

Parry J Guilford

List of Publications by Year in descending order

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Version: 2024-02-01

78
papers

9,748
citations

71061

41
h-index

74108

75
g-index

80
all docs

80
docs citations

80
times ranked

8691
citing authors

#	ARTICLE	IF	CITATIONS
1	E-Cadherin-Deficient Cells Are Sensitive to the Multikinase Inhibitor Dasatinib. <i>Cancers</i> , 2022, 14, 1609.	1.7	4
2	Loss of E-Cadherin Leads to Druggable Vulnerabilities in Sphingolipid Metabolism and Vesicle Trafficking. <i>Cancers</i> , 2022, 14, 102.	1.7	6
3	E-Cadherin-Deficient Epithelial Cells Are Sensitive to HDAC Inhibitors. <i>Cancers</i> , 2022, 14, 175.	1.7	8
4	Modelling hereditary diffuse gastric cancer initiation using transgenic mouse-derived gastric organoids and single-cell sequencing. <i>Journal of Pathology</i> , 2021, 254, 254-264.	2.1	11
5	Updated perspective and directions on hereditary diffuse gastric cancer. , 2021, , 217-258.		1
6	Discovery of AL-GDa62 as a Potential Synthetic Lethal Lead for the Treatment of Gastric Cancer. <i>Journal of Medicinal Chemistry</i> , 2021, 64, 18114-18142.	2.9	4
7	Hereditary diffuse gastric cancer: updated clinical practice guidelines. <i>Lancet Oncology</i> , The, 2020, 21, e386-e397.	5.1	237
8	Circulating tumor DNA is a sensitive marker for routine monitoring of treatment response in advanced colorectal cancer. <i>Carcinogenesis</i> , 2020, 41, 1507-1517.	1.3	11
9	Cohesin mutations are synthetic lethal with stimulation of WNT signaling. <i>ELife</i> , 2020, 9, .	2.8	22
10	An estimate of limited duration cancer prevalence in New Zealand using 'big' data. <i>New Zealand Medical Journal</i> , 2020, 133, 49-62.	0.5	1
11	E-cadherin-deficient cells have synthetic lethal vulnerabilities in plasma membrane organisation, dynamics and function. <i>Gastric Cancer</i> , 2019, 22, 273-286.	2.7	24
12	Comparison of Roche Cell-Free DNA collection Tubes to Streck Cell-Free DNA BCT s for sample stability using healthy volunteers. <i>Practical Laboratory Medicine</i> , 2019, 16, e00125.	0.6	18
13	Identification of c.1531C>T Pathogenic Variant in the CDH1 Gene as a Novel Germline Mutation of Hereditary Diffuse Gastric Cancer. <i>International Journal of Molecular Sciences</i> , 2019, 20, 4980.	1.8	12
14	A high-throughput screen to identify novel synthetic lethal compounds for the treatment of E-cadherin-deficient cells. <i>Scientific Reports</i> , 2019, 9, 12511.	1.6	13
15	Allosteric AKT Inhibitors Target Synthetic Lethal Vulnerabilities in E-Cadherin-Deficient Cells. <i>Cancers</i> , 2019, 11, 1359.	1.7	22
16	Clinical spectrum and pleiotropic nature of CDH1 germline mutations. <i>Journal of Medical Genetics</i> , 2019, 56, 199-208.	1.5	74
17	Hereditary gastric cancer: what's new? Update 2013-2018. <i>Familial Cancer</i> , 2019, 18, 363-367.	0.9	44
18	Germline CDH1 mutations are a significant contributor to the high frequency of early-onset diffuse gastric cancer cases in New Zealand Māori. <i>Familial Cancer</i> , 2019, 18, 83-90.	0.9	33

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19	E-cadherin signal sequence disruption: a novel mechanism underlying hereditary cancer. <i>Molecular Cancer</i> , 2018, 17, 112.	7.9	11
20	Clinical comparison of noninvasive urine tests for ruling out recurrent urothelial carcinoma. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2017, 35, 531.e15-531.e22.	0.8	94
21	Performance Characteristics of a Multigene Urine Biomarker Test for Monitoring for Recurrent Urothelial Carcinoma in a Multicenter Study. <i>Journal of Urology</i> , 2017, 197, 1419-1426.	0.2	82
22	Genome-wide methylation analysis identifies a core set of hypermethylated genes in CIMP-H colorectal cancer. <i>BMC Cancer</i> , 2017, 17, 228.	1.1	32
23	Risk of stomach cancer in Aotearoa/New Zealand: A Māori population based case-control study. <i>PLoS ONE</i> , 2017, 12, e0181581.	1.1	15
24	Where to from here? Posthumous healthcare data, digital e(lectronic)-mortality and New Zealand's healthcare future. <i>New Zealand Medical Journal</i> , 2017, 130, 64-70.	0.5	0
25	Hereditary diffuse gastric cancer: updated clinical guidelines with an emphasis on germline <i>CDH1</i> mutation carriers. <i>Journal of Medical Genetics</i> , 2015, 52, 361-374.	1.5	479
26	Hereditary Diffuse Gastric Cancer Syndrome. <i>JAMA Oncology</i> , 2015, 1, 23.	3.4	540
27	Synthetic Lethal Screens Identify Vulnerabilities in GPCR Signaling and Cytoskeletal Organization in E-Cadherin Deficient Cells. <i>Molecular Cancer Therapeutics</i> , 2015, 14, 1213-1223.	1.9	30
28	A Comparison of Real-Time and Endpoint Cell Viability Assays for Improved Synthetic Lethal Drug Validation. <i>Journal of Biomolecular Screening</i> , 2015, 20, 1286-1293.	2.6	51
29	Culture, law, ethics, and social implications: Is society ready for advanced. <i>Australasian Medical Journal</i> , 2014, 7, 200-202.	0.1	5
30	E-cadherin loss alters cytoskeletal organization and adhesion in non-malignant breast cells but is insufficient to induce an epithelial-mesenchymal transition. <i>BMC Cancer</i> , 2014, 14, 552.	1.1	108
31	Molecular Mechanisms of Hereditary Diffuse Gastric Cancer Initiation and Progression. , 2013, , 51-76.		0
32	Early events in cell adhesion and polarity during epithelial-mesenchymal transition. <i>Journal of Cell Science</i> , 2012, 125, 4417-4422.	1.2	286
33	A Multigene Urine Test for the Detection and Stratification of Bladder Cancer in Patients Presenting with Hematuria. <i>Journal of Urology</i> , 2012, 188, 741-747.	0.2	128
34	Hereditary diffuse gastric cancer: translation of CDH1 germline mutations into clinical practice. <i>Gastric Cancer</i> , 2010, 13, 1-10.	2.7	143
35	Oligonucleotide array outperforms SNP array on formalin-fixed paraffin-embedded clinical samples. <i>Cancer Genetics and Cytogenetics</i> , 2010, 198, 1-6.	1.0	10
36	Hereditary diffuse gastric cancer: updated consensus guidelines for clinical management and directions for future research. <i>Journal of Medical Genetics</i> , 2010, 47, 436-444.	1.5	495

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37	The International Testicular Cancer Linkage Consortium: A clinicopathologic descriptive analysis of 461 familial malignant testicular germ cell tumor kindred. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2010, 28, 492-499.	0.8	42
38	E-Cadherin Deficiency Initiates Gastric Signet-Ring Cell Carcinoma in Mice and Man. <i>Cancer Research</i> , 2009, 69, 2050-2056.	0.4	147
39	Epigenetic silencing in non-neoplastic epithelia identifies E-cadherin (<i>CDH1</i>) as a target for chemoprevention of lobular neoplasia. <i>Journal of Pathology</i> , 2009, 218, 265-272.	2.1	41
40	Hereditary diffuse gastric cancer: A manifestation of lost cell polarity. <i>Cancer Science</i> , 2009, 100, 1151-1157.	1.7	78
41	Increased levels of active c-Src distinguish invasive from in situ lobular lesions. <i>Breast Cancer Research</i> , 2009, 11, R45.	2.2	28
42	Analysis of the <i>DND1</i> gene in men with sporadic and familial testicular germ cell tumors. <i>Genes Chromosomes and Cancer</i> , 2008, 47, 247-252.	1.5	37
43	A novel diffuse gastric cancer susceptibility variant in E-cadherin (<i>CDH1</i>) intron 2: A case control study in an Italian population. <i>BMC Cancer</i> , 2008, 8, 138.	1.1	13
44	Predicting Clinical Outcome through Molecular Profiling in Stage III Melanoma. <i>Clinical Cancer Research</i> , 2008, 14, 5173-5180.	3.2	62
45	Secreted CXCL1 Is a Potential Mediator and Marker of the Tumor Invasion of Bladder Cancer. <i>Clinical Cancer Research</i> , 2008, 14, 2579-2587.	3.2	95
46	Development of a Multiplex RNA Urine Test for the Detection and Stratification of Transitional Cell Carcinoma of the Bladder. <i>Clinical Cancer Research</i> , 2008, 14, 742-749.	3.2	74
47	Hereditary diffuse gastric cancer and lost cell polarity: a short path to cancer. <i>Future Oncology</i> , 2008, 4, 229-239.	1.1	13
48	Genomic characterization of multiple clinical phenotypes of cancer using multivariate linear regression models. <i>Bioinformatics</i> , 2007, 23, 732-738.	1.8	7
49	Destabilized Adhesion in the Gastric Proliferative Zone and c-Src Kinase Activation Mark the Development of Early Diffuse Gastric Cancer. <i>Cancer Research</i> , 2007, 67, 2480-2489.	0.4	114
50	Multiple Gene Expression Classifiers from Different Array Platforms Predict Poor Prognosis of Colorectal Cancer. <i>Clinical Cancer Research</i> , 2007, 13, 498-507.	3.2	114
51	A short guide to hereditary diffuse gastric cancer. <i>Hereditary Cancer in Clinical Practice</i> , 2007, 5, 183.	0.6	36
52	P21-Activated Kinase 1: A New Molecular Marker for Intravesical Recurrence After Transurethral Resection of Bladder Cancer. <i>Journal of Urology</i> , 2007, 178, 1073-1079.	0.2	44
53	Expression of Krüppel-like factor 5 in human gastric carcinomas. <i>Journal of Cancer Research and Clinical Oncology</i> , 2007, 134, 163-167.	1.2	42
54	Hereditary Diffuse Gastric Cancer: Diagnosis and Management. <i>Clinical Gastroenterology and Hepatology</i> , 2006, 4, 262-275.	2.4	163

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55	Genome-wide linkage screen for testicular germ cell tumour susceptibility loci. <i>Human Molecular Genetics</i> , 2006, 15, 443-451.	1.4	138
56	The Y Deletion gr/gr and Susceptibility to Testicular Germ Cell Tumor. <i>American Journal of Human Genetics</i> , 2005, 77, 1034-1043.	2.6	197
57	Prognostic Analysis of E-Cadherin Gene Promoter Hypermethylation in Patients with Surgically Resected, Node-Positive, Diffuse Gastric Cancer. <i>Clinical Cancer Research</i> , 2004, 10, 2784-2789.	3.2	83
58	N-Terminal E-Cadherin Peptides Act as Decoy Receptors for <i>Listeria monocytogenes</i> . <i>Infection and Immunity</i> , 2003, 71, 1580-1583.	1.0	19
59	Novel germline CDH1 mutations in hereditary diffuse gastric cancer families. <i>Human Mutation</i> , 2002, 19, 518-525.	1.1	63
60	Association of CDH1 haplotypes with susceptibility to sporadic diffuse gastric cancer. <i>Oncogene</i> , 2002, 21, 8192-8195.	2.6	91
61	Gastric Cancer: Inherited Predisposition. , 2002, , 253-258.		0
62	Incidence of gastric cancer and breast cancer in CDH1 (E-cadherin) mutation carriers from hereditary diffuse gastric cancer families. <i>Gastroenterology</i> , 2001, 121, 1348-1353.	0.6	579
63	Germline E-cadherin gene mutations. <i>Cancer</i> , 2001, 92, 181-187.	2.0	131
64	Hereditary diffuse gastric cancer. <i>Advances in Cancer Research</i> , 2001, 83, 55-65.	1.9	35
65	Methylation of the CDH1 promoter as the second genetic hit in hereditary diffuse gastric cancer. <i>Nature Genetics</i> , 2000, 26, 16-17.	9.4	420
66	E-cadherin downregulation in cancer: fuel on the fire?. <i>Trends in Molecular Medicine</i> , 1999, 5, 172-177.	2.6	104
67	E-cadherin germline mutations define an inherited cancer syndrome dominated by diffuse gastric cancer. <i>Human Mutation</i> , 1999, 14, 249-255.	1.1	247
68	E-cadherin unlikely to be a common 'low penetrance' gene for colorectal cancer. <i>American Journal of Medical Genetics Part A</i> , 1999, 84, 169-171.	2.4	3
69	E-cadherin germline mutations in familial gastric cancer. <i>Nature</i> , 1998, 392, 402-405.	13.7	1,542
70	Sequence Characterization of a Newly Identified Human $\hat{\pm}$ -Tubulin Gene (TUBA2). <i>Genomics</i> , 1998, 47, 125-130.	1.3	6
71	Defective myosin VIIA gene responsible for Usher syndrome type IB. <i>Nature</i> , 1995, 374, 60-61.	13.7	1,101
72	A YAC Contig and an EST Map in the Pericentromeric Region of Chromosome 13 Surrounding the Loci for Neurosensory Nonsyndromic Deafness (DFNB1 and DFNA3) and Limb-Girdle Muscular Dystrophy Type 2C (LGMD2C). <i>Genomics</i> , 1995, 29, 163-169.	1.3	25

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73	A gene responsible for a dominant form of neurosensory non-syndromic deafness maps to the NSRD1 recessive deafness gene interval. <i>Human Molecular Genetics</i> , 1994, 3, 2219-2222.	1.4	93
74	A non-syndromic form of neurosensory, recessive deafness maps to the pericentromeric region of chromosome 13q. <i>Nature Genetics</i> , 1994, 6, 24-28.	9.4	262
75	A human gene responsible for neurosensory, non-syndromic recessive deafness is a candidate homologue of the mouse sh-1 gene. <i>Human Molecular Genetics</i> , 1994, 3, 989-993.	1.4	134
76	Mutation and replacement of the 16-kDa protein gene IN RNA-1 of tobacco rattle virus. <i>Virology</i> , 1991, 182, 607-614.	1.1	16
77	Triple gene block proteins of white clover mosaic potexvirus are required for transport. <i>Virology</i> , 1991, 183, 695-702.	1.1	200
78	Organization and intervirial homologies of the coat protein gene of white clover mosaic virus. <i>Virology</i> , 1988, 162, 459-465.	1.1	17