Anne M Bowcock

List of Publications by Year
in descending order

173	21,551	13099 68	9589 142
papers	citations	h-index	g-index

186	186	186	23024
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Kinetic Characterization of ASXL1/2-Mediated Allosteric Regulation of the BAP1 Deubiquitinase. Molecular Cancer Research, 2021, 19, 1099-1112.	3.4	1
2	Canonical and Interior Circular RNAs Function as Competing Endogenous RNAs in Psoriatic Skin. International Journal of Molecular Sciences, 2021, 22, 5182.	4.1	8
3	Abstract 81: Studying Ghanian Cancer Genomes Using Cell-free DNA. , 2021, , .		0
4	Clinicopathological features and associations in a series of South African acral melanomas. Pigment Cell and Melanoma Research, 2021, 34, 1120-1122.	3.3	2
5	CARD14â€associated papulosquamous eruption (CAPE) in pediatric patients: Three additional cases and review of the literature. Pediatric Dermatology, 2021, 38, 1237-1242.	0.9	15
6	Circulating tumor DNA is readily detectable among Ghanaian breast cancer patients supporting non-invasive cancer genomic studies in Africa. Npj Precision Oncology, 2021, 5, 83.	5.4	4
7	The tumor genetics of acral melanoma: What should a dermatologist know?. JAAD International, 2020, 1, 135-147.	2.2	18
8	Global expression and CpG methylation analysis of primary endothelial cells before and after TNFa stimulation reveals gene modules enriched in inflammatory and infectious diseases and associated DMRs. PLoS ONE, 2020, 15, e0230884.	2.5	11
9	CARD14E138A signalling in keratinocytes induces TNF-dependent skin and systemic inflammation. ELife, 2020, 9, .	6.0	16
10	CARD14-Mediated Psoriasis and Pityriasis Rubra Piliaris (PRP). , 2020, , 92-95.		0
11	Gain of function p.E138A alteration in Card14 leads to psoriasiform skin inflammation and implicates genetic modifiers in disease severity. Experimental and Molecular Pathology, 2019, 110, 104286.	2.1	13
12	A Breath of Fresh Air: Opening up the Lung Cancer Genome. Cancer Research, 2019, 79, 4808-4810.	0.9	0
13	Integrative Copy Number Analysis of Uveal Melanoma Reveals Novel Candidate Genes Involved in Tumorigenesis Including a Tumor Suppressor Role for PHF10/BAF45a. Clinical Cancer Research, 2019, 25, 5156-5166.	7.0	16
14	Familial and Somatic BAP1 Mutations Inactivate ASXL1/2-Mediated Allosteric Regulation of BAP1 Deubiquitinase by Targeting Multiple Independent Domains. Cancer Research, 2018, 78, 1200-1213.	0.9	24
15	Punctuated evolution of canonical genomic aberrations in uveal melanoma. Nature Communications, 2018, 9, 116.	12.8	144
16	CARD14-associated papulosquamous eruption: A spectrum including features of psoriasis and pityriasis rubra pilaris. Journal of the American Academy of Dermatology, 2018, 79, 487-494.	1.2	82
17	CARD14-Mediated Psoriasis and Pityriasis Rubra Piliaris (PRP). , 2018, , 1-4.		0
18	The Molecular Revolution in Cutaneous Biology: The Era of Genome-Wide Association Studies and Statistical, BigÂData, and Computational Topics. Journal of Investigative Dermatology, 2017, 137, e113-e118.	0.7	14

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19	Molecular profiling of colorectal pulmonary metastases and primary tumours: implications for targeted treatment. Oncotarget, 2017, 8, 64999-65008.	1.8	14
20	Driver Mutations in Uveal Melanoma. JAMA Ophthalmology, 2016, 134, 728.	2.5	192
21	Psoriasis mutations disrupt CARD14 autoinhibition promoting BCL10-MALT1-dependent NF-Î²B activation. Biochemical Journal, 2016, 473, 1759-1768.	3.7	62
22	Association of Melanocortin-1 Receptor Variants with Pigmentary Traits in Humans: AÂPooled Analysis from the M-Skip Project. Journal of Investigative Dermatology, 2016, 136, 1914-1917.	0.7	16
23	Management of resectable colorectal lung metastases. Clinical and Experimental Metastasis, 2016, 33, 285-296.	3.3	36
24	Epigenetic reprogramming and aberrant expression of PRAME are associated with increased metastatic risk in Class 1 and Class 2 uveal melanomas. Oncotarget, 2016, 7, 59209-59219.	1.8	94
25	Biology of advanced uveal melanoma and next steps for clinical therapeutics. Pigment Cell and Melanoma Research, 2015, 28, 135-147.	3.3	81
26	Genome-wide meta-analysis identifies multiple novel associations and ethnic heterogeneity of psoriasis susceptibility. Nature Communications, 2015, 6, 6916.	12.8	154
27	The immunogenetics of Psoriasis: A comprehensive review. Journal of Autoimmunity, 2015, 64, 66-73.	6.5	447
28	Genome-wide Association Analysis of Psoriatic Arthritis and Cutaneous Psoriasis Reveals Differences in Their Genetic Architecture. American Journal of Human Genetics, 2015, 97, 816-836.	6.2	245
29	Fine mapping of eight psoriasis susceptibility loci. European Journal of Human Genetics, 2015, 23, 844-853.	2.8	25
30	AKT Inhibitors Promote Cell Death in Cervical Cancer through Disruption of mTOR Signaling and Glucose Uptake. PLoS ONE, 2014, 9, e92948.	2.5	68
31	DNA copy number changes as diagnostic tools for lung cancer. Thorax, 2014, 69, 496-497.	5.6	21
32	Fine Mapping Major Histocompatibility Complex Associations in Psoriasis and Its Clinical Subtypes. American Journal of Human Genetics, 2014, 95, 162-172.	6.2	182
33	CARD14 Expression in Dermal Endothelial Cells in Psoriasis. PLoS ONE, 2014, 9, e111255.	2.5	52
34	Recurrent mutations at codon 625 of the splicing factor SF3B1 in uveal melanoma. Nature Genetics, 2013, 45, 133-135.	21.4	447
35	BAP1 deficiency causes loss of melanocytic cell identity in uveal melanoma. BMC Cancer, 2013, 13, 371.	2.6	123
36	Primary Ciliary Dyskinesia-Causing Mutations in Amish and Mennonite Communities. Journal of Pediatrics, 2013, 163, 383-387.	1.8	19

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37	Association of Cardiovascular and Metabolic Disease Genes with Psoriasis. Journal of Investigative Dermatology, 2013, 133, 836-839.	0.7	62
38	Noncanonical microRNAs and endogenous siRNAs in normal and psoriatic human skin. Human Molecular Genetics, 2013, 22, 737-748.	2.9	43
39	Deletion of the activating <scp>NKG</scp>2C receptor and a functional polymorphism in its ligand <scp>HLA</scp>â€Ł in psoriasis susceptibility. Experimental Dermatology, 2013, 22, 679-681.	2.9	31
40	Runx Transcription Factors Repress Human and Murine c-Myc Expression in a DNA-Binding and C-Terminally Dependent Manner. PLoS ONE, 2013, 8, e69083.	2.5	9
41	Psoriasis Patients Are Enriched for Genetic Variants That Protect against HIV-1 Disease. PLoS Genetics, 2012, 8, e1002514.	3.5	66
42	Targeting Skin: Vitiligo and Autoimmunity. Journal of Investigative Dermatology, 2012, 132, 13-15.	0.7	13
43	Protective Effect of Human Endogenous Retrovirus K dUTPase Variants on Psoriasis Susceptibility. Journal of Investigative Dermatology, 2012, 132, 1833-1840.	0.7	22
44	Histone Deacetylase Inhibitors Induce Growth Arrest and Differentiation in Uveal Melanoma. Clinical Cancer Research, 2012, 18, 408-416.	7.0	241
45	A Subset of Methylated CpG Sites Differentiate Psoriatic from Normal Skin. Journal of Investigative Dermatology, 2012, 132, 583-592.	0.7	138
46	Mutations in the Gene PRRT2 Cause Paroxysmal Kinesigenic Dyskinesia with Infantile Convulsions. Cell Reports, 2012, 1, 2-12.	6.4	250
47	Identification of 15 new psoriasis susceptibility loci highlights the role of innate immunity. Nature Genetics, 2012, 44, 1341-1348.	21.4	848
48	Genome-Wide Meta-Analysis of Psoriatic Arthritis Identifies Susceptibility Locus at REL. Journal of Investigative Dermatology, 2012, 132, 1133-1140.	0.7	99
49	PSORS2 Is Due to Mutations in CARD14. American Journal of Human Genetics, 2012, 90, 784-795.	6.2	365
50	Rare and Common Variants in CARD14, Encoding an Epidermal Regulator of NF-kappaB, in Psoriasis. American Journal of Human Genetics, 2012, 90, 796-808.	6.2	306
51	Nonlesional atopic dermatitis skin is characterized by broad terminal differentiation defects and variable immune abnormalities. Journal of Allergy and Clinical Immunology, 2011, 127, 954-964.e4.	2.9	375
52	Reversal of atopic dermatitis with narrow-band UVB phototherapy and biomarkers for therapeutic response. Journal of Allergy and Clinical Immunology, 2011, 128, 583-593.e4.	2.9	182
53	Deep sequencing of small RNAs from human skin reveals major alterations in the psoriasis miRNAome. Human Molecular Genetics, 2011, 20, 4025-4040.	2.9	213
54	A Genetic Risk Score Combining Ten Psoriasis Risk Loci Improves Disease Prediction. PLoS ONE, 2011, 6, e19454.	2.5	84

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55	Use of pharmacogenomics in psoriasis. Clinical Investigation, 2011, 1, 399-411.	0.0	4
56	Meta-Analysis Confirms the LCE3C_LCE3B Deletion as a Risk Factor for Psoriasis in Several Ethnic Groups and Finds Interaction with HLA-Cw6. Journal of Investigative Dermatology, 2011, 131, 1105-1109.	0.7	89
57	Influence of CrohnÊ¼s Disease Risk Alleles and Smoking on Disease Location. Diseases of the Colon and Rectum, 2011, 54, 1020-1025.	1.3	15
58	Primary Ciliary Dyskinesia in Amish Communities. Journal of Pediatrics, 2010, 156, 1023-1025.	1.8	13
59	Psoriasis genetics: breaking the barrier. Trends in Genetics, 2010, 26, 415-423.	6.7	203
60	Association analyses identify six new psoriasis susceptibility loci in the Chinese population. Nature Genetics, 2010, 42, 1005-1009.	21.4	287
61	Genome-wide association analysis identifies three psoriasis susceptibility loci. Nature Genetics, 2010, 42, 1000-1004.	21.4	313
62	Genome-Wide Association Studies and Infectious Disease. Critical Reviews in Immunology, 2010, 30, 305-309.	0.5	9
63	Carriers of Rare Missense Variants in IFIH1 Are Protected from Psoriasis. Journal of Investigative Dermatology, 2010, 130, 2768-2772.	0.7	65
64	Frequent Mutation of BAP1 in Metastasizing Uveal Melanomas. Science, 2010, 330, 1410-1413.	12.6	1,242
65	Multiple Loci within the Major Histocompatibility Complex Confer Risk of Psoriasis. PLoS Genetics, 2009, 5, e1000606.	3.5	141
66	Psoriasis Bench to Bedside. Archives of Dermatology, 2009, 145, 462-4.	1.4	29
67	Further Genetic Evidence for Three Psoriasis-Risk Genes: ADAM33, CDKAL1, and PTPN22. Journal of Investigative Dermatology, 2009, 129, 629-634.	0.7	67
68	Genome-wide scan reveals association of psoriasis with IL-23 and NF-Î°B pathways. Nature Genetics, 2009, 41, 199-204.	21.4	1,229
69	Deletion of the late cornified envelope LCE3B and LCE3C genes as a susceptibility factor for psoriasis. Nature Genetics, 2009, 41, 211-215.	21.4	482
70	Contribution of genetic factors for melanoma susceptibility in sporadic US melanoma patients. Experimental Dermatology, 2009, 18, 485-487.	2.9	17
71	Broad defects in epidermal cornification in atopic dermatitis identified through genomic analysis. Journal of Allergy and Clinical Immunology, 2009, 124, 1235-1244.e58.	2.9	231
72	New insights into the pathogenesis and genetics of psoriatic arthritis. Nature Clinical Practice Rheumatology, 2009, 5, 83-91.	3.2	112

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73	Genetic Linkage Localizes an Adolescent Idiopathic Scoliosis and Pectus Excavatum Gene to Chromosome 18 q. Spine, 2009, 34, E94-E100.	2.0	66
74	Asymmetric Lower-Limb Malformations in Individuals with Homeobox PITX1 Gene Mutation. American Journal of Human Genetics, 2008, 83, 616-622.	6.2	125
75	W1122 Serology Testing, Genotype, and Other Risk Factors for Repeat Ileocolic Resection in Crohn's Disease: A Time-to-Event Analysis. Gastroenterology, 2008, 134, A-637-A-638.	1.3	2
76	A Genome-Wide Association Study of Psoriasis and Psoriatic Arthritis Identifies New Disease Loci. PLoS Genetics, 2008, 4, e1000041.	3.5	572
77	Oncogenic Mutations inGNAQOccur Early in Uveal Melanoma. , 2008, 49, 5230.		329
78	Disruption of Sodium Bicarbonate Transporter SLC4A10 in a Patient With Complex Partial Epilepsy and Mental Retardation. Archives of Neurology, 2008, 65, 550.	4.5	61
79	Absence of HOXD10 Mutations in Idiopathic Clubfoot and Sporadic Vertical Talus. Clinical Orthopaedics and Related Research, 2007, 462, 27-31.	1.5	19
80	Loss of Heterozygosity of Chromosome 3 Detected with Single Nucleotide Polymorphisms Is Superior to Monosomy 3 for Predicting Metastasis in Uveal Melanoma. Clinical Cancer Research, 2007, 13, 2923-2927.	7.0	122
81	CHD7 Gene Polymorphisms Are Associated with Susceptibility to Idiopathic Scoliosis. American Journal of Human Genetics, 2007, 80, 957-965.	6.2	142
82	Two novel point mutations in the long-range SHH enhancer in three families with triphalangeal thumb and preaxial polydactyly. American Journal of Medical Genetics, Part A, 2007, 143A, 27-32.	1.2	101
83	Completing the map of human genetic variation. Nature, 2007, 447, 161-165.	27.8	178
84	Guilt by association. Nature, 2007, 447, 645-646.	27.8	40
85	Pathogenesis and therapy of psoriasis. Nature, 2007, 445, 866-873.	27.8	1,543
86	Additional evidence of a locus for complex febrile and afebrile seizures on chromosome 12q22-23.3. Neurogenetics, 2007, 8, 61-63.	1.4	7
87	Investigation of the Chromosome 17q25 PSORS2 Locus in Atopic Dermatitis. Journal of Investigative Dermatology, 2006, 126, 603-606.	0.7	16
88	HOXD10 M319K mutation in a family with isolated congenital vertical talus. Journal of Orthopaedic Research, 2006, 24, 448-453.	2.3	48
89	Peptidoglycan recognition proteins Pglyrp3 and Pglyrp4 are encoded from the epidermal differentiation complex and are candidate genes for the Psors4 locus on chromosome 1q21. Human Genetics, 2006, 119, 113-125.	3.8	45
90	Evidence for an additional locus for split hand/foot malformation in chromosome region 8q21.11â€“q22.3. American Journal of Medical Genetics, Part A, 2006, 140A, 1744-1748.	1.2	17

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91	A Regulatory Role for 1-Acylglycerol-3-phosphate-O-acyltransferase 2 in Adipocyte Differentiation. Journal of Biological Chemistry, 2006, 281, 11082-11089.	3.4	112
92	The Tetratricopeptide Repeat Domain 7 Gene is Mutated in Flaky Skin Mice: A Model for Psoriasis, Autoimmunity, and Anemia. Experimental Biology and Medicine, 2005, 230, 659-667.	2.4	26
93	Variable hand and foot abnormalities in family with congenital vertical talus and CDMP-1 gene mutation. Journal of Orthopaedic Research, 2005, 23, 1490-1494.	2.3	19
94	Understanding the Pathogenesis of Psoriasis, Psoriatic Arthritis, and Autoimmunity via a Fusion of Molecular Genetics and Immunology. Immunologic Research, 2005, 32, 045-056.	2.9	18
95	Direct genomic selection. Nature Methods, 2005, 2, 63-69.	19.0	81
96	Getting under the skin: the immunogenetics of psoriasis. Nature Reviews Immunology, 2005, 5, 699-711.	22.7	416
97	Localization of PSORS1 to a haplotype block harboring HLA-C and distinct from corneodesmosin and HCR. Human Genetics, 2005, 118, 466-476.	3.8	61
98	THE GENETICS OF PSORIASIS AND AUTOIMMUNITY. Annual Review of Genomics and Human Genetics, 2005, 6, 93-122.	6.2	117
99	Activating Killer Cell Immunoglobulin-Like Receptor Gene KIR2DS1 Is Associated With Psoriatic Arthritis. Human Immunology, 2005, 66, 836-841.	2.4	84
100	Interferon Î±-Inducible Protein 27 (IFI27) is Upregulated in Psoriatic Skin and Certain Epithelial Cancers. Journal of Investigative Dermatology, 2004, 122, 717-721.	0.7	93
101	Psoriasis Genetics: The Way Forward. Journal of Investigative Dermatology, 2004, 122, xv-xvii.	0.7	18
102	A transmission disequilibrium test for general pedigrees that is robust to the presence of random genotyping errors and any number of untyped parents. European Journal of Human Genetics, 2004, 12, 752-761.	2.8	57
103	SNTG1, the gene encoding ?1-syntrophin: a candidate gene for idiopathic scoliosis. Human Genetics, 2004, 115, 81-89.	3.8	64
104	The genetics of psoriasis, psoriatic arthritis and atopic dermatitis. Human Molecular Genetics, 2004, 13, 43R-55.	2.9	215
105	Psoriasis vulgaris: cutaneous lymphoid tissue supports T-cell activation and â€˜Type 1â€™ inflammatory gene expression. Trends in Immunology, 2004, 25, 295-305.	6.8	255
106	The Immunogenetics of Inflammatory Skin Disease. , 2004, , 55-73.		0
107	Novel mechanisms of T-cell and dendritic cell activation revealed by profiling of psoriasis on the 63,100-element oligonucleotide array. Physiological Genomics, 2003, 13, 69-78.	2.3	282
108	Novel immunoglobulin superfamily gene cluster, mapping to a region of human chromosome 17q25, linked to psoriasis susceptibility. Human Genetics, 2003, 112, 34-41.	3.8	74

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109	A putative RUNX1 binding site variant between SLC9A3R1 and NAT9 is associated with susceptibility to psoriasis. Nature Genetics, 2003, 35, 349-356.	21.4	284
110	Genetics of psoriasis: the potential impact on new therapies. Journal of the American Academy of Dermatology, 2003, 49, 51-56.	1.2	70
111	Risk Factors for Diabetes in Familial Partial Lipodystrophy, Dunnigan Variety. Diabetes Care, 2003, 26, 1350-1355.	8.6	68
112	Multisystem dystrophy syndrome due to novel missense mutations in the amino-terminal head and alpha-helical rod domains of the lamin A/C gene. American Journal of Medicine, 2002, 112, 549-555.	1.5	138
113	AGPAT2 is mutated in congenital generalized lipodystrophy linked to chromosome 9q34. Nature Genetics, 2002, 31, 21-23.	21.4	475
114	Phenotypic Heterogeneity in Patients with Familial Partial Lipodystrophy (Dunnigan Variety) Related to the Site of Missense Mutations in Lamin A/C Gene1. Journal of Clinical Endocrinology and Metabolism, 2001, 86, 59-65.	3.6	97
115	Zeroing in on tolerance. Nature Medicine, 2001, 7, 279-281.	30.7	7
116	Human Population Expansion and Microsatellite Variation. Molecular Biology and Evolution, 2000, 17, 757-767.	8.9	85
117	Localization of Susceptibility to Familial Idiopathic Scoliosis. Spine, 2000, 25, 2372-2380.	2.0	124
118	Localization of a gene for familial recurrent arthritis. Arthritis and Rheumatism, 2000, 43, 2041-2045.	6.7	70
119	Mutational and Haplotype Analyses of Families with Familial Partial Lipodystrophy (Dunnigan Variety) Reveal Recurrent Missense Mutations in the Globular C-Terminal Domain of Lamin A/C. American Journal of Human Genetics, 2000, 66, 1192-1198.	6.2	260
120	JM2, encoding a fork headâ€“related protein, is mutated in X-linked autoimmunityâ€“allergic disregulation syndrome. Journal of Clinical Investigation, 2000, 106, R75-R81.	8.2	792
121	A Gene for Congenital Generalized Lipodystrophy Maps to Human Chromosome 9q34. Journal of Clinical Endocrinology and Metabolism, 1999, 84, 3390-3394.	3.6	167
122	Genomic structure, mapping, and expression analysis of the mammalian Lunatic, Manic, and Radical fringe genes. Mammalian Genome, 1999, 10, 535-541.	2.2	24
123	In vitro transformation of cell lines from human salivary gland tumors. , 1999, 81, 793-798.		18
124	Molecular Cloning and Characterization of the HumanCLOCKGene: Expression in the Suprachiasmatic Nuclei. Genomics, 1999, 57, 189-200.	2.9	115
125	Hereditary Breast Cancer Genes. , 1999, , 199-224.		1
126	A Gene for Congenital Generalized Lipodystrophy Maps to Human Chromosome 9q34. Journal of Clinical Endocrinology and Metabolism, 1999, 84, 3390-3394.	3.6	56

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127	Conservation of function and primary structure in the BRCA1-associated RING domain (BARD1) protein. Oncogene, 1998, 17, 2143-2148.	5.9	36
128	Localization of the gene for familial partial lipodystrophy (Dunnigan variety) to chromosome 1q21â€“22. Nature Genetics, 1998, 18, 292-295.	21.4	151
129	Markers and Methods for Reconstructing Modern Human History. DNA Sequence, 1998, 8, 329-342.	0.7	0
130	The C-terminal (BRCT) Domains of BRCA1 Interact in Vivo with CtIP, a Protein Implicated in the CtBP Pathway of Transcriptional Repression. Journal of Biological Chemistry, 1998, 273, 25388-25392.	3.4	343
131	Differential Contributions ofBRCA1andBRCA2to Early-Onset Breast Cancer. New England Journal of Medicine, 1997, 336, 1416-1422.	27.0	197
132	JAK3 Maps to Human Chromosome 19p12 within a Cluster of Proto-oncogenes and Transcription Factors. Genomics, 1997, 43, 109-111.	2.9	12
133	Isolation of Two Novel WNT Genes, WNT14 and WNT15, One of Which (WNT15) Is Closely Linked to WNT3 on Human Chromosome 17q21. Genomics, 1997, 46, 450-458.	2.9	50
134	Association Mapping of Disease Loci, by Use of a Pooled DNA Genomic Screen. American Journal of Human Genetics, 1997, 61, 734-747.	6.2	228
135	Breast Cancer Genes. Breast Journal, 1997, 3, 1-6.	1.0	9
136	Infrequency of BRCA2 alterations in head and neck squamous cell carcinoma. Oncogene, 1997, 14, 2189-2193.	5.9	16
137	Association mapping with a pooled DNA genomic screen of the non-HLA genes in IDDM, RA and MS. Human Immunology, 1996, 47, 152.	2.4	0
138	Characterization ofEZH1,a Human Homolog ofDrosophila Enhancer of zestenearBRCA1. Genomics, 1996, 37, 161-171.	2.9	49
139	Identification of a RING protein that can interact in vivo with the BRCA1 gene product. Nature Genetics, 1996, 14, 430-440.	21.4	683
140	Structure and Characterization of the Human Tissue Inhibitor of Metalloproteinases-2 Gene. Journal of Biological Chemistry, 1996, 271, 25498-25505.	3.4	103
141	Genetic Locus for Psoriasis Identified. Annals of Medicine, 1995, 27, 183-186.	3.8	12
142	Direct selection of expressed sequences within a 1-Mb region flanking BRCA1 on human chromosome 17q21. Genomics, 1995, 25, 248-255.	2.9	21
143	A YAC-, P1-, and cosmid-based physical Map of the BRCA1 region on chromosome 17q21. Genomics, 1995, 25, 264-273.	2.9	15
144	The Gene for Pancreatic Polypeptide (PPY) and the Anonymous Marker D17S78 Are within 45 kb of Each Other on Chromosome 17q21. Genomics, 1994, 21, 458-460.	2.9	3

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145	Molecular cloning of BRCA1: a gene for early onset familial breast and ovarian cancer. Breast Cancer Research and Treatment, 1993, 28, 121-135.	2.5	18
146	Microsatellite Polymorphism Linkage Map of Human Chromosome 13q. Genomics, 1993, 15, 376-386.	2.9	55
147	The CEPH Consortium Linkage Map of Human Chromosome 13. Genomics, 1993, 16, 486-496.	2.9	55
148	High-Density Genetic Map of the BRCA1 Region of Chromosome 17q12-q21. Genomics, 1993, 17, 618-623.	2.9	87
149	The CA repeat marker D17S791 islocated within 40 kb of the WNT3 gene on chromosome 17q. Genomics, 1993, 18, 728-729.	2.9	2
150	A somatic cell hybrid map of human chromosome 13. Genomics, 1993, 18, 486-495.	2.9	6
151	A linkage map of human chromosome 15 with an average resolution of 2 cM and containing 55 polymorphic microsatellites. Human Molecular Genetics, 1993, 2, 2019-2030.	2.9	54
152	Evolution of Modern Humans: Evidence From Nuclear DNA Polymorphisms. , 1993, , 69-83.		17
153	Dinucleotide repeat polymorphism at the D1S167 locus. Human Molecular Genetics, 1992, 1, 138-138.	2.9	4
154	Dinucleotide repeat polymorphism at the D6S223 locus. Human Molecular Genetics, 1992, 1, 66-66.	2.9	1
155	Detection of a polymorphism within the pepsinogen C gene with PCR: Construction of a linkage map around PGC from 6p11â€“6p21.3. Genomics, 1992, 14, 398-402.	2.9	5
156	The human cationic amino acid transporter (ATRC1): Physical and genetic mapping to 13q12â€“q14. Genomics, 1992, 12, 430-434.	2.9	42
157	Linkage of Tunisian autosomal recessive Duchenneâ€“like muscular dystrophy to the pericentromeric region of chromosome 13q. Nature Genetics, 1992, 2, 315-317.	21.4	186
158	The study of variation in the human genome. Genomics, 1991, 11, 491-498.	2.9	43
159	A contiguous linkage map of chromosome 13q with 39 distinct loci separated on average by 5.1 centimorgans. Genomics, 1991, 11, 517-529.	2.9	25
160	Loss of heterozygosity for chromosome 22 DNA sequences in human meningioma. Cancer Genetics and Cytogenetics, 1991, 53, 271-277.	1.0	18
161	A Highly Informative Polymorphism of the Pepsinogen C Gene Detected by Polymerase Chain Reaction. Advances in Experimental Medicine and Biology, 1991, 306, 95-99.	1.6	3
162	Polymorphism and mapping of the IGF1 gene, and absence of association with stature among African Pygmies. Human Genetics, 1990, 85, 349-54.	3.8	17

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163	An SspI RFLP at the D13S25 locus identified by the anonymous single copy probe H2â€“42. Nucleic Acids Research, 1990, 18, 7194-7194.	14.5	4
164	An alpha satellite DNA polymorphism specific for the centromeric region of chromosome 13. Genomics, 1990, 7, 110-114.	2.9	16
165	The anonymous probe pF5A identifying the locus D13S61 detects RFLPs with XmnI and BanII. Nucleic Acids Research, 1989, 17, 8397-8397.	14.5	1
166	The anonymous DNA probe p7-26 identifying the locus [D7S17] reveals an XmnI polymorphism. Nucleic Acids Research, 1989, 17, 1787-1787.	14.5	0
167	The Î± chain of human propionyl CoA carboxylase (PCCA) (mapped to chromosome 13) detects an RFLP with XmnI. Nucleic Acids Research, 1989, 17, 8400-8400.	14.5	1
168	The anonymous probe pR1â€“4 which identifies the locus D13S59 detects a BanII RFLP. Nucleic Acids Research, 1989, 17, 8396-8396.	14.5	3
169	Rapid detection and sequencing of alleles in the 3â€² flanking region of the Interleukin-6 gene. Nucleic Acids Research, 1989, 17, 6855-6864.	14.5	83
170	Predicting genotypes at loci for autosomal recessive disorders using linked genetic markers: application to Wilson's disease. Human Genetics, 1988, 79, 109-117.	3.8	20
171	The NGF and kallikrein genes of mouse, the African rat Mastomys natalensis and man: their distribution and mode of expression in the salivary gland. Molecular Brain Research, 1988, 3, 165-172.	2.3	11
172	The haemoglobin H disease mental retardation syndrome: molecular studies on the South African case. British Journal of Haematology, 1984, 56, 69-78.	2.5	16
173	Hereditary Breast Cancer Genes. , 0, , 199-224.		0