

Mapping cis- and trans-regulatory effects across multiple

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Extent, Causes, and Consequences of Small RNA Expression Variation in Human Adipose Tissue. PLoS Genetics, 2012, 8, e1002704.	1.5	48
2	Tissue specificity of genetic regulation of gene expression. Nature Genetics, 2012, 44, 1077-1078.	9.4	16
3	Omics technologies and the study of human ageing. Nature Reviews Genetics, 2013, 14, 601-607.	7.7	108
4	Genome-wide association analysis identifies 13 new risk loci for schizophrenia. Nature Genetics, 2013, 45, 1150-1159.	9.4	1,395
5	Variability in the immune system: of vaccine responses and immune states. Current Opinion in Immunology, 2013, 25, 542-547.	2.4	45
6	Where genotype is not predictive of phenotype: towards an understanding of the molecular basis of reduced penetrance in human inherited disease. Human Genetics, 2013, 132, 1077-1130.	1.8	528
7	Association of Parkinson Disease with Structural and Regulatory Variants in the HLA Region. American Journal of Human Genetics, 2013, 93, 984-993.	2.6	145
8	Coordinated Effects of Sequence Variation on DNA Binding, Chromatin Structure, and Transcription. Science, 2013, 342, 744-747.	6.0	364
9	Cell-type, allelic, and genetic signatures in the human pancreatic beta cell transcriptome. Genome Research, 2013, 23, 1554-1562.	2.4	161
10	Expression quantitative trait loci: present and future. Philosophical Transactions of the Royal Society B: Biological Sciences, 2013, 368, 20120362.	1.8	363
11	Transcriptome study of differential expression in schizophrenia. Human Molecular Genetics, 2013, 22, 5001-5014.	1.4	73
12	Melanoma risk loci as determinants of melanoma recurrence and survival. Journal of Translational Medicine, 2013, 11, 279.	1.8	30
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14	Systematic identification of trans eQTLs as putative drivers of known disease associations. Nature Genetics, 2013, 45, 1238-1243.	9.4	1,544
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18	Next-generation sequencing in understanding complex neurological disease. Expert Review of Neurotherapeutics, 2013, 13, 215-227.	1.4	18

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19	Sherlock: Detecting Gene-Disease Associations by Matching Patterns of Expression QTL and GWAS. <i>American Journal of Human Genetics</i> , 2013, 92, 667-680.	2.6	243
20	Dense genotyping of immune-related disease regions identifies 14 new susceptibility loci for juvenile idiopathic arthritis. <i>Nature Genetics</i> , 2013, 45, 664-669.	9.4	337
21	rs2072135, a low- ϵ penetrance variant for chronic lymphocytic leukaemia?. <i>British Journal of Haematology</i> , 2013, 162, 221-228.	1.2	1
22	How next-generation sequencing is transforming complex disease genetics. <i>Trends in Genetics</i> , 2013, 29, 23-30.	2.9	70
23	The <i>CTRB1/2</i> Locus Affects Diabetes Susceptibility and Treatment via the Incretin Pathway. <i>Diabetes</i> , 2013, 62, 3275-3281.	0.3	96
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25	Genetic Determinants of Trabecular and Cortical Volumetric Bone Mineral Densities and Bone Microstructure. <i>PLoS Genetics</i> , 2013, 9, e1003247.	1.5	100
26	Genotype-Environment Interactions Reveal Causal Pathways That Mediate Genetic Effects on Phenotype. <i>PLoS Genetics</i> , 2013, 9, e1003803.	1.5	72
27	Global Properties and Functional Complexity of Human Gene Regulatory Variation. <i>PLoS Genetics</i> , 2013, 9, e1003501.	1.5	55
28	Congruence of Additive and Non-Additive Effects on Gene Expression Estimated from Pedigree and SNP Data. <i>PLoS Genetics</i> , 2013, 9, e1003502.	1.5	79
29	Preferential Binding to Elk-1 by SLE-Associated IL10 Risk Allele Upregulates IL10 Expression. <i>PLoS Genetics</i> , 2013, 9, e1003870.	1.5	36
30	Trans-Ancestral Studies Fine Map the SLE-Susceptibility Locus TNFSF4. <i>PLoS Genetics</i> , 2013, 9, e1003554.	1.5	50
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32	Genome-wide expression quantitative trait loci analysis in asthma. <i>Current Opinion in Allergy and Clinical Immunology</i> , 2013, 13, 487-494.	1.1	19
33	CDKN2B expression in adipose tissue of familial combined hyperlipidemia patients. <i>Journal of Lipid Research</i> , 2013, 54, 3491-3505.	2.0	23
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56	A Role for Noncoding Variation in Schizophrenia. <i>Cell Reports</i> , 2014, 9, 1417-1429.	2.9	225
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70	Genetic Determinants of Circulating Interleukin-1 Receptor Antagonist Levels and Their Association With Glycemic Traits. <i>Diabetes</i> , 2014, 63, 4343-4359.	0.3	40
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134	Expression Quantitative Trait Loci Information Improves Predictive Modeling of Disease Relevance of Non-Coding Genetic Variation. <i>PLoS ONE</i> , 2015, 10, e0140758.	1.1	17
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144	Association of breast cancer risk loci with breast cancer survival. <i>International Journal of Cancer</i> , 2015, 137, 2837-2845.	2.3	33
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154	Variation in <i>PPARG</i> Is Associated With Longitudinal Change in Insulin Resistance in Mexican Americans at Risk for Type 2 Diabetes. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2015, 100, 1187-1195.	1.8	18
155	The role of regulatory variation in complex traits and disease. <i>Nature Reviews Genetics</i> , 2015, 16, 197-212.	7.7	864
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161	Dissecting the genetics of the human transcriptome identifies novel trait-related <i>trans</i> -eQTLs and corroborates the regulatory relevance of non-protein coding loci. <i>Human Molecular Genetics</i> , 2015, 24, 4746-4763.	1.4	94
162	A trans-ethnic genome-wide association study identifies gender-specific loci influencing pediatric aBMD and BMC at the distal radius. <i>Human Molecular Genetics</i> , 2015, 24, 5053-5059.	1.4	48
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