Sarah G Buxbaum

List of Publications by Year in descending order

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172457 206112 4,714 50 29 48 citations h-index g-index papers 50 50 50 10604 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	HMG-coenzyme A reductase inhibition, type 2 diabetes, and bodyweight: evidence from genetic analysis and randomised trials. Lancet, The, 2015, 385, 351-361.	13.7	562
2	Association between alcohol and cardiovascular disease: Mendelian randomisation analysis based on individual participant data. BMJ, The, 2014, 349, g4164-g4164.	6.0	528
3	Novel Loci for Adiponectin Levels and Their Influence on Type 2 Diabetes and Metabolic Traits: A Multi-Ethnic Meta-Analysis of 45,891 Individuals. PLoS Genetics, 2012, 8, e1002607.	3 . 5	419
4	The landscape of recombination in African Americans. Nature, 2011, 476, 170-175.	27.8	319
5	A Quantitative-Trait Analysis of Human Plasma–Dopamine β-Hydroxylase Activity: Evidence for a Major Functional Polymorphism at the DBH Locus. American Journal of Human Genetics, 2001, 68, 515-522.	6.2	253
6	Large-Scale Gene-Centric Meta-analysis across 32 Studies Identifies Multiple Lipid Loci. American Journal of Human Genetics, 2012, 91, 823-838.	6.2	227
7	A Whole-Genome Scan for Obstructive Sleep Apnea and Obesity. American Journal of Human Genetics, 2003, 72, 340-350.	6.2	212
8	Causal Effects of Body Mass Index on Cardiometabolic Traits and Events: A Mendelian Randomization Analysis. American Journal of Human Genetics, 2014, 94, 198-208.	6.2	199
9	Admixture Mapping of White Cell Count: Genetic Locus Responsible for Lower White Blood Cell Count in the Health ABC and Jackson Heart Studies. American Journal of Human Genetics, 2008, 82, 81-87.	6.2	172
10	Whole Genome Scan for Obstructive Sleep Apnea and Obesity in African-American Families. American Journal of Respiratory and Critical Care Medicine, 2004, 169, 1314-1321.	5.6	163
11	Genome Scan for Familial Abdominal Aortic Aneurysm Using Sex and Family History as Covariates Suggests Genetic Heterogeneity and Identifies Linkage to Chromosome 19q13. Circulation, 2004, 109, 2103-2108.	1.6	120
12	Genetics of the apnea hypopnea index in Caucasians and African Americans: I. Segregation analysis. Genetic Epidemiology, 2002, 22, 243-253.	1.3	116
13	Mendelian Randomization Studies Do Not Support a Causal Role for Reduced Circulating Adiponectin Levels in Insulin Resistance and Type 2 Diabetes. Diabetes, 2013, 62, 3589-3598.	0.6	116
14	Enhanced Statistical Tests for GWAS in Admixed Populations: Assessment using African Americans from CARe and a Breast Cancer Consortium. PLoS Genetics, 2011, 7, e1001371.	3.5	110
15	Genome-wide meta-analyses of smoking behaviors in African Americans. Translational Psychiatry, 2012, 2, e119-e119.	4.8	94
16	Interaction of sleep quality and psychosocial stress on obesity in African Americans: the Cardiovascular Health Epidemiology Study (CHES). BMC Public Health, 2010, 10, 581.	2.9	91
17	The Structure of Linkage Disequilibrium at the DBH Locus Strongly Influences the Magnitude of Association between Diallelic Markers and Plasma Dopamine \hat{l}^2 -Hydroxylase Activity. American Journal of Human Genetics, 2003, 72, 1389-1400.	6.2	81
18	Association of Genetic Loci with Sleep Apnea in European Americans and African-Americans: The Candidate Gene Association Resource (CARe). PLoS ONE, 2012, 7, e48836.	2. 5	64

#	Article	IF	CITATIONS
19	Genome-wide association study of coronary artery calcified atherosclerotic plaque in African Americans with type 2 diabetes. BMC Genetics, 2017, 18, 105.	2.7	54
20	The Triglyceride/High-Density Lipoprotein Cholesterol Ratio Fails to Predict Insulin Resistance in African-American Women: An Analysis of Jackson Heart Study. Metabolic Syndrome and Related Disorders, 2010, 8, 511-514.	1.3	51
21	Screening for familial and hereditary prostate cancer. International Journal of Cancer, 2016, 138, 2579-2591.	5.1	49
22	Leptinemia and its association with stroke and coronary heart disease in the Jackson Heart Study. Clinical Endocrinology, 2010, 72, 32-37.	2.4	48
23	Novel Loci Associated With PR Interval in a Genome-Wide Association Study of 10 African American Cohorts. Circulation: Cardiovascular Genetics, 2012, 5, 639-646.	5.1	48
24	Multi-Ethnic Analysis of Lipid-Associated Loci: The NHLBI CARe Project. PLoS ONE, 2012, 7, e36473.	2.5	46
25	A Single Nucleotide Polymorphism at DBH, Possibly Associated with Attention-Deficit/Hyperactivity Disorder, Associates with Lower Plasma Dopamine β-Hydroxylase Activity and is in Linkage Disequilibrium with Two Putative Functional Single Nucleotide Polymorphisms. Biological Psychiatry, 2006, 60, 1034-1038.	1.3	42
26	Clinical Correlates and Heritability of QT Interval Duration in Blacks. Circulation: Arrhythmia and Electrophysiology, 2009, 2, 427-432.	4.8	40
27	Common variants in <i>DRD2</i> i>are associated with sleep duration: the CARe consortium. Human Molecular Genetics, 2016, 25, 167-179.	2.9	40
28	Gene-Centric Meta-Analysis of Lipid Traits in African, East Asian and Hispanic Populations. PLoS ONE, 2012, 7, e50198.	2.5	40
29	Impact of Ancestry and Common Genetic Variants on QT Interval in African Americans. Circulation: Cardiovascular Genetics, 2012, 5, 647-655.	5.1	38
30	Fine-mapping, novel loci identification, and SNP association transferability in a genome-wide association study of QRS duration in African Americans. Human Molecular Genetics, 2016, 25, 4350-4368.	2.9	37
31	Screening and kinetic analysis of recombinant anti-CEA antibody fragments. Journal of Immunological Methods, 1995, 183, 119-125.	1.4	30
32	Common genetic variation near the connexin-43 gene is associated with resting heart rate in African Americans: A genome-wide association study of 13,372 participants. Heart Rhythm, 2013, 10, 401-408.	0.7	29
33	Fine-Mapping in African Americans of 8 Recently Discovered Genetic Loci for Plasma Lipids. Circulation: Cardiovascular Genetics, 2010, 3, 358-364.	5.1	28
34	Interactive Influences of Ethnicity, Endothelin-1 Gene, and Everyday Discrimination Upon Nocturnal Ambulatory Blood Pressure. Annals of Behavioral Medicine, 2013, 45, 377-386.	2.9	28
35	SYSTEMS BIOLOGY ANALYSES OF GENE EXPRESSION AND GENOME WIDE ASSOCIATION STUDY DATA IN OBSTRUCTIVE SLEEP APNEA. , 2010, , 14-25.		25
36	Adhesion molecules, endothelin-1 and lung function in seven population-based cohorts. Biomarkers, 2013, 18, 196-203.	1.9	23

#	Article	IF	CITATIONS
37	Lipids, obesity and gallbladder disease in women: insights from genetic studies using the cardiovascular gene-centric 50K SNP array. European Journal of Human Genetics, 2016, 24, 106-112.	2.8	23
38	SCN5A Variation Is Associated With Electrocardiographic Traits in the Jackson Heart Study. Circulation: Cardiovascular Genetics, 2011, 4, 139-144.	5.1	22
39	Communication and technology in genetic counseling for familial cancer. Clinical Genetics, 2014, 85, 213-222.	2.0	21
40	Using the PRECEDE-PROCEED model for an online peer-to-peer suicide prevention and awareness for depression (SPAD) intervention among African American college students: experimental study. Health Promotion Perspectives, 2018, 8, 15-24.	1.9	21
41	Comparative Study of Multipoint Methods for Genotype Error Detection. Human Heredity, 2004, 58, 175-189.	0.8	17
42	Testing for Colon Neoplasia Susceptibility Variants at the Human COX2 Locus. Journal of the National Cancer Institute, 2001, 93, 635-639.	6.3	15
43	Associations of adiponectin with individual European ancestry in African Americans: the Jackson Heart Study. Frontiers in Genetics, 2014, 5, 22.	2.3	13
44	HIV/AIDS Knowledge of Undergraduate Students at a Historically Black College and University. Diseases (Basel, Switzerland), 2018, 6, 98.	2.5	12
45	Genetic Variability of Smoking Persistence in African Americans. Cancer Prevention Research, 2011, 4, 729-734.	1.5	11
46	Determination of binding constants of diabodies directed against prostate-specific antigen using electrochemiluminescence-based immunoassays., 1996, 9, 456-461.		7
47	Using Social Cognitive Theory to Predict Medication Compliance Behavior in Patients with Depression in Southern United States in 2016 in a Cross-Sectional Study. Journal of Caring Sciences, 2018, 7, 1-8.	1.0	6
48	Predicting Quantitative Trait Levels by Modeling SNP Interaction. Genetic Epidemiology, 2001, 21, S608-S613.	1.3	2
49	Admixture Mapping of White Cell Count: Genetic Locus Responsible for Lower White Blood Cell Count in the Health ABC and Jackson Heart Studies. American Journal of Human Genetics, 2008, 82, 532.	6.2	1
50	Genetic Analysis of a Population Heavy Drinking Phenotype Identifies Risk Variants in Whites. Journal of Clinical Psychopharmacology, 2013, 33, 206-210.	1.4	1