

# Genome-wide meta-analysis identifies 11 new loci for an insights into genetic architecture

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Citation Report

#	ARTICLE	IF	CITATIONS
3	Meta-analysis of Gene-Level Associations for Rare Variants Based on Single-Variant Statistics. American Journal of Human Genetics, 2013, 93, 236-248.	2.6	60
4	Bipolar disorder risk alleles in children with ADHD. Journal of Neural Transmission, 2013, 120, 1611-1617.	1.4	15
5	The Next-Generation Sequencing Revolution and Its Impact on Genomics. Cell, 2013, 155, 27-38.	13.5	856
6	Contribution of 32 GWAS-Identified Common Variants to Severe Obesity in European Adults Referred for Bariatric Surgery. PLoS ONE, 2013, 8, e70735.	1.1	39
7	A genomewide study of body mass index and its genetic correlation with thromboembolic risk. Thrombosis and Haemostasis, 2014, 112, 1036-1043.	1.8	7
8	Marker-Based Estimation of Genetic Parameters in Genomics. PLoS ONE, 2014, 9, e102715.	1.1	6
9	Genome-Wide Association Studies Identify the Loci for 5 Exterior Traits in a Large White Ā— Minzhu Pig Population. PLoS ONE, 2014, 9, e103766.	1.1	32
10	Genome-Wide Association Studies and Heritability Estimates of Body Mass Index Related Phenotypes in Bangladeshi Adults. PLoS ONE, 2014, 9, e105062.	1.1	19
11	Mitochondrial Genetic Variants Identified to Be Associated with BMI in Adults. PLoS ONE, 2014, 9, e105116.	1.1	34
12	Secular Trends in Growth and Nutritional Status of Mozambican School-Aged Children and Adolescents. PLoS ONE, 2014, 9, e114068.	1.1	26
13	Reading and Language Disorders: The Importance of Both Quantity and Quality. Genes, 2014, 5, 285-309.	1.0	41
14	The Association of Common Variants in PCSK1 With Obesity: A HuGE Review and Meta-Analysis. American Journal of Epidemiology, 2014, 180, 1051-1065.	1.6	45
15	How Fatty Acids and Common Genetic Variants Together Affect the Inflammation of Adipose Tissue. Current Cardiovascular Risk Reports, 2014, 8, 1.	0.8	1
16	Combined effect of established BMI loci on obesity-related traits in an Algerian population sample. BMC Genetics, 2014, 15, 128.	2.7	5
17	Meta-analysis of genome-wide association studies in East Asian-ancestry populations identifies four new loci for body mass index. Human Molecular Genetics, 2014, 23, 5492-5504.	1.4	192
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19	Whole-exome imputation of sequence variants identified two novel alleles associated with adult body height in African Americans. Human Molecular Genetics, 2014, 23, 6607-6615.	1.4	14
20	Principal Component Analysis Characterizes Shared Pathogenetics from Genome-Wide Association Studies. PLoS Computational Biology, 2014, 10, e1003820.	1.5	19

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21	Phenome-wide association study (PheWAS) in EMR-linked pediatric cohorts, genetically links PLCL1 to speech language development and IL5-IL13 to Eosinophilic Esophagitis. <i>Frontiers in Genetics</i> , 2014, 5, 401.	1.1	70
22	RTeQTL: Real-Time Online Engine for Expression Quantitative Trait Loci Analyses. <i>Database: the Journal of Biological Databases and Curation</i> , 2014, 2014, bau066-bau066.	1.4	7
23	Novel insights from genetic and epigenetic studies in understanding the complex uraemic phenotype. <i>Nephrology Dialysis Transplantation</i> , 2014, 29, 964-971.	0.4	9
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