

# Albert de la Chapelle

## List of Publications by Year in descending order

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Version: 2024-02-01

586  
papers

78,756  
citations

668

122  
h-index

529

266  
g-index

596  
all docs

596  
docs citations

596  
times ranked

49606  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identifying Novel Susceptibility Genes for Colorectal Cancer Risk From a Transcriptome-Wide Association Study of 125,478 Subjects. <i>Gastroenterology</i> , 2021, 160, 1164-1178.e6.	1.3	36
2	Poor Survival and Differential Impact of Genetic Features of Black Patients with Acute Myeloid Leukemia. <i>Cancer Discovery</i> , 2021, 11, 626-637.	9.4	41
3	Double somatic mismatch repair gene pathogenic variants as common as Lynch syndrome among endometrial cancer patients. <i>Gynecologic Oncology</i> , 2021, 160, 161-168.	1.4	24
4	Genetically predicted circulating concentrations of micronutrients and risk of colorectal cancer among individuals of European descent: a Mendelian randomization study. <i>American Journal of Clinical Nutrition</i> , 2021, 113, 1490-1502.	4.7	27
5	Genetic architectures of proximal and distal colorectal cancer are partly distinct. <i>Gut</i> , 2021, 70, 1325-1334.	12.1	44
6	Multiethnic genome-wide association study of differentiated thyroid cancer in the EPITHYR consortium. <i>International Journal of Cancer</i> , 2021, 148, 2935-2946.	5.1	11
7	Gene expression signature predicts relapse in adult patients with cytogenetically normal acute myeloid leukemia. <i>Blood Advances</i> , 2021, 5, 1474-1482.	5.2	20
8	Response to Li and Hopper. <i>American Journal of Human Genetics</i> , 2021, 108, 527-529.	6.2	5
9	A novel essential splice site variant in SPTB in a large hereditary spherocytosis family. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2021, 9, e1641.	1.2	2
10	Prospective Statewide Study of Universal Screening for Hereditary Colorectal Cancer: The Ohio Colorectal Cancer Prevention Initiative. <i>JCO Precision Oncology</i> , 2021, 5, 779-791.	3.0	31
11	Transcriptome analysis discloses dysregulated genes in normal appearing tumor-adjacent thyroid tissues from patients with papillary thyroid carcinoma. <i>Scientific Reports</i> , 2021, 11, 14126.	3.3	9
12	Variation in the risk of colorectal cancer in families with Lynch syndrome: a retrospective cohort study. <i>Lancet Oncology</i> , The, 2021, 22, 1014-1022.	10.7	58
13	Cumulative Burden of Colorectal Cancer-Associated Genetic Variants Is More Strongly Associated With Early-Onset vs Late-Onset Cancer. <i>Gastroenterology</i> , 2020, 158, 1274-1286.e12.	1.3	110
14	Circulating Levels of Insulin-like Growth Factor 1 and Insulin-like Growth Factor Binding Protein 3 Associate With Risk of Colorectal Cancer Based on Serologic and Mendelian Randomization Analyses. <i>Gastroenterology</i> , 2020, 158, 1300-1312.e20.	1.3	90
15	Inherited myeloproliferative neoplasm risk affects haematopoietic stem cells. <i>Nature</i> , 2020, 586, 769-775.	27.8	101
16	Mutational landscape and clinical outcome of patients with de novo acute myeloid leukemia and rearrangements involving 11q23/ <i>KMT2A</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 26340-26346.	7.1	59
17	Characterizing the function of EPB41L4A in the predisposition to papillary thyroid carcinoma. <i>Scientific Reports</i> , 2020, 10, 19984.	3.3	3
18	Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 432-444.	6.2	124

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19	GWAS of thyroid stimulating hormone highlights pleiotropic effects and inverse association with thyroid cancer. <i>Nature Communications</i> , 2020, 11, 3981.	12.8	86
20	Adiposity, metabolites, and colorectal cancer risk: Mendelian randomization study. <i>BMC Medicine</i> , 2020, 18, 396.	5.5	76
21	Hemochromatosis risk genotype is not associated with colorectal cancer or age at its diagnosis. <i>Human Genetics and Genomics Advances</i> , 2020, 1, 100010.	1.7	3
22	Additional gene mutations may refine the 2017 European LeukemiaNet classification in adult patients with de novo acute myeloid leukemia aged <math>\leq 60</math> years. <i>Leukemia</i> , 2020, 34, 3215-3227.	7.2	66
23	Assessing thyroid cancer risk using polygenic risk scores. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 5997-6002.	7.1	39
24	Thyroid Carcinomas That Occur in Familial Adenomatous Polyposis Patients Recurrently Harbor Somatic Variants in <i>APC</i> , <i>BRAF</i> , and <i>KTM2D</i> . <i>Thyroid</i> , 2020, 30, 380-388.	4.5	18
25	A Truncating Germline Mutation of <i>TINF2</i> in Individuals with Thyroid Cancer or Melanoma Results in Longer Telomeres. <i>Thyroid</i> , 2020, 30, 204-213.	4.5	27
26	Variants in <i>LRRC34</i> reveal distinct mechanisms for predisposition to papillary thyroid carcinoma. <i>Journal of Medical Genetics</i> , 2020, 57, 519-527.	3.2	3
27	Genetic Characterization and Prognostic Relevance of Acquired Uniparental Disomies in Cytogenetically Normal Acute Myeloid Leukemia. <i>Clinical Cancer Research</i> , 2019, 25, 6524-6531.	7.0	12
28	Methylated SEPTIN9 plasma test for colorectal cancer detection may be applicable to Lynch syndrome. <i>BMJ Open Gastroenterology</i> , 2019, 6, e000299.	2.7	9
29	Implementation of standardized variant-calling nomenclature in the age of next-generation sequencing: where do we stand?. <i>Leukemia</i> , 2019, 33, 809-810.	7.2	1
30	Risk Haplotypes Uniquely Associated with Radioiodine-Refractory Thyroid Cancer Patients of High African Ancestry. <i>Thyroid</i> , 2019, 29, 530-539.	4.5	8
31	Clinical characteristics of patients with colorectal cancer with double somatic mismatch repair mutations compared with Lynch syndrome. <i>Journal of Medical Genetics</i> , 2019, 56, 462-470.	3.2	61
32	Identification of Rare Variants Predisposing to Thyroid Cancer. <i>Thyroid</i> , 2019, 29, 946-955.	4.5	41
33	Microsatellite Instability Occurs in a Subset of Follicular Thyroid Cancers. <i>Thyroid</i> , 2019, 29, 523-529.	4.5	31
34	Neuroblastoma RAS viral oncogene homolog mRNA is differentially spliced to give five distinct isoforms: implications for melanoma therapy. <i>Melanoma Research</i> , 2019, 29, 491-500.	1.2	6
35	Fine mapping of 14q13 reveals novel variants associated with different histological subtypes of papillary thyroid carcinoma. <i>International Journal of Cancer</i> , 2019, 144, 503-512.	5.1	4
36	Discovery of common and rare genetic risk variants for colorectal cancer. <i>Nature Genetics</i> , 2019, 51, 76-87.	21.4	377

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37	Genome-wide association study identifies an acute myeloid leukemia susceptibility locus near BICRA. <i>Leukemia</i> , 2019, 33, 771-775.	7.2	15
38	The 2017 European Leukemianet Genetic Risk Classification Performs Poorly in Older Patients with Acute Myeloid Leukemia (AML) and Should be Refined to Identify Patients Requiring Additional or Alternative Treatment. <i>Blood</i> , 2019, 134, 2681-2681.	1.4	1
39	Clinical implications of GWAS variants associated with differentiated thyroid cancer. <i>Endokrynologia Polska</i> , 2019, 70, 423-429.	1.0	7
40	Distinct Gene Expression Profiles and Mutations Associate with Outcome in Younger Adults with De Novo Cytogenetically Normal Acute Myeloid Leukemia (CN-AML) (Alliance). <i>Blood</i> , 2019, 134, 1247-1247.	1.4	1
41	Mutation patterns identify adult patients with de novo acute myeloid leukemia aged 60 years or older who respond favorably to standard chemotherapy: an analysis of Alliance studies. <i>Leukemia</i> , 2018, 32, 1338-1348.	7.2	80
42	The Role of NRG1 in the Predisposition to Papillary Thyroid Carcinoma. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2018, 103, 1369-1379.	3.6	23
43	The role of SMAD3 in the genetic predisposition to papillary thyroid carcinoma. <i>Genetics in Medicine</i> , 2018, 20, 927-935.	2.4	12
44	Assessment of Tumor Sequencing as a Replacement for Lynch Syndrome Screening and Current Molecular Tests for Patients With Colorectal Cancer. <i>JAMA Oncology</i> , 2018, 4, 806.	7.1	136
45	Penetrance of a rare familial mutation predisposing to papillary thyroid cancer. <i>Familial Cancer</i> , 2018, 17, 431-434.	1.9	7
46	Cancer Risks for <i>PMS2</i> -Associated Lynch Syndrome. <i>Journal of Clinical Oncology</i> , 2018, 36, 2961-2968.	1.6	147
47	Identification of a Recurrent <i>LMO7</i> - <i>BRAF</i> Fusion in Papillary Thyroid Carcinoma. <i>Thyroid</i> , 2018, 28, 748-754.	4.5	19
48	NF1 mutations are recurrent in adult acute myeloid leukemia and confer poor outcome. <i>Leukemia</i> , 2018, 32, 2536-2545.	7.2	33
49	Additional Gene Mutations Refine the 2017 European Leukemianet (ELN) Classification of Adult Patients (Pts) with De Novo Acute Myeloid Leukemia (AML) Aged <60 Years: An Analysis of Alliance for Clinical Trials in Oncology (Alliance) Studies. <i>Blood</i> , 2018, 132, 2740-2740.	1.4	1
50	Mutations in Genes Associated with Familial Predisposition to Myeloid Neoplasms: Their Frequency and Associations with Pretreatment Characteristics in Adult Patients (Pts) with Presumably Sporadic De Novo Acute Myeloid Leukemia (AML). <i>Blood</i> , 2018, 132, 1478-1478.	1.4	13
51	Uniparental Disomies (UPD) of Chromosome 13q Is Associated with Shorter Disease-Free Survival in Adult Patients (Pts) with De Novo Cytogenetically Normal Acute Myeloid Leukemia (CN-AML). <i>Blood</i> , 2018, 132, 2777-2777.	1.4	0
52	Genome-Wide Association Study (GWAS) Identifies a Significant Acute Myeloid Leukemia (AML) Susceptibility Locus Near BICRA. <i>Blood</i> , 2018, 132, 85-85.	1.4	0
53	A genome-wide association study yields five novel thyroid cancer risk loci. <i>Nature Communications</i> , 2017, 8, 14517.	12.8	117
54	Comprehensive population-wide analysis of Lynch syndrome in Iceland reveals founder mutations in MSH6 and PMS2. <i>Nature Communications</i> , 2017, 8, 14755.	12.8	96

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55	No evidence for microsatellite instability in acute myeloid leukemia. <i>Leukemia</i> , 2017, 31, 1474-1476.	7.2	11
56	The mutational oncoprint of recurrent cytogenetic abnormalities in adult patients with de novo acute myeloid leukemia. <i>Leukemia</i> , 2017, 31, 2211-2218.	7.2	69
57	MYH9 binds to lncRNA gene <i>PTCSC2</i> and regulates <i>FOXE1</i> in the 9q22 thyroid cancer risk locus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 474-479.	7.1	80
58	Prevalence and Spectrum of Germline Cancer Susceptibility Gene Mutations Among Patients With Early-Onset Colorectal Cancer. <i>JAMA Oncology</i> , 2017, 3, 464.	7.1	510
59	Identification of NRAS isoform 2 overexpression as a mechanism facilitating BRAF inhibitor resistance in malignant melanoma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 9629-9634.	7.1	16
60	Mutation Frequencies in Patients With Early-Onset Colorectal Cancer—Reply. <i>JAMA Oncology</i> , 2017, 3, 1587.	7.1	5
61	Mutations in the CCND1 and CCND2 genes are frequent events in adult patients with t(8;21)(q22;q22) acute myeloid leukemia. <i>Leukemia</i> , 2017, 31, 1278-1285.	7.2	47
62	Mutational Landscape and Gene Expression Patterns in Adult Acute Myeloid Leukemias with Monosomy 7 as a Sole Abnormality. <i>Cancer Research</i> , 2017, 77, 207-218.	0.9	23
63	The early years of the ESHG leading to the reform of 1988 and the spirit of the Sestri Levante school. <i>European Journal of Human Genetics</i> , 2017, 25, S6-S12.	2.8	1
64	Variants in microRNA genes in familial papillary thyroid carcinoma. <i>Oncotarget</i> , 2017, 8, 6475-6482.	1.8	8
65	HABP2 G534E Variant in Papillary Thyroid Carcinoma. <i>PLoS ONE</i> , 2016, 11, e0146315.	2.5	31
66	The Olympic Games and Athletic Sex Assignment. <i>JAMA - Journal of the American Medical Association</i> , 2016, 316, 1359.	7.4	12
67	Colon Cancer Germline Genetics: The Unbelievable Year 1993 and Thereafter. <i>Cancer Research</i> , 2016, 76, 4025-4027.	0.9	1
68	Genome-Wide Expression Screening Discloses Long Noncoding RNAs Involved in Thyroid Carcinogenesis. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2016, 101, 4005-4013.	3.6	47
69	Adult acute myeloid leukemia with trisomy 11 as the sole abnormality is characterized by the presence of five distinct gene mutations: MLL-PTD, DNMT3A, U2AF1, FLT3-ITD and IDH2. <i>Leukemia</i> , 2016, 30, 2254-2258.	7.2	16
70	Structural characterization of NRAS isoform 5. <i>Protein Science</i> , 2016, 25, 1069-1074.	7.6	5
71	Primary Cell Culture Systems for Human Thyroid Studies. <i>Thyroid</i> , 2016, 26, 1131-1140.	4.5	14
72	Frequent PIK3CA Mutations in Colorectal and Endometrial Tumors With 2 or More Somatic Mutations in Mismatch Repair Genes. <i>Gastroenterology</i> , 2016, 151, 440-447.e1.	1.3	36

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73	Papillary Thyroid Carcinoma: Association Between Germline DNA Variant Markers and Clinical Parameters. <i>Thyroid</i> , 2016, 26, 1276-1284.	4.5	32
74	Dissection of the Major Hematopoietic Quantitative Trait Locus in Chromosome 6q23.3 Identifies miR-3662 as a Player in Hematopoiesis and Acute Myeloid Leukemia. <i>Cancer Discovery</i> , 2016, 6, 1036-1051.	9.4	14
75	Patients with colorectal cancer associated with Lynch syndrome and MLH1 promoter hypermethylation have similar prognoses. <i>Genetics in Medicine</i> , 2016, 18, 863-868.	2.4	30
76	PMS2 monoallelic mutation carriers: the known unknown. <i>Genetics in Medicine</i> , 2016, 18, 13-19.	2.4	51
77	The Mutational Patterns Associated with Cytogenetic Subsets of De Novo Acute Myeloid Leukemia (AML): A Study of 1603 Adult Patients (Pts). <i>Blood</i> , 2016, 128, 287-287.	1.4	0
78	CCND1 and CCND2 Mutations Are Frequent in Adults with Core-Binding Factor Acute Myeloid Leukemia (CBF-AML) with t(8;21)(q22;q22). <i>Blood</i> , 2016, 128, 2740-2740.	1.4	0
79	Somatic <i>MED12</i> mutations are associated with poor prognosis markers in chronic lymphocytic leukemia. <i>Oncotarget</i> , 2015, 6, 1884-1888.	1.8	49
80	PD-1 Blockade in Tumors with Mismatch-Repair Deficiency. <i>New England Journal of Medicine</i> , 2015, 372, 2509-2520.	27.0	7,696
81	A germline mutation in <i>SRRM2</i> , a splicing factor gene, is implicated in papillary thyroid carcinoma predisposition. <i>Scientific Reports</i> , 2015, 5, 10566.	3.3	83
82	Multiple functional variants in long-range enhancer elements contribute to the risk of SNP rs965513 in thyroid cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 6128-6133.	7.1	79
83	Immunoglobulin transcript sequence and somatic hypermutation computation from unselected RNA-seq reads in chronic lymphocytic leukemia. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 4322-4327.	7.1	38
84	PTCSC3 Is Involved in Papillary Thyroid Carcinoma Development by Modulating <i>S100A4</i> Gene Expression. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2015, 100, E1370-E1377.	3.6	65
85	MicroRNA-3151 inactivates TP53 in <i>BRAF</i> -mutated human malignancies. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E6744-51.	7.1	17
86	Genetic Predisposition to Papillary Thyroid Carcinoma: Involvement of <i>FOXE1</i> , <i>TSHR</i> , and a Novel lincRNA Gene, <i>PTCSC2</i> . <i>Journal of Clinical Endocrinology and Metabolism</i> , 2015, 100, E164-E172.	3.6	93
87	Identification of Novel Genetic Loci Associated with Thyroid Peroxidase Antibodies and Clinical Thyroid Disease. <i>PLoS Genetics</i> , 2014, 10, e1004123.	3.5	150
88	Prostate cancer incidence in males with Lynch syndrome. <i>Genetics in Medicine</i> , 2014, 16, 553-557.	2.4	88
89	Intronic <i>miR-3151</i> Within <i>BAALC</i> Drives Leukemogenesis by Deregulating the TP53 Pathway. <i>Science Signaling</i> , 2014, 7, ra36.	3.6	18
90	MicroRNA-related sequence variations in human cancers. <i>Human Genetics</i> , 2014, 133, 463-469.	3.8	27

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91	Variants in the <i>ATM</i> - <i>CHEK2</i> - <i>BRCA1</i> axis determine genetic predisposition and clinical presentation of papillary thyroid carcinoma. <i>Genes Chromosomes and Cancer</i> , 2014, 53, 516-523.	2.8	50
92	Biallelic <i>MUTYH</i> mutations can mimic Lynch syndrome. <i>European Journal of Human Genetics</i> , 2014, 22, 1334-1337.	2.8	87
93	Colon and Endometrial Cancers With Mismatch Repair Deficiency Can Arise From Somatic, Rather Than Germline, Mutations. <i>Gastroenterology</i> , 2014, 147, 1308-1316.e1.	1.3	328
94	<i>NRAS</i> isoforms differentially affect downstream pathways, cell growth, and cell transformation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 4179-4184.	7.1	27
95	Micro RNA-3662 Located within the Major Hematopoietic Differentiation Quantitative Trait Locus on Chromosome 6q23 Acts As a Tumor Suppressor By Direct Targeting of the NF- $\kappa$ B Transcriptional Activation Pathway. <i>Blood</i> , 2014, 124, 2216-2216.	1.4	0
96	Immunodepletion Plasma Proteomics by TripleTOF 5600 and Orbitrap Elite/LTQ-Orbitrap Velos/Q Exactive Mass Spectrometers. <i>Journal of Proteome Research</i> , 2013, 12, 4351-4365.	3.7	43
97	Cumulative Risk Impact of Five Genetic Variants Associated with Papillary Thyroid Carcinoma. <i>Thyroid</i> , 2013, 23, 1532-1540.	4.5	63
98	Recurrent and founder mutations in the <i>PMS2</i> gene. <i>Clinical Genetics</i> , 2013, 83, 238-243.	2.0	16
99	How do we approach the goal of identifying everybody with Lynch Syndrome?. <i>Familial Cancer</i> , 2013, 12, 313-317.	1.9	58
100	<i>SRGAP1</i> Is a Candidate Gene for Papillary Thyroid Carcinoma Susceptibility. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2013, 98, E973-E980.	3.6	74
101	In-Depth Characterization of the MicroRNA Transcriptome in Normal Thyroid and Papillary Thyroid Carcinoma. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2013, 98, E1401-E1409.	3.6	125
102	Unraveling the Genetic Predisposition to Differentiated Thyroid Carcinoma. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2013, 98, 3974-3976.	3.6	5
103	Germline Allele-Specific Expression of <i>DAPK1</i> in Chronic Lymphocytic Leukemia. <i>PLoS ONE</i> , 2013, 8, e55261.	2.5	24
104	Ultra-Rare Mutation in Long-Range Enhancer Predisposes to Thyroid Carcinoma with High Penetrance. <i>PLoS ONE</i> , 2013, 8, e61920.	2.5	36
105	Characterization of a New Chronic Lymphocytic Leukemia Cell Line for Mechanistic In Vitro and In Vivo Studies Relevant to Disease. <i>PLoS ONE</i> , 2013, 8, e76607.	2.5	51
106	Performance of <i>PREMM1,2,6</i> , <i>MMRpredict</i> , and <i>MMRpro</i> in detecting Lynch syndrome among endometrial cancer cases. <i>Genetics in Medicine</i> , 2012, 14, 670-680.	2.4	40
107	Heritable polymorphism predisposes to high <i>BAALC</i> expression in acute myeloid leukemia. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 6668-6673.	7.1	23
108	MicroRNA Signature in Thyroid Fine Needle Aspiration Cytology Applied to $\neq$ Atypia of Undetermined Significance Cases. <i>Thyroid</i> , 2012, 22, 9-16.	4.5	92

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109	Characterization of the colorectal cancer-associated enhancer MYC-335 at 8q24: the role of rs67491583. <i>Cancer Genetics</i> , 2012, 205, 25-33.	0.4	24
110	Identification of Lynch Syndrome Among Patients With Colorectal Cancer. <i>JAMA - Journal of the American Medical Association</i> , 2012, 308, 1555.	7.4	443
111	Discovery of common variants associated with low TSH levels and thyroid cancer risk. <i>Nature Genetics</i> , 2012, 44, 319-322.	21.4	208
112	Cancer Risks for Relatives of Patients With Serrated Polyposis. <i>American Journal of Gastroenterology</i> , 2012, 107, 770-778.	0.4	80
113	miR-3151 interplays with its host gene BAALC and independently affects outcome of patients with cytogenetically normal acute myeloid leukemia. <i>Blood</i> , 2012, 120, 249-258.	1.4	64
114	The polymorphism rs944289 predisposes to papillary thyroid carcinoma through a large intergenic noncoding RNA gene of tumor suppressor type. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 8646-8651.	7.1	237
115	Microcephalic osteodysplastic primordial dwarfism type I with biallelic mutations in the <i>RNU4ATAC</i> gene. <i>Clinical Genetics</i> , 2012, 82, 140-146.	2.0	31
116	An American founder mutation in <i>MLH1</i> . <i>International Journal of Cancer</i> , 2012, 130, 2088-2095.	5.1	12
117	Evaluation of Allele-Specific Somatic Changes of Genome-Wide Association Study Susceptibility Alleles in Human Colorectal Cancers. <i>PLoS ONE</i> , 2012, 7, e37672.	2.5	8
118	Variants in the Netrin-1 Receptor <i>UNC5C</i> Prevent Apoptosis and Increase Risk of Familial Colorectal Cancer. <i>Gastroenterology</i> , 2011, 141, 2039-2046.	1.3	28
119	MicroRNAs in Thyroid Cancer. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2011, 96, 3326-3336.	3.6	115
120	Thyroid Hormone Receptor $\beta$ (THRB) Is a Major Target Gene for MicroRNAs Deregulated in Papillary Thyroid Carcinoma (PTC). <i>Journal of Clinical Endocrinology and Metabolism</i> , 2011, 96, E546-E553.	3.6	82
121	Telomere Length and Telomerase Reverse Transcriptase Gene Copy Number in Patients with Papillary Thyroid Carcinoma. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2011, 96, E1876-E1880.	3.6	18
122	Mutations in U4atac snRNA, a Component of the Minor Spliceosome, in the Developmental Disorder MOPD I. <i>Science</i> , 2011, 332, 238-240.	12.6	223
123	The Search for Unaffected Individuals with Lynch Syndrome: Do the Ends Justify the Means?. <i>Cancer Prevention Research</i> , 2011, 4, 1-5.	1.5	138
124	MIR-3151, a Novel MicroRNA Embedded in BAALC, Is Only Weakly Co-Expressed with Its Host Gene and Independently Impacts on the Clinical Outcome of Older Patients (Pts) with De Novo Cytogenetically Normal Acute Myeloid Leukemia (CN-AML). <i>Blood</i> , 2011, 118, 1462-1462.	1.4	0
125	289 Sex and Gender in Sport: Fallacy of the "Level Playing Field". <i>Pediatric Research</i> , 2010, 68, 149-149.	2.3	2
126	Reprogramming of miRNA networks in cancer and leukemia. <i>Genome Research</i> , 2010, 20, 589-599.	5.5	331



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127	Phenotypic diversity in patients with multiple serrated polyps: a genetics clinic study. <i>International Journal of Colorectal Disease</i> , 2010, 25, 703-712.	2.2	48
128	Low cancer incidence rates in Ohio Amish. <i>Cancer Causes and Control</i> , 2010, 21, 69-75.	1.8	29
129	Risk Factors for Colorectal Cancer in Patients with Multiple Serrated Polyps: A Cross-Sectional Case Series from Genetics Clinics. <i>PLoS ONE</i> , 2010, 5, e11636.	2.5	68
130	Role of PTPRJ genotype in papillary thyroid carcinoma risk. <i>Endocrine-Related Cancer</i> , 2010, 17, 1001-1006.	3.1	25
131	The genetic basis of colorectal cancer in a population-based incident cohort with a high rate of familial disease. <i>Gut</i> , 2010, 59, 1369-1377.	12.1	65
132	Clinical Relevance of Microsatellite Instability in Colorectal Cancer. <i>Journal of Clinical Oncology</i> , 2010, 28, 3380-3387.	1.6	273
133	Risks of Lynch Syndrome Cancers for MSH6 Mutation Carriers. <i>Journal of the National Cancer Institute</i> , 2010, 102, 193-201.	6.3	328
134	Allele-specific expression of TGFBR1 in colon cancer patients. <i>Carcinogenesis</i> , 2010, 31, 1800-1804.	2.8	18
135	Reply to L.H. Jensen et al and S. Jahn et al. <i>Journal of Clinical Oncology</i> , 2009, 27, e225-e225.	1.6	1
136	Infrequent Detection of Germline Allele-Specific Expression of TGFBR1 in Lymphoblasts and Tissues of Colon Cancer Patients. <i>Cancer Research</i> , 2009, 69, 4959-4961.	0.9	20
137	Polymorphic mature microRNAs from passenger strand of pre-miR-146a contribute to thyroid cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 1502-1505.	7.1	311
138	<i>Tgfr1</i> Haploinsufficiency Is a Potent Modifier of Colorectal Cancer Development. <i>Cancer Research</i> , 2009, 69, 678-686.	0.9	52
139	A Susceptibility Locus for Papillary Thyroid Carcinoma on Chromosome 8q24. <i>Cancer Research</i> , 2009, 69, 625-631.	0.9	133
140	Identifying Lynch syndrome. <i>International Journal of Cancer</i> , 2009, 125, 1492-1493.	5.1	32
141	Epitope-positive truncating MLH1 mutation and loss of PMS2: implications for IHC-directed genetic testing for lynch syndrome. <i>Familial Cancer</i> , 2009, 8, 501-504.	1.9	16
142	Common variants on 9q22.33 and 14q13.3 predispose to thyroid cancer in European populations. <i>Nature Genetics</i> , 2009, 41, 460-464.	21.4	353
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