Claire Bellis

List of Publications by Year in descending order

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44 papers ci

9,891 citations

257450 24 h-index 223800 46 g-index

48 all docs 48 docs citations

48 times ranked 18149 citing authors

#	Article	IF	CITATIONS
1	Large-Scale Whole-Genome Sequencing of Three Diverse Asian Populations in Singapore. Cell, 2019, 179, 736-749.e15.	28.9	126
2	BDNF Variants May Modulate Long-Term Visual Memory Performance in a Healthy Cohort. International Journal of Molecular Sciences, 2017, 18, 655.	4.1	19
3	Genome-wide physical activity interactions in adiposity ― A meta-analysis of 200,452 adults. PLoS Genetics, 2017, 13, e1006528.	3.5	158
4	A principal component meta-analysis on multiple anthropometric traits identifies novel loci for body shape. Nature Communications, 2016, 7, 13357.	12.8	74
5	New loci for body fat percentage reveal link between adiposity and cardiometabolic disease risk. Nature Communications, 2016, 7, 10495.	12.8	245
6	Genome-wide meta-analysis uncovers novel loci influencing circulating leptin levels. Nature Communications, 2016, 7, 10494.	12.8	153
7	Association of the microRNA-Single Nucleotide Polymorphism rs2910164 in miR146a with sporadic breast cancer susceptibility: A case control study. Gene, 2016, 576, 256-260.	2.2	20
8	Serum bilirubin concentration is modified by UGT1A1 Haplotypes and influences risk of Type-2 diabetes in the Norfolk Island genetic isolate. BMC Genetics, 2015, 16, 136.	2.7	7
9	â€~Mutiny on the Bounty': the genetic history of Norfolk Island reveals extreme gender-biased admixture. Investigative Genetics, 2015, 6, 11.	3.3	9
10	The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. PLoS Genetics, 2015, 11, e1005378.	3.5	331
11	New genetic loci link adipose and insulin biology to body fat distribution. Nature, 2015, 518, 187-196.	27.8	1,328
12	Genetic studies of body mass index yield new insights for obesity biology. Nature, 2015, 518, 197-206.	27.8	3,823
13	Interaction between Common Genetic Variants and Total Fat Intake on Low-Density Lipoprotein Peak Particle Diameter: A Genome-Wide Association Study. Journal of Nutrigenetics and Nutrigenomics, 2015, 8, 44-53.	1.3	24
14	Novel loci associated with usual sleep duration: the CHARGE Consortium Genome-Wide Association Study. Molecular Psychiatry, 2015, 20, 1232-1239.	7.9	112
15	A Phenomic Scan of the Norfolk Island Genetic Isolate Identifies a Major Pleiotropic Effect Locus Associated with Metabolic and Renal Disorder Markers. PLoS Genetics, 2015, 11, e1005593.	3.5	3
16	Human Plasma Lipidome Is Pleiotropically Associated With Cardiovascular Risk Factors and Death. Circulation: Cardiovascular Genetics, 2014, 7, 854-863.	5.1	56
17	Geneâ€byâ€age effects on BMI from birth to adulthood: The fels longitudinal study. Obesity, 2014, 22, 875-881.	3.0	18
18	Plasma dihydroceramide species associate with waist circumference in Mexican American families. Obesity, 2014, 22, 950-956.	3.0	32

#	Article	IF	Citations
19	Plasma lipidome is independently associated with variability in metabolic syndrome in Mexican American families. Journal of Lipid Research, 2014, 55, 939-946.	4.2	12
20	Defining the role of common variation in the genomic and biological architecture of adult human height. Nature Genetics, 2014, 46, 1173-1186.	21.4	1,818
21	Variability in Associations of Phosphatidylcholine Molecular Species with Metabolic Syndrome in Mexican–American Families. Lipids, 2013, 48, 497-503.	1.7	15
22	Mapping eQTLs in the Norfolk Island Genetic Isolate Identifies Candidate Genes for CVD Risk Traits. American Journal of Human Genetics, 2013, 93, 1087-1099.	6.2	28
23	Genetic risk for earlier menarche also influences peripubertal body mass index. American Journal of Physical Anthropology, 2013, 150, 10-20.	2.1	18
24	Plasma Lipidomic Profile Signature of Hypertension in Mexican American Families. Hypertension, 2013, 62, 621-626.	2.7	87
25	A variant in the <i>LRRFIP1</i> gene is associated with adiposity and inflammation. Obesity, 2013, 21, 185-192.	3.0	29
26	The Positive Association of Obesity Variants with Adulthood Adiposity Strengthens over an 80-Year Period: A Gene-by-Birth Year Interaction. Human Heredity, 2013, 75, 175-185.	0.8	43
27	A genome-wide analysis of 'Bounty' descendants implicates several novel variants in migraine susceptibility. Neurogenetics, 2012, 13, 261-266.	1.4	32
28	Heritability and genome-wide linkage analysis of migraine in the genetic isolate of Norfolk Island. Gene, 2012, 494, 119-123.	2.2	18
29	An X Chromosome Association Scan of the Norfolk Island Genetic Isolate Provides Evidence for a Novel Migraine Susceptibility Locus at Xq12. PLoS ONE, 2012, 7, e37903.	2.5	11
30	A genome-wide approach accounting for body mass index identifies genetic variants influencing fasting glycemic traits and insulin resistance. Nature Genetics, 2012, 44, 659-669.	21.4	762
31	Large scale mitochondrial sequencing in Mexican Americans suggests a reappraisal of Native American origins. BMC Evolutionary Biology, 2011, 11, 293.	3.2	77
32	Variants in the human potassium channel gene (KCNN3) are associated with migraine in a high risk genetic isolate. Journal of Headache and Pain, 2011, 12, 603-608.	6.0	12
33	The Norfolk Island Eye Study (NIES): Rationale, Methodology and Distribution of Ocular Biometry (Biometry of the Bounty). Twin Research and Human Genetics, 2011, 14, 42-52.	0.6	11
34	Legacy of mutiny on the Bounty: founder effect and admixture on Norfolk Island. European Journal of Human Genetics, 2010, 18, 67-72.	2.8	26
35	European and Polynesian admixture in the Norfolk Island population. Heredity, 2010, 105, 229-234.	2.6	18
36	Principal Component and Linkage Analysis of Cardiovascular Risk Traits in the Norfolk Isolate. Human Heredity, 2009, 68, 55-64.	0.8	18

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#	Article	IF	CITATION
37	Linkage mapping of CVD risk traits in the isolated Norfolk Island population. Human Genetics, 2008, 124, 543-552.	3.8	19
38	Linkage disequilibrium analysis in the genetically isolated Norfolk Island population. Heredity, 2008, 100, 366-373.	2.6	24
39	Association between a 19Âbp deletion polymorphism at the dopamine beta-hydroxylase (DBH) locus and migraine with aura. Journal of the Neurological Sciences, 2006, 251, 118-123.	0.6	64
40	A genome-wide scan provides evidence for loci influencing a severe heritable form of common migraine. Neurogenetics, 2005, 6, 67-72.	1.4	41
41	Phenotypical Characterisation of the Isolated Norfolk Island Population Focusing on Epidemiological Indicators of Cardiovascular Disease. Human Heredity, 2005, 60, 211-219.	0.8	37
42	Investigation of the low-density lipoprotein receptor gene and cholesterol as a risk factor for migraine. Journal of the Neurological Sciences, 2004, 227, 95-100.	0.6	13
43	A molecular genetic approach for forensic animal species identification. Forensic Science International, 2003, 134, 99-108.	2.2	89
44	A Variant in the LRRFIP1 Gene Is Associated With Adiposity and Inflammation. Obesity, 0, , .	3.0	4