

Catherine Bourgain

List of Publications by Year in descending order

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Version: 2024-02-01

42
papers

1,477
citations

430874

18
h-index

315739

38
g-index

44
all docs

44
docs citations

44
times ranked

2051
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic susceptibility in practice: The regulatory trajectory of non-rare thrombophilia (NRT) genetic tests in the clinical management of venous thrombo-embolism (VTE). <i>Social Science and Medicine</i> , 2022, 304, 112903.	3.8	7
2	Les fœlicités du capital en santé. <i>Revue Française De Socio-Économie</i> , 2021, n° 26, 127-147.	0.2	1
3	Appraising screening, making risk in/visible. The medical debate over Non-Rare Thrombophilia (NRT) testing before prescribing the pill. <i>Sociology of Health and Illness</i> , 2021, 43, 1627-1642.	2.1	5
4	Imperfect biomarkers for adjuvant chemotherapy in early stage breast cancer with good prognosis. <i>Social Science and Medicine</i> , 2020, 246, 112735.	3.8	5
5	Geneva Statement on Heritable Human Genome Editing: The Need for Course Correction. <i>Trends in Biotechnology</i> , 2020, 38, 351-354.	9.3	37
6	Adopt a moratorium on heritable genome editing. <i>Nature</i> , 2019, 567, 165-168.	27.8	314
7	Searching for secondary findings: considering actionability and preserving the right not to know. <i>European Journal of Human Genetics</i> , 2019, 27, 1481-1484.	2.8	13
8	It has to become true genetics™: tumour genetics and the division of diagnostic labour in the clinic. <i>Sociology of Health and Illness</i> , 2019, 41, 643-657.	2.1	7
9	Nouveaux usages de l'ADN en matière pénale: promesses, savoirs et pratiques. <i>Cahiers Droit Sciences & Technologies</i> , 2019, , 83-93.	0.1	1
10	De la génétique clinique à la médecine génomique. <i>Cahiers Droit Sciences & Technologies</i> , 2019, , 15-29.	0.1	2
11	Iron reinforcements in Beauvais and Metz Cathedrals: from bloomery or finery? The use of logistic regression for differentiating smelting processes. <i>Journal of Archaeological Science</i> , 2014, 42, 315-333.	2.4	55
12	Quantitative damage-benefit evaluation of drug effects: major discrepancies between the general population, users and experts. <i>Journal of Psychopharmacology</i> , 2013, 27, 590-599.	4.0	11
13	Genetic and Environmental Factors Influencing the Placental Growth Factor (PGF) Variation in Two Populations. <i>PLoS ONE</i> , 2012, 7, e42537.	2.5	11
14	Association between a polymorphism in the promoter of a glutamate receptor subunit gene (<i>GRIN2A</i>) and alcoholism. <i>Addiction Biology</i> , 2012, 17, 783-785.	2.6	31
15	A damage/benefit evaluation of addictive product use. <i>Addiction</i> , 2012, 107, 441-450.	3.3	24
16	Genetics of VEGF Serum Variation in Human Isolated Populations of Cilento: Importance of VEGF Polymorphisms. <i>PLoS ONE</i> , 2011, 6, e16982.	2.5	68
17	Family History of Thyroid Cancer and the Risk of Differentiated Thyroid Cancer in French Polynesia. <i>Thyroid</i> , 2010, 20, 393-400.	4.5	26
18	A common SNP near BMP2 is associated with severity of the iron burden in HFE p.C282Y homozygous patients: A follow-up study. <i>Blood Cells, Molecules, and Diseases</i> , 2010, 44, 34-37.	1.4	32

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19	Linkage Analysis with Dense SNP Maps in Isolated Populations. <i>Human Heredity</i> , 2009, 68, 87-97.	0.8	16
20	Comprehensive Linkage and Association Analyses Identify Haplotype, Near to the TNFSF15 Gene, Significantly Associated with Spondyloarthritis. <i>PLoS Genetics</i> , 2009, 5, e1000528.	3.5	55
21	Epidemiology, Pathology, and Genetics of Histiocytic Sarcoma in the Bernese Mountain Dog Breed. <i>Journal of Heredity</i> , 2009, 100, S19-S27.	2.4	84
22	A multiple splitting approach to linkage analysis in large pedigrees identifies a linkage to asthma on chromosome 12. <i>Genetic Epidemiology</i> , 2009, 33, 207-216.	1.3	13
23	PEL: an unbiased method for estimating age-dependent genetic disease risk from pedigree data unselected for family history. <i>Genetic Epidemiology</i> , 2009, 33, 379-385.	1.3	17
24	Progressive Retinal Atrophy in the Border Collie: A new XLPR. <i>BMC Veterinary Research</i> , 2008, 4, 10.	1.9	18
25	Identification and Replication of a Novel Obesity Locus on Chromosome 1q24 in Isolated Populations of Cilento. <i>Diabetes</i> , 2008, 57, 783-790.	0.6	16
26	Facteurs acquis et g�n�tiques de modulation de la p�n�trance de lâ€™h�mochromatose HFE. <i>Bulletin De L'Academie Nationale De Medecine</i> , 2008, 192, 873-881.	0.0	0
27	Inheritance of arterial lesions in renal fibromuscular dysplasia. <i>Journal of Human Hypertension</i> , 2007, 21, 393-400.	2.2	99
28	Common Variants in the BMP2, BMP4, and HJV Genes of the Hepcidin Regulation Pathway Modulate HFE Hemochromatosis Penetrance. <i>American Journal of Human Genetics</i> , 2007, 81, 799-807.	6.2	120
29	A mixture model approach to multiple testing for the genetic analysis of gene expression. <i>BMC Proceedings</i> , 2007, 1, S141.	1.6	5
30	Power of genome-wide association studies in the presence of interacting loci. <i>Genetic Epidemiology</i> , 2007, 31, 748-762.	1.3	17
31	Are genome-wide association studies all that we need to dissect the genetic component of complex human diseases?. <i>European Journal of Human Genetics</i> , 2007, 15, 260-263.	2.8	25
32	New susceptibility locus for hypertension on chromosome 8q by efficient pedigree-breaking in an Italian isolate. <i>Human Molecular Genetics</i> , 2006, 15, 1735-1743.	2.9	39
33	Complex trait mapping in isolated populations: Are specific statistical methods required?. <i>European Journal of Human Genetics</i> , 2005, 13, 698-706.	2.8	28
34	Impact of the diagnosis definition on linkage detection. <i>BMC Genetics</i> , 2005, 6, S140.	2.7	1
35	Detection of susceptibility loci by genome-wide linkage analysis. <i>BMC Genetics</i> , 2005, 6, S18.	2.7	5
36	Modeling the effect of a genetic factor for a complex trait in a simulated population. <i>BMC Genetics</i> , 2005, 6, S87.	2.7	1

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37	Comparing strategies for association mapping in samples with related individuals. BMC Genetics, 2005, 6, S98.	2.7	9
38	Testing for Hardy-Weinberg Equilibrium in Samples With Related Individuals. Genetics, 2004, 168, 2349-2361.	2.9	59
39	Are common disease susceptibility alleles the same in outbred and founder populations?. European Journal of Human Genetics, 2004, 12, 584-590.	2.8	31
40	Novel Case-Control Test in a Founder Population Identifies P-Selectin as an Atopy-Susceptibility Locus. American Journal of Human Genetics, 2003, 73, 612-626.	6.2	148
41	Comparison of family based haplotype methods using intragenic SNPs in candidate genes. European Journal of Human Genetics, 2002, 10, 313-319.	2.8	2
42	Maximum Identity Length Contrast: A Powerful Method For Susceptibility Gene Detection in Isolated Populations. Genetic Epidemiology, 2001, 21, S560-4.	1.3	15