## Conversion of diploidy to haploidy

Nature 403, 723-724 DOI: 10.1038/35001659

**Citation Report** 

#	Article	IF	CITATIONS
1	Genetics testing for colon cancer: Joint statement of the American College of Medical Genetics and American Society of Human Genetics: Joint Test and Technology Transfer Committee Working Group. Genetics in Medicine, 2000, 2, 362-366.	1.1	36
3	Sequence variability and candidate gene analysis in complex disease: association of micro opioid receptor gene variation with substance dependence. Human Molecular Genetics, 2000, 9, 2895-2908.	1.4	276
4	Recurrent germline mutation in MSH2 arises frequently de novo. Journal of Medical Genetics, 2000, 37, 646-652.	1.5	72
5	Screening for Genomic Rearrangements in Families with Breast and Ovarian Cancer Identifies BRCA1 Mutations Previously Missed by Conformation-Sensitive Gel Electrophoresis or Sequencing. American Journal of Human Genetics, 2000, 67, 841-850.	2.6	149
6	Genetics of hereditary colon cancer— a basis for prevention?. European Journal of Cancer, 2000, 36, 1215-1223.	1.3	9
7	GENETICS: Genetic Testing- Present and Future. Science, 2000, 289, 1890-1892.	6.0	51
8	Efficient detection of hereditary nonpolyposis colorectal cancer gene carriers by screening for tumor microsatellite instability before germline genetic testing. Gastroenterology, 2001, 120, 21-30.	0.6	149
9	Germline mutations of EXO1 gene in patients with hereditary nonpolyposis colorectal cancer (HNPCC) and atypical HNPCC forms. Gastroenterology, 2001, 120, 1580-1587.	0.6	138
10	AGA technical review on hereditary colorectal cancer and genetic testing. Gastroenterology, 2001, 121, 198-213.	0.6	318
11	The colon cancer burden of genetically defined hereditary nonpolyposis colon cancer. Gastroenterology, 2001, 121, 830-838.	0.6	236
12	Gastric Cancer and H. pylori: Host Genetics Open the Way. Gastroenterology, 2001, 121, 1002-1012.	0.6	50
13	HNPCC: An uncommon but important diagnosis. Gastroenterology, 2001, 121, 1005-1008.	0.6	17
14	Deficient DNA mismatch repair: a common etiologic factor for colon cancer. Human Molecular Genetics, 2001, 10, 735-740.	1.4	429
15	STK11/LKB1 Peutz-Jeghers Gene Inactivation in Intraductal Papillary-Mucinous Neoplasms of the Pancreas. American Journal of Pathology, 2001, 159, 2017-2022.	1.9	251
16	PTEN mutations and Proteus syndrome. Lancet, The, 2001, 358, 2079-2080.	6.3	59
17	Adhesion molecules in lymphocyte trafficking and colitis. Gastroenterology, 2001, 121, 1008-1010.	0.6	4
18	Identification and Characterization of Disease-Related Genes: Focus on Endocrine Neoplasias. , 2001, 28, 20-49.		0
19	Coincidence, coevolution, or causation? DNA content, cell size, and the C-value enigma. Biological Reviews, 2001, 76, 65-101.	4.7	590

#	Article	IF	CITATIONS
20	Molecular Screening for Hereditary Nonpolyposis Colorectal Cancer: A Prospective, Population-Based Study. Journal of Clinical Oncology, 2001, 19, 3944-3950.	0.8	101
21	BRCA1/2 Testing: Complex Themes in Result Interpretation. Journal of Clinical Oncology, 2001, 19, 2555-2565.	0.8	59
22	Hybrids monosomal for human chromosome 5 reveal the presence of a spinal muscular atrophy (SMA) carrier with two SMN1 copies on one chromosome. Human Genetics, 2001, 108, 109-115.	1.8	38
23	Methylation pattern of different regions of theMLH1 promoter and silencing of gene expression in hereditary and sporadic colorectal cancer. Genes Chromosomes and Cancer, 2001, 31, 357-361.	1.5	53
24	Mutations of the 'minor' mismatch repair gene MSH6 in typical and atypical hereditary nonpolyposis colorectal cancer. Familial Cancer, 2001, 1, 95-101.	0.9	24
25	Clinical and biologic heterogeneity of hereditary nonpolyposis colorectal cancer. International Journal of Cancer, 2001, 95, 323-328.	2.3	19
26	Experimentally-derived haplotypes substantially increase the efficiency of linkage disequilibrium studies. Nature Genetics, 2001, 28, 361-364.	9.4	150
27	At the interfaces of epidemiology, genetics and genomics. Nature Reviews Genetics, 2001, 2, 142-147.	7.7	68
28	Genetic susceptibility for breast cancer?Risk assessment and counseling. Seminars in Oncology, 2001, 28, 419-433.	0.8	25
29	Facilitating haplotype analysis by fully automated analysis of all chromosomes in human-mouse hybrid cell lines. Cytogenetic and Genome Research, 2001, 93, 11-15.	0.6	10
30	Mutations in cis can confound genotype-phenotype correlations in hypertrophic cardiomyopathy. Journal of Medical Genetics, 2001, 38, 385-388.	1.5	32
31	Allelic association and disease mapping. Briefings in Bioinformatics, 2001, 2, 375-387.	3.2	7
32	Associating Genes to Drug Response. Drug Information Journal, 2002, 36, 751-761.	0.5	2
33	Haploinsufficiency of Flap endonuclease (Fen1) leads to rapid tumor progression. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 9924-9929.	3.3	227
34	Exploring and Exploiting Instability. Cancer Biology and Therapy, 2002, 1, 212-225.	1.5	14
35	Characterisation of the growth regulating gene IMP3, a candidate for Silver-Russell syndrome. Journal of Medical Genetics, 2002, 39, 575-581.	1.5	49
36	Integrating pharmacogenomics into drug development. Pharmacogenomics, 2002, 3, 453-467.	0.6	32
37	Haplotype-Specific Effects on Endothelial NO Synthase Promoter Efficiency. Arteriosclerosis, Thrombosis, and Vascular Biology, 2002, 22, e1-4.	1.1	129

#	Article	IF	CITATIONS
38	Mismatch Repair Genes hMLH1 and hMSH2 and Colorectal Cancer: A HuGE Review. American Journal of Epidemiology, 2002, 156, 885-902.	1.6	128
39	Determination of a Novel Haplotype of β2-adrenergic Receptor in the Japanese Population by the Combination of the Electronic Microchip Assay Using the NanoChip System with Allele-specific PCR. Drug Metabolism and Pharmacokinetics, 2002, 17, 532-539.	1.1	2
40	Turning SNPs into Useful Markers of Drug Response. , 0, , 35-55.		1
41	GENETIC ANDEPIGENETICALTERATIONS INCOLONCANCER. Annual Review of Genomics and Human Genetics, 2002, 3, 101-128.	2.5	261
42	Functional analysis of hMLH1 variants and HNPCC-related mutations using a human expression system. Gastroenterology, 2002, 122, 211-219.	0.6	179
43	Hereditary nonpolyposis colorectal cancer in young colorectal cancer patients: High-risk clinic versus population-based registry. Gastroenterology, 2002, 122, 940-947.	0.6	29
44	Score Tests for Association between Traits and Haplotypes when Linkage Phase Is Ambiguous. American Journal of Human Genetics, 2002, 70, 425-434.	2.6	1,656
45	Focus on colon cancer. Cancer Cell, 2002, 1, 233-236.	7.7	127
46	Determination of ?2-adrenergic receptor (ADRB2) haplotypes by a multiplexed polymerase chain reaction assay. Human Mutation, 2002, 20, 479-479.	1.1	34
47	A 10-Mb paracentric inversion of chromosome arm 2p inactivatesMSH2 and is responsible for hereditary nonpolyposis colorectal cancer in a North-American kindred. Genes Chromosomes and Cancer, 2002, 35, 49-57.	1.5	57
48	Identification and functional characterization of the promoter region of the humanMSH6 gene. Genes Chromosomes and Cancer, 2002, 33, 36-46.	1.5	20
49	Genetic testing and risk assessment for spinal muscular atrophy (SMA). Human Genetics, 2002, 111, 477-500.	1.8	167
50	Genomic rearrangements of EYA1 account for a large fraction of families with BOR syndrome. European Journal of Human Genetics, 2002, 10, 757-766.	1.4	45
51	An endonuclease/ligase based mutation scanning method especially suited for analysis of neoplastic tissue. Oncogene, 2002, 21, 1909-1921.	2.6	40
52	Small changes in expression affect predisposition to tumorigenesis. Nature Genetics, 2002, 30, 25-26.	9.4	234
53	Chromosome 7p disruptions in Silver Russell syndrome: delineating an imprinted candidate gene region Human Genetics, 2002, 111, 376-387.	1.8	79
54	Correlations between Phenotype and Microsatellite Instability in HNPCC: Implications for Genetic Testing. Familial Cancer, 2002, 3, 117-121.	0.9	8
55	Genetic mutation associated with meiotic metaphase-specific apoptosis in MRL/MpJ mice. Molecular Reproduction and Development, 2003, 64, 179-188.	1.0	17

#	Article	IF	CITATIONS
56	Detection and assignment ofTP53 mutations in tumor DNA using peptide mass signature genotyping. Human Mutation, 2003, 22, 158-165.	1.1	14
57	Identification and characterization of genomic rearrangements ofMSH2 andMLH1 in Lynch syndrome (HNPCC) by novel techniques. Human Mutation, 2003, 22, 258-258.	1.1	87
58	Clinical and molecular delineation of the Greig cephalopolysyndactyly contiguous gene deletion syndrome and its distinction from acrocallosal syndrome. American Journal of Medical Genetics Part A, 2003, 123A, 236-242.	2.4	50
59	Hereditary nonpolyposis colorectal cancer and related conditions. American Journal of Medical Genetics Part A, 2003, 122A, 325-334.	2.4	70
60	Identification of a deletion in the mismatch repair gene, MSH2, using mouse-human cell hybrids monosomal for chromosome 2. Clinical Genetics, 2003, 63, 215-218.	1.0	4
61	Germline hMLH1 promoter mutation in a Newfoundland HNPCC kindred. Clinical Genetics, 2003, 64, 220-227.	1.0	20
62	Increased expression and no mutation of the Flap endonuclease (FEN1) gene in human lung cancer. Oncogene, 2003, 22, 7243-7246.	2.6	64
63	Genetic testing for high-risk colon cancer patients1 1Abbreviations used in this paper: FAP, familial adenomatous polyposis; HMPS, hereditary mixed polyposis syndrome; HNPCC, hereditary nonpolyposis colon cancer; JPS, juvenile polyposis; MMR, mutation mismatch repair; MSI, microsatellite instability; PIS. Peutz-leghers syndrome: TGF, transforming growth factor., Gastroenterology, 2003, 124, 1574-1594.	0.6	194
64	Diagnosis and Management of Hereditary Non-Polyposis Colon Cancer. Gastrointestinal Endoscopy, 2003, 58, 390-408.	0.5	28
65	Hereditary Colorectal Cancer. New England Journal of Medicine, 2003, 348, 919-932.	13.0	1,870
		10.7	
66	Hereditary nonpolyposis colorectal cancer: preventive management. Cancer Treatment Reviews, 2003, 29, 461-470.	3.4	53
66 67	Hereditary nonpolyposis colorectal cancer: preventive management. Cancer Treatment Reviews, 2003, 29, 461-470. Sensitive and Efficient Detection of RB1 Gene Mutations Enhances Care for Families with Retinoblastoma. American Journal of Human Genetics, 2003, 72, 253-269.	3.4 2.6	53 285
66 67 68	<ul> <li>Hereditary nonpolyposis colorectal cancer: preventive management. Cancer Treatment Reviews, 2003, 29, 461-470.</li> <li>Sensitive and Efficient Detection of RB1 Gene Mutations Enhances Care for Families with Retinoblastoma. American Journal of Human Genetics, 2003, 72, 253-269.</li> <li>Inherited Human Diseases: Victories, Challenges, Disappointments**Previously presented at the annual meeting of The American Society of Human Genetics, in Baltimore, on October 18, 2002 American Journal of Human Genetics, 2003, 72, 236-240.</li> </ul>	3.4 2.6 2.6	53 285 7
66 67 68 69	<ul> <li>Hereditary nonpolyposis colorectal cancer: preventive management. Cancer Treatment Reviews, 2003, 29, 461-470.</li> <li>Sensitive and Efficient Detection of RB1 Gene Mutations Enhances Care for Families with Retinoblastoma. American Journal of Human Genetics, 2003, 72, 253-269.</li> <li>Inherited Human Diseases: Victories, Challenges, Disappointments**Previously presented at the annual meeting of The American Society of Human Genetics, in Baltimore, on October 18, 2002 American Journal of Human Genetics, 2003, 72, 236-240.</li> <li>Molecular Analysis of Hereditary Nonpolyposis Colorectal Cancer in the United States: High Mutation Detection Rate among Clinically Selected Families and Characterization of an American Founder Genomic Deletion of the MSH2 Gene. American Journal of Human Genetics, 2003, 72, 1088-1100.</li> </ul>	3.4 2.6 2.6 2.6	53 285 7 195
66 67 68 69 70	<ul> <li>Hereditary nonpolyposis colorectal cancer: preventive management. Cancer Treatment Reviews, 2003, 29, 461-470.</li> <li>Sensitive and Efficient Detection of RB1 Gene Mutations Enhances Care for Families with Retinoblastoma. American Journal of Human Genetics, 2003, 72, 253-269.</li> <li>Inherited Human Diseases: Victories, Challenges, Disappointments**Previously presented at the annual meeting of The American Society of Human Genetics, in Baltimore, on October 18, 2002 American Journal of Human Genetics, 2003, 72, 236-240.</li> <li>Molecular Analysis of Hereditary Nonpolyposis Colorectal Cancer in the United States: High Mutation Detection Rate among Clinically Selected Families and Characterization of an American Founder Genomic Deletion of the MSH2 Gene. American Journal of Human Genetics, 2003, 72, 1088-1100.</li> <li>Confirmation of the Type 2 Myotonic Dystrophy (CCTG) Expansion Mutation in Patients with Proximal Myotonic Myopathy/Proximal Myotonic Dystrophy of Different European Origins: A Single Shared Haplotype Indicates an Ancestral Founder Effect. American Journal of Human Genetics, 2003, 73, 835-848.</li> </ul>	3.4 2.6 2.6 2.6 2.6	53 285 7 195 132
<ul> <li>66</li> <li>67</li> <li>68</li> <li>69</li> <li>70</li> <li>71</li> </ul>	<ul> <li>Hereditary nonpolyposis colorectal cancer: preventive management. Cancer Treatment Reviews, 2003, 29, 461-470.</li> <li>Sensitive and Efficient Detection of RB1 Gene Mutations Enhances Care for Families with Retinoblastoma. American Journal of Human Genetics, 2003, 72, 253-269.</li> <li>Inherited Human Diseases: Victories, Challenges, Disappointments**Previously presented at the annual meeting of The American Society of Human Genetics, in Baltimore, on October 18, 2002 American Journal of Human Genetics, 2003, 72, 236-240.</li> <li>Molecular Analysis of Hereditary Nonpolyposis Colorectal Cancer in the United States: High Mutation Detection Rate among Clinically Selected Families and Characterization of an American Founder Genomic Deletion of the MSH2 Gene. American Journal of Human Genetics, 2003, 72, 1088-1100.</li> <li>Confirmation of the Type 2 Myotonic Dystrophy (CCTG) Expansion Mutation in Patients with Proximal Myotonic Myopathyl/Proximal Myotonic Dystrophy of Different European Origins: A Single Shared Haplotype Indicates an Ancestral Founder Effect. American Journal of Human Genetics, 2003, 73, 835-848.</li> <li>Altered Expression of MLH1, MSH2, and MSH6 in Predisposition to Hereditary Nonpolyposis Colorectal Cancer. Journal of Clinical Oncology, 2003, 21, 3629-3637.</li> </ul>	3.4 2.6 2.6 2.6 2.6 0.8	53 285 7 195 132 88
<ul> <li>66</li> <li>67</li> <li>68</li> <li>69</li> <li>70</li> <li>71</li> <li>73</li> </ul>	<ul> <li>Hereditary nonpolyposis colorectal cancer: preventive management. Cancer Treatment Reviews, 2003, 29, 461-470.</li> <li>Sensitive and Efficient Detection of RB1 Gene Mutations Enhances Care for Families with Retinoblastoma. American Journal of Human Cenetics, 2003, 72, 253-269.</li> <li>Inherited Human Diseases: Victories, Challenges, Disappointments**Previously presented at the annual meeting of The American Society of Human Genetics, in Baltimore, on October 18, 2002 American Journal of Human Genetics, 2003, 72, 253-269.</li> <li>Molecular Analysis of Hereditary Nonpolyposis Colorectal Cancer in the United States: High Mutation Detection Rate among Clinically Selected Families and Characterization of an American Founder Genomic Deletion of the MSH2 Gene. American Journal of Human Genetics, 2003, 72, 1088-1100.</li> <li>Confirmation of the Type 2 Myotonic Dystrophy (CCTG) Expansion Mutation in Patients with Proximal Myotonic Myopathy/Proximal Myotonic Dystrophy of Different European Origins: A Single Shared Haplotype Indicates an Ancestral Founder Effect. American Journal of Human Genetics, 2003, 73, 835-848.</li> <li>Altered Expression of MLH1, MSH2, and MSH6 in Predisposition to Hereditary Nonpolyposis Colorectal Cancer. Journal of Clinical Oncology, 2003, 21, 3629-3637.</li> <li>Multiple Primary Cancer, Including Transitional Cell Carcinoma of the Upper Uroepithelial Tract in a Multigeneration Hipcc Family: Molecular Genetic, Diagnostic, and Management Implications. American Journal of Gastroenterology, 2003, 98, 664-670.</li> </ul>	3.4 2.6 2.6 2.6 2.6 2.6 0.8 0.2	<ul> <li>53</li> <li>285</li> <li>7</li> <li>195</li> <li>132</li> <li>88</li> <li>15</li> </ul>

#	Article	IF	CITATIONS
76	Analysis of the ARMD1 locus: evidence that a mutation in HEMICENTIN-1 is associated with age-related macular degeneration in a large family. Human Molecular Genetics, 2003, 12, 3315-3323.	1.4	175
77	Conversion Technology and Cancer Predispositions. , 2003, 223, 415-424.		0
78	Hereditary Colon Cancer Genes. , 2003, 222, 059-083.		4
79	Conversion technology and its role in genetic testing of inherited diseases. Expert Review of Molecular Diagnostics, 2003, 3, 497-506.	1.5	4
80	Hereditary Breast Cancer Considering Cowden Syndrome. Cancer Nursing, 2003, 26, 370-375.	0.7	4
81	The Hereditary Nonpolyposis Colorectal Cancer Syndrome: Genetics and Clinical Implications. Annals of Internal Medicine, 2003, 138, 560.	2.0	256
82	HNPCC (Lynch Syndrome): Differential Diagnosis, Molecular Genetics and Management - a Review. Hereditary Cancer in Clinical Practice, 2003, 1, 7.	0.6	19
83	Frequent genomic disorganisation of MLH1 in hereditary non-polyposis colorectal cancer (HNPCC) screened by RT-PCR on puromycin treated samples. Journal of Medical Genetics, 2003, 40, 30e-30.	1.5	9
84	Robust Dosage-PCR for Detection of Heterozygous Chromosomal Deletions. BioTechniques, 2003, 34, 558-570.	0.8	15
85	The Genetics of Hereditary Non-Polyposis Colorectal Cancer. Journal of the National Comprehensive Cancer Network: JNCCN, 2003, 1, 137-144.	2.3	11
86	Lynch Syndrome: History and Current Status. Disease Markers, 2004, 20, 181-198.	0.6	49
88	Clinical genetic counselling for familial cancers requires reliable data on familial cancer risks and general action plans. Journal of Medical Genetics, 2004, 41, 801-807.	1.5	45
89	An Alu-mediated partial SDHC deletion causes familial and sporadic paraganglioma. Journal of Medical Genetics, 2004, 41, 703-709.	1.5	74
90	Mismatch Repair Gene PMS2. Cancer Research, 2004, 64, 4721-4727.	0.4	149
91	Spinal muscular atrophy: molecular genetics and diagnostics. Expert Review of Molecular Diagnostics, 2004, 4, 15-29.	1.5	124
92	Genetic aetiology of diffuse gastric cancer: so near, yet so far. Journal of Medical Genetics, 2004, 41, 481-483.	1.5	1
93	A Founder Mutation of the <emph type="ITAL">MSH2</emph> Gene and Hereditary Nonpolyposis Colorectal Cancer in the United States. JAMA - Journal of the American Medical Association, 2004, 291, 718.	3.8	75
94	A NOTCH4 association with multiple sclerosis is secondary to HLA-DR*1501. Tissue Antigens, 2004, 63, 13-20.	1.0	25

#	Article	IF	CITATIONS
95	Novel germline hMSH2 genomic deletion and somatic hMSH2 mutations in a hereditary nonpolyposis colorectal cancer family. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2004, 548, 19-25.	0.4	4
96	Genomic instability and colon cancer. Cancer and Metastasis Reviews, 2004, 23, 11-27.	2.7	280
97	The utility of immunohistochemical detection of DNA mismatch repair gene proteins. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2004, 445, 431-441.	1.4	96
98	Fine mapping of the Schnyder?s crystalline corneal dystrophy locus. Human Genetics, 2004, 114, 594-600.	1.8	21
99	Mapping genomic deletions down to the base: a quantitative copy number scanning approach used to characterise and clone the breakpoints of a recurrent 7p14.2p15.3 deletion. Human Genetics, 2004, 115, 459-467.	1.8	26
100	Cancer in Jews: introduction and overview. Familial Cancer, 2004, 3, 177-192.	0.9	24
101	A new algorithm for haplotype-based association analysis: the Stochastic-EM algorithm. Annals of Human Genetics, 2004, 68, 165-177.	0.3	258
102	Inherited predisposition to cancer: A historical overview. American Journal of Medical Genetics Part A, 2004, 129C, 5-22.	2.4	44
103	Power of direct vs. indirect haplotyping in association studies. Genetic Epidemiology, 2004, 26, 116-124.	0.6	13
104	Haplotype Block Partitioning and Tag SNP Selection Using Genotype Data and Their Applications to Association Studies. Genome Research, 2004, 14, 908-916.	2.4	143
106	Long-Range (17.7 kb) Allele-Specific Polymerase Chain Reaction Method for Direct Haplotyping of R117H and IVS-8 Mutations of the Cystic Fibrosis Transmembrane Regulator Gene. Journal of Molecular Diagnostics, 2004, 6, 264-270.	1.2	17
107	Incorporating Genotyping Uncertainty in Haplotype Inference for Single-Nucleotide Polymorphisms. American Journal of Human Genetics, 2004, 74, 495-510.	2.6	45
108	DNA/RNA Helicase Gene Mutations in a Form of Juvenile Amyotrophic Lateral Sclerosis (ALS4). American Journal of Human Genetics, 2004, 74, 1128-1135.	2.6	717
109	An Alu-mediated 31.5-kb deletion as the cause of factor XI deficiency in 2 unrelated patients. Blood, 2004, 104, 2394-2396.	0.6	12
110	Direct molecular haplotyping by melting curve analysis of hybridization probes: beta 2-adrenergic receptor haplotypes as an example. Nucleic Acids Research, 2005, 33, e89-e89.	6.5	30
111	Value of Immunohistochemical Detection of DNA Mismatch Repair Proteins in Predicting Germline Mutation in Hereditary Colorectal Neoplasms. American Journal of Surgical Pathology, 2005, 29, 96-104.	2.1	136
112	Morphogenetic investigation of metaphase-specific cell death in meiotic spermatocytes in mice. Kaibogaku Zasshi Journal of Anatomy, 2005, 80, 141-152.	1.2	4
114	Hereditary Colorectal Cancer-Part II. Current Problems in Surgery, 2005, 42, 267-333.	0.6	31

#	Article	IF	CITATIONS
115	Molecular characterization of the spectrum of genomic deletions in the mismatch repair genesMSH2,MLH1,MSH6, andPMS2 responsible for hereditary nonpolyposis colorectal cancer (HNPCC). Genes Chromosomes and Cancer, 2005, 44, 123-138.	1.5	112
116	Linear allele-specific long-range amplification: a novel method of long-range molecular haplotyping. Human Mutation, 2005, 26, 393-394.	1.1	10
117	Spectrum and frequencies of mutations inMSH2 andMLH1 identified in 1,721 German families suspected of hereditary nonpolyposis colorectal cancer. International Journal of Cancer, 2005, 116, 692-702.	2.3	113
118	Molecular pathogenesis of colorectal cancer. Cancer, 2005, 104, 2035-2047.	2.0	138
119	Use of Microsatellite Instability and Immunohistochemistry Testing for the Identification of Individuals at Risk for Lynch Syndrome. Familial Cancer, 2005, 4, 255-265.	0.9	109
120	Evolution of the Nomenclature for the Hereditary Colorectal Cancer Syndromes. Familial Cancer, 2005, 4, 211-218.	0.9	118
122	Approach to common chronic disorders of adulthood. , 2005, , .		0
123	Progress in Genetic Testing, Classification, and Identification of Lynch Syndrome. JAMA - Journal of the American Medical Association, 2005, 293, 2028.	3.8	50
124	Direct Molecular Haplotyping of the IVS-8 Poly(TG) and PolyT Repeat Tracts in the Cystic Fibrosis Gene by Melting Curve Analysis of Hybridization Probes. Clinical Chemistry, 2005, 51, 1619-1623.	1.5	15
125	Accuracy of MSI testing in predicting germline mutations of MSH2 and MLH1: a case study in Bayesian meta-analysis of diagnostic tests without a gold standard. Biostatistics, 2005, 6, 450-464.	0.9	15
126	History and Molecular Genetics of Lynch Syndrome in Family G. JAMA - Journal of the American Medical Association, 2005, 294, 2195.	3.8	70
127	Conversion Analysis for Mutation Detection in <emph>MLH1</emph> and <emph>MSH2</emph> in Patients With Colorectal Cancer. JAMA - Journal of the American Medical Association, 2005, 293, 799.	3.8	93
128	Shotgun haplotyping: a novel method for surveying allelic sequence variation. Nucleic Acids Research, 2005, 33, e152-e152.	6.5	7
129	Identification of disease genes by whole genome CGH arrays. Human Molecular Genetics, 2005, 14, R215-R223.	1.4	140
130	Comprehensive analysis of CDKN2A (p16INK4A/p14ARF) and CDKN2B genes in 53 melanoma index cases considered to be at heightened risk of melanoma. Journal of Medical Genetics, 2005, 43, 39-47.	1.5	50
131	Inherited Susceptibility to Colorectal Cancer. Annual Review of Medicine, 2005, 56, 539-554.	5.0	80
132	Immunohistochemical Analysis Reveals High Frequency of PMS2 Defects in Colorectal Cancer. Gastroenterology, 2005, 128, 1160-1171.	0.6	166
133	Characterization of hMLH1 and hMSH2 Gene Dosage Alterations in Lynch Syndrome Patients. Gastroenterology, 2005, 129, 846-854.	0.6	49

		CITATION R	REPORT	
#	Article		IF	CITATIONS
134	Definition and Clinical Importance of Haplotypes. Annual Review of Medicine, 2005, 56	5, 303-320.	5.0	283
135	New Developments in Lynch Syndrome (Hereditary Nonpolyposis Colorectal Cancer) a Repair Gene Testing. Gastroenterology, 2006, 130, 577-587.	nd Mismatch	0.6	58
137	DNA mismatch repair and Lynch syndrome. Journal of Molecular Histology, 2006, 37, 2	71-283.	1.0	17
138	Surveillance Colonoscopy in Individuals at Risk for Hereditary Nonpolyposis Colorectal Evidence-Based Review. Diseases of the Colon and Rectum, 2006, 49, 80-95.	Cancer: An	0.7	20
139	Direct molecular haplotyping of multiple polymorphisms within exon 4 of the human catechol-O-methyltransferase gene by liquid chromatography–electrospray ionizatio mass spectrometry. Analytical and Bioanalytical Chemistry, 2006, 386, 83-91.	n time-of-flight	1.9	12
140	The genetics of HNPCC: Application to diagnosis and screening. Critical Reviews in Oncology/Hematology, 2006, 58, 208-220.		2.0	89
141	A likelihood-based method for haplotype association studies of case-control data with uncertainty. Science in China Series A: Mathematics, 2006, 49, 130-144.	genotyping	0.5	2
142	HaploRec: efficient and accurate large-scale reconstruction of haplotypes. BMC Bioinfo 7, 542.	rmatics, 2006,	1.2	42
143	A comparison of several methods for haplotype frequency estimation and haplotype re for tightly linked markers from general pedigrees. Genetic Epidemiology, 2006, 30, 423	construction 3-437.	0.6	24
144	Long-range PCR facilitates the identification of PMS2-specific mutations. Human Mutat 490-495.	tion, 2006, 27,	1.1	90
145	Prediction of Germline Mutations and Cancer Risk in the Lynch Syndrome. JAMA - Jourr American Medical Association, 2006, 296, 1479.	al of the	3.8	328
146	DM2 intronic expansions: evidence for CCUG accumulation without flanking sequence ZNF9 mRNA processing or protein expression. Human Molecular Genetics, 2006, 15, 1	or effects on 808-1815.	1.4	99
147	Haplotype inference for present–absent genotype data using previously identified ha haplotype patterns. Bioinformatics, 2007, 23, 2399-2406.	aplotypes and	1.8	23
148	An efficient method for multi-locus molecular haplotyping. Nucleic Acids Research, 200	)7, 35, e6-e6.	6.5	17
149	Incorporating Genotyping Uncertainty in Haplotype Frequency Estimation in Pedigree Heredity, 2007, 64, 172-181.	Studies. Human	0.4	8
150	Human SULT1A1 gene: copy number differences and functional implications. Human N Genetics, 2007, 16, 463-470.	1olecular	1.4	102
151	MALDI-MS of Nucleic Acids and Practical Implementations in Genomics and Genetics. ,	0, , 131-179.		3
152	DLX5 and DLX6 Expression Is Biallelic and Not Modulated by MeCP2 Deficiency. Ameri Human Genetics, 2007, 81, 492-506.	can Journal of	2.6	48

#	Article	IF	CITATIONS
153	Hereditary Nonpolyposis Colorectal Cancer. , 2007, , 223-232.		2
154	Molecular Pathology in Clinical Practice. , 2007, , .		8
155	Corticotropin releasing hormone ( <i>CRH</i> ) gene variation: Comprehensive resequencing for variant and molecular haplotype discovery in monosomic hybrid cell lines. DNA Sequence, 2007, 18, 434-444.	0.7	17
157	Understanding the accuracy of statistical haplotype inference with sequence data of known phase. Genetic Epidemiology, 2007, 31, 659-671.	0.6	64
158	Direct haplotyping of bi-allelic SNPs using ARMS and RFLP analysis techniques. New Biotechnology, 2007, 24, 609-612.	2.7	7
159	Rapid and reliable genotyping of polymorphic loci modifying correct splicing of CFTR pre-mRNA using mass spectrometry. European Journal of Human Genetics, 2007, 15, 53-61.	1.4	1
160	Coincidence, coevolution, or causation? DNA content, cellsize, and the Câ€value enigma. Biological Reviews, 2001, 76, 65-101.	4.7	67
161	Total Parenteral Nutrition Leads to Alteration of Hepatocyte Cell Cycle Gene Expression and Proliferation in the Mouse. Digestive Diseases and Sciences, 2007, 52, 920-930.	1.1	4
162	Germline missense mutations in mismatch-repair genes and genetic testing for HNPCC. Current Colorectal Cancer Reports, 2007, 3, 191-198.	1.0	0
163	Accurate classification of <i>MLH1/MSH2</i> missense variants with multivariate analysis of protein polymorphisms-mismatch repair (MAPP-MMR). Human Mutation, 2008, 29, 852-860.	1.1	101
164	Myofibrillar myopathy with arrhythmogenic right ventricular cardiomyopathy 7: corroboration and narrowing of the critical region on 10q22.3. European Journal of Human Genetics, 2008, 16, 367-373.	1.4	18
165	Clinical characterization and genetic mapping of North Carolina macular dystrophy. Vision Research, 2008, 48, 470-477.	0.7	28
166	Evaluation of two methods for computational HLA haplotypes inference using a real dataset. BMC Bioinformatics, 2008, 9, 68.	1.2	13
167	Haplotypeâ€Association Analysis. Advances in Genetics, 2008, 60, 335-405.	0.8	116
169	Origins and Prevalence of the American Founder Mutation of <i>MSH2</i> . Cancer Research, 2008, 68, 2145-2153.	0.4	34
170	SNP-specific extraction of haplotype-resolved targeted genomic regions. Nucleic Acids Research, 2008, 36, e94-e94.	6.5	25
171	Distinct effects of the recurrent Mlh1 <sup>G67R</sup> mutation on MMR functions, cancer, and meiosis. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 4247-4252.	3.3	39
172	A frame-shift mutation of PMS2 is a widespread cause of Lynch syndrome. Journal of Medical Genetics, 2008, 45, 340-345.	1.5	47

#	Article	IF	CITATIONS
173	A Novel Duplication Confirms the Involvement of 5q23.2 in Autosomal Dominant Leukodystrophy. Archives of Neurology, 2008, 65, 1496.	4.9	32
175	hnRNP H enhances skipping of a nonfunctional exon P3A in CHRNA1 and a mutation disrupting its binding causes congenital myasthenic syndrome. Human Molecular Genetics, 2008, 17, 4022-4035.	1.4	54
176	Glucocorticoid Receptor Gene Variant in the 3′ Untranslated Region Is Associated with Multiple Measures of Blood Pressure. Journal of Clinical Endocrinology and Metabolism, 2009, 94, 268-276.	1.8	22
177	Is it Williams syndrome? <i>GTF2IRD1</i> implicated in visual–spatial construction and <i>GTF2I</i> in sociability revealed by high resolution arrays. American Journal of Medical Genetics, Part A, 2009, 149A, 302-314.	0.7	100
178	Fine mapping and association studies in a candidate region for autism on chromosome 2q31–q32. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2009, 150B, 535-544.	1.1	12
179	Large genomic rearrangements and germline epimutations in Lynch syndrome. International Journal of Cancer, 2009, 124, 2333-2340.	2.3	80
180	Haplotype Inference for Population Data with Genotyping Errors. Biometrical Journal, 2009, 51, 644-658.	0.6	1
181	Characterization of a t(5;8)(q31;q21) translocation in a patient with mental retardation and congenital heart disease: implications for involvement of RUNX1T1 in human brain and heart development. European Journal of Human Genetics, 2009, 17, 1010-1018.	1.4	20
182	Review of the Lynch syndrome: history, molecular genetics, screening, differential diagnosis, and medicolegal ramifications. Clinical Genetics, 2009, 76, 1-18.	1.0	672
183	Self-organizing map approaches for the haplotype assembly problem. Mathematics and Computers in Simulation, 2009, 79, 3026-3037.	2.4	8
184	Random Effects Models in a Meta-Analysis of the Accuracy of Two Diagnostic Tests Without a Gold Standard. Journal of the American Statistical Association, 2009, 104, 512-523.	1.8	71
185	Generation of Medaka Fish Haploid Embryonic Stem Cells. Science, 2009, 326, 430-433.	6.0	157
187	Colorectal Cancer Due to Deficiency in DNA Mismatch Repair Function. Advances in Anatomic Pathology, 2009, 16, 405-417.	2.4	132
189	Medaka fish stem cells and their applications. Science China Life Sciences, 2010, 53, 426-434.	2.3	19
190	Detection of genetic alterations in hereditary colorectal cancer screeningâ~†. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2010, 693, 19-31.	0.4	29
191	Direct determination of molecular haplotypes by chromosome microdissection. Nature Methods, 2010, 7, 299-301.	9.0	66
192	Confirmation of Linkage to and Localization of Familial Colon Cancer Risk Haplotype on Chromosome 9q22. Cancer Research, 2010, 70, 5409-5418.	0.4	42
193	Hereditary Colorectal Cancer. , 2010, , .		1

#	Article	IF	CITATIONS
194	Forward and Reverse Genetics through Derivation of Haploid Mouse Embryonic Stem Cells. Cell Stem Cell, 2011, 9, 563-574.	5.2	208
195	The Extent of Linkage Disequilibrium and Computational Challenges of Single Nucleotide Polymorphisms in Genome-Wide Association Studies. Current Drug Metabolism, 2011, 12, 498-506.	0.7	5
196	Completely phased genome sequencing through chromosome sorting. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 12-17.	3.3	93
197	A clinically complex form of dominant optic atrophy (OPA8) maps on chromosome 16. Human Molecular Genetics, 2011, 20, 1893-1905.	1.4	36
198	Cilia and Models for Studying Structure and Function. Proceedings of the American Thoracic Society, 2011, 8, 423-429.	3.5	39
199	Sequence and expression analysis of gaps in human chromosome 20. Nucleic Acids Research, 2012, 40, 6660-6672.	6.5	5
200	Generation of Genetically Modified Mice by Oocyte Injection of Androgenetic Haploid Embryonic Stem Cells. Cell, 2012, 149, 605-617.	13.5	168
201	A Simple PCR–RFLP Method for Genetic Phase Determination in Compound Heterozygotes. Frontiers in Genetics, 2012, 2, 108.	1.1	1
202	An American founder mutation in <i>MLH1</i> . International Journal of Cancer, 2012, 130, 2088-2095.	2.3	12
203	The History of Lynch Syndrome. Familial Cancer, 2013, 12, 145-157.	0.9	76
203 204	The History of Lynch Syndrome. Familial Cancer, 2013, 12, 145-157. Complementarity of Binding Motifs is a General Property of HLA-A and HLA-B Molecules and Does Not Seem to Effect HLA Haplotype Composition. Frontiers in Immunology, 2013, 4, 374.	0.9	<b>76</b> 5
203 204 205	The History of Lynch Syndrome. Familial Cancer, 2013, 12, 145-157.         Complementarity of Binding Motifs is a General Property of HLA-A and HLA-B Molecules and Does Not Seem to Effect HLA Haplotype Composition. Frontiers in Immunology, 2013, 4, 374.         HnRNP L and hnRNP LL antagonistically modulate PTB-mediated splicing suppression of CHRNA1 pre-mRNA. Scientific Reports, 2013, 3, 2931.	0.9 2.2 1.6	76 5 41
203 204 205 207	The History of Lynch Syndrome. Familial Cancer, 2013, 12, 145-157.         Complementarity of Binding Motifs is a General Property of HLA-A and HLA-B Molecules and Does Not Seem to Effect HLA Haplotype Composition. Frontiers in Immunology, 2013, 4, 374.         HnRNP L and hnRNP LL antagonistically modulate PTB-mediated splicing suppression of CHRNA1 pre-mRNA. Scientific Reports, 2013, 3, 2931.         Colon: Colorectal adenocarcinoma. Atlas of Genetics and Cytogenetics in Oncology and Haematology, 2013,	0.9 2.2 1.6 0.1	76 5 41 3
203 204 205 207 208	The History of Lynch Syndrome. Familial Cancer, 2013, 12, 145-157.         Complementarity of Binding Motifs is a General Property of HLA-A and HLA-B Molecules and Does Not Seem to Effect HLA Haplotype Composition. Frontiers in Immunology, 2013, 4, 374.         HnRNP L and hnRNP LL antagonistically modulate PTB-mediated splicing suppression of CHRNA1 pre-mRNA. Scientific Reports, 2013, 3, 2931.         Colon: Colorectal adenocarcinoma. Atlas of Genetics and Cytogenetics in Oncology and Haematology, 2013,         Lynch Syndrome 101 (Years, That Is). American Society of Clinical Oncology Educational Book / ASCO American Society of Clinical Oncology Meeting, 2014,, 27-32.	0.9 2.2 1.6 0.1 1.8	<ul> <li>76</li> <li>5</li> <li>41</li> <li>3</li> <li>2</li> </ul>
203 204 205 207 208	The History of Lynch Syndrome. Familial Cancer, 2013, 12, 145-157.         Complementarity of Binding Motifs is a General Property of HLA-A and HLA-B Molecules and Does Not Seem to Effect HLA Haplotype Composition. Frontiers in Immunology, 2013, 4, 374.         HnRNP L and hnRNP LL antagonistically modulate PTB-mediated splicing suppression of CHRNA1 pre-mRNA. Scientific Reports, 2013, 3, 2931.         Colon: Colorectal adenocarcinoma. Atlas of Genetics and Cytogenetics in Oncology and Haematology, 2013,         Lynch Syndrome 101 (Years, That Is). American Society of Clinical Oncology Educational Book / ASCO American Society of Clinical Oncology Meeting, 2014, , 27-32.         A hidden Markov model for haplotype inference for present-absent data of clustered genes using identified haplotypes and haplotype patterns. Frontiers in Genetics, 2014, 5, 267.	0.9 2.2 1.6 0.1 1.8 1.1	<ul> <li>76</li> <li>5</li> <li>41</li> <li>3</li> <li>2</li> <li>1</li> </ul>
203 204 205 207 208 209	The History of Lynch Syndrome. Familial Cancer, 2013, 12, 145-157.         Complementarity of Binding Motifs is a General Property of HLA-A and HLA-B Molecules and Does Not Seem to Effect HLA Haplotype Composition. Frontiers in Immunology, 2013, 4, 374.         HnRNP L and hnRNP LL antagonistically modulate PTB-mediated splicing suppression of CHRNA1 pre-mRNA. Scientific Reports, 2013, 3, 2931.         Colon: Colorectal adenocarcinoma. Atlas of Genetics and Cytogenetics in Oncology and Haematology, 2013,         Lynch Syndrome 101 (Years, That Is). American Society of Clinical Oncology Educational Book / ASCO American Society of Clinical Oncology Meeting, 2014,, 27-32.         A hidden Markov model for haplotype inference for present-absent data of clustered genes using identified haplotypes and haplotype patterns. Frontiers in Genetics, 2014, 5, 267.         Molecular endocrinology and endocrine genetics, 2014,, 9-33.e1.	0.9 2.2 1.6 0.1 1.8 1.1	<ul> <li>76</li> <li>5</li> <li>41</li> <li>3</li> <li>2</li> <li>1</li> <li>1</li> </ul>
<ul> <li>203</li> <li>204</li> <li>205</li> <li>207</li> <li>208</li> <li>209</li> <li>210</li> <li>211</li> </ul>	The History of Lynch Syndrome. Familial Cancer, 2013, 12, 145-157.         Complementarity of Binding Motifs is a General Property of HLA-A and HLA-B Molecules and Does Not Seem to Effect HLA Haplotype Composition. Frontiers in Immunology, 2013, 4, 374.         HnRNP L and hnRNP LL antagonistically modulate PTB-mediated splicing suppression of CHRNA1 pre-mRNA. Scientific Reports, 2013, 3, 2931.         Colon: Colorectal adenocarcinoma. Atlas of Genetics and Cytogenetics in Oncology and Haematology, 2013,         Lynch Syndrome 101 (Years, That Is). American Society of Clinical Oncology Educational Book / ASCO American Society of Clinical Oncology Meeting, 2014, 1, 27-32.         A hidden Markov model for haplotype inference for present-absent data of clustered genes using identified haplotypes and haplotype patterns. Frontiers in Genetics, 2014, 5, 267.         Molecular endocrinology and endocrine genetics. , 2014, , 9-33.e1.         Boveri at 100: Theodor Boveri and genetic predisposition to cancer. Journal of Pathology, 2014, 234, 142-145.	0.9 2.2 1.6 0.1 1.8 1.1 2.1	<ul> <li>76</li> <li>5</li> <li>41</li> <li>3</li> <li>2</li> <li>1</li> <li>18</li> </ul>

		Citation R	EPORT	
#	Article		IF	CITATIONS
213	A simple method for gene phasing using mate pair sequencing. BMC Medical Genetics,	2014, 15, 19.	2.1	8
214	Genome wide functional genetics in haploid cells. FEBS Letters, 2014, 588, 2415-2421		1.3	20
215	Milestones of Lynch syndrome: 1895–2015. Nature Reviews Cancer, 2015, 15, 181-	194.	12.8	603
216	Haplotype-resolved genome sequencing: experimental methods and applications. Natu Genetics, 2015, 16, 344-358.	re Reviews	7.7	156
217	Quantitative haplotyping of PCR products by nonsynchronous pyrosequencing with di- Analytical and Bioanalytical Chemistry, 2016, 408, 8263-8271.	base addition.	1.9	3
218	Haplotype-based Statistical Inference for Population-based Case–control and Cross- with Complex Sample Designs. Journal of Survey Statistics and Methodology, 2016, 4,	Sectional Studies 188-214.	0.5	2
219	Conditional Displacement Hybridization Assay for Multiple SNP Phasing. Analytical Che 9961-9966.	mistry, 2017, 89,	3.2	9
220	Late-time growth rate, mixing, and anisotropy in the multimode narrowband Richtmyer instability: The <i><math>\hat{J}_{s}</math>/i&gt;-group collaboration. Physics of Fluids, 2017, 29, .</i>	'–Meshkov	1.6	79
221	Classification of Genetic Variants. , 2018, , 257-280.			0
222	Direct numerical simulation of the multimode narrowband Richtmyer–Meshkov insta Computers and Fluids, 2019, 194, 104309.	bility.	1.3	17
223	Turbulent transport and mixing in the multimode narrowband Richtmyer-Meshkov inst of Fluids, 2019, 31, .	ability. Physics	1.6	26
224	A New Fast Phasing Method Based On Haplotype Subtraction. Journal of Molecular Dia 21, 427-436.	gnostics, 2019,	1.2	1
225	Current advances in haploid stem cells. Protein and Cell, 2020, 11, 23-33.		4.8	9
226	A More Open Approach Is Needed to Develop Cell-Based Fish Technology: It Starts with Earth, 2020, 3, 54-64.	n Zebrafish. One	3.6	31
227	Theranostics Approaches to Gastric and Colon Cancer. Diagnostics and Therapeutic Ac Malignancies, 2020, , .	vances in Gl	0.2	2
229	Historical Aspects of Lynch Syndrome. , 2010, , 15-42.			4
230	The Genetics of Colorectal Cancer. , 2013, , 1-24.			2
231	Molecular Endocrinology and Endocrine Genetics. , 2008, , 1-25.			1

#	Article	IF	CITATIONS
232	Identification of a large rearrangement of theBRCA1 gene using colour bar code on combed DNA in an American breast/ovarian cancer family previously studied by direct sequencing. Journal of Medical Genetics, 2001, 38, 388-392.	1.5	45
233	SNP Haplotype Mapping in a Small ALS Family. PLoS ONE, 2009, 4, e5687.	1.1	5
234	Mismatch repair protein expression in colorectal cancer. Journal of Gastrointestinal Oncology, 2013, 4, 397-408.	0.6	42
235	Haploid Analysis (Monosomal Hybrid Technique). , 2004, , 565-569.		0
236	Hereditary colorectal cancer and brain tumor syndromes. , 2006, , 993-1001.		0
237	MOLECULAR BIOLOGY OF COLORECTAL CANCER. , 2008, , 867-896.		0
238	Molecular Genetics and Cancer Risks in Lynch Syndrome. , 2008, , 129-147.		0
239	Colon: Colorectal adenocarcinoma. Atlas of Genetics and Cytogenetics in Oncology and Haematology, 2011, , .	0.1	1
240	Genetics of Colon Cancer Susceptibility. , 2012, , 23-45.		0
241	Historical Development of Lynch Syndrome. , 2013, , 1-24.		0
242	Hereditary Cancer. , 2017, , 335-353.		1
243	Understanding Colorectal Cancer: The Basics. Diagnostics and Therapeutic Advances in Gl Malignancies, 2020, , 93-115.	0.2	0
244	Paraganglioma and Pheochromocytoma. , 2008, , 165-212.		0
245	The evolution of colorectal cancer genetics-Part 1: from discovery to practice. Journal of Gastrointestinal Oncology, 2014, 5, 326-35.	0.6	15
247	Entamoeba histolytica and Probable Effect on Production Microsatellite Instability in Colorectal Cancer. Current Microbiology, 2022, 79, 111.	1.0	1
248	Experimental method for haplotype phasing across the entire length of chromosome 21 in trisomy 21 cells using a chromosome elimination technique. Journal of Human Genetics, 2022, 67, 565-572.	1.1	2