57, 218-228.	0.6	89
16	Multiple Variants in Vascular Endothelial Growth Factor (VEGFA) Are Risk Factors for Time to Severe Retinopathy in Type 1 Diabetes. Diabetes, 2007, 56, 2161-2168.	0.6	88
17	Genome-Wide Linkage Scan of a Large Family with IgA Nephropathy Localizes a Novel Susceptibility Locus to Chromosome 2q36. Journal of the American Society of Nephrology: JASN, 2007, 18, 2408-2415.	6.1	112
18	The Novel Neuronal Ceroid Lipofuscinosis Gene MFSD8 Encodes a Putative Lysosomal Transporter. American Journal of Human Genetics, 2007, 81, 136-146.	6.2	180

#	Article	IF	Citations
19	Mapping autism risk loci using genetic linkage and chromosomal rearrangements. Nature Genetics, 2007, 39, 319-328.	21.4	1,272
20	The role of Self-Defined Race/Ethnicity in Population Structure Control. Annals of Human Genetics, 2006, 70, 496-505.	0.8	24
21	Assessing the validity of the association between the SUMO4 M55V variant and risk of type 1 diabetes. Nature Genetics, 2005, 37, 111-112.	21.4	47
22	A genome scan for parent-of-origin linkage effects in alcoholism. BMC Genetics, 2005, 6, S160.	2.7	11
23	Genetic analysis of common factors underlying cardiovascular disease-related traits. BMC Genetics, 2003, 4, S56.	2.7	11
24	Dimensions of Religiosity and Their Relationship to Lifetime Psychiatric and Substance Use Disorders. American Journal of Psychiatry, 2003, 160, 496-503.	7.2	425
25	Stepwise formation of a SMAD activity gradient during dorsal-ventral patterning of the Drosophila embryo. Development (Cambridge), 2003, 130, 5705-5716.	2.5	66