

# Douglas R Stewart

## List of Publications by Year in descending order

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Version: 2024-02-01

80  
papers

4,084  
citations

117625

34  
h-index

128289

60  
g-index

83  
all docs

83  
docs citations

83  
times ranked

5166  
citing authors

#	ARTICLE	IF	CITATIONS
1	ACMG SF v3.0 list for reporting of secondary findings in clinical exome and genome sequencing: a policy statement of the American College of Medical Genetics and Genomics (ACMG). <i>Genetics in Medicine</i> , 2021, 23, 1381-1390.	2.4	356
2	Histopathologic evaluation of atypical neurofibromatous tumors and their transformation into malignant peripheral nerve sheath tumor in patients with neurofibromatosis 1—a consensus overview. <i>Human Pathology</i> , 2017, 67, 1-10.	2.0	275
3	<i>DICER1</i> and Associated Conditions: Identification of At-risk Individuals and Recommended Surveillance Strategies. <i>Clinical Cancer Research</i> , 2018, 24, 2251-2261.	7.0	260
4	Pleuropulmonary blastoma: A report on 350 central pathology-confirmed pleuropulmonary blastoma cases by the International Pleuropulmonary Blastoma Registry. <i>Cancer</i> , 2015, 121, 276-285.	4.1	242
5	Recommendations for reporting of secondary findings in clinical exome and genome sequencing, 2021 update: a policy statement of the American College of Medical Genetics and Genomics (ACMG). <i>Genetics in Medicine</i> , 2021, 23, 1391-1398.	2.4	145
6	Care of adults with neurofibromatosis type 1: a clinical practice resource of the American College of Medical Genetics and Genomics (ACMG). <i>Genetics in Medicine</i> , 2018, 20, 671-682.	2.4	128
7	Temporal order of RNase IIIb and loss-of-function mutations during development determines phenotype in <i>DICER1</i> syndrome: a unique variant of the two-hit tumor suppression model. <i>F1000Research</i> , 2015, 4, 214.	1.6	125
8	Glomus Tumors in Neurofibromatosis Type 1: Genetic, Functional, and Clinical Evidence of a Novel Association. <i>Cancer Research</i> , 2009, 69, 7393-7401.	0.9	122
9	Quantification of Thyroid Cancer and Multinodular Goiter Risk in the <i>DICER1</i> Syndrome: A Family-Based Cohort Study. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2017, 102, 1614-1622.	3.6	120
10	Neoplasm Risk Among Individuals With a Pathogenic Germline Variant in <i>DICER1</i> . <i>Journal of Clinical Oncology</i> , 2019, 37, 668-676.	1.6	107
11	<i>DICER1</i> -Pleuropulmonary Blastoma Familial Tumor Predisposition Syndrome. , 2014, 19, 90-100.		97
12	<i>DICER1</i> -related Sertoli-Leydig cell tumor and gynandroblastoma: Clinical and genetic findings from the International Ovarian and Testicular Stromal Tumor Registry. <i>Gynecologic Oncology</i> , 2017, 147, 521-527.	1.4	87
13	Malignant Peripheral Nerve Sheath Tumors State of the Science: Leveraging Clinical and Biological Insights into Effective Therapies. <i>Sarcoma</i> , 2017, 2017, 1-10.	1.3	84
14	Nasal chondromesenchymal hamartomas arise secondary to germline and somatic mutations of <i>DICER1</i> in the pleuropulmonary blastoma tumor predisposition disorder. <i>Human Genetics</i> , 2014, 133, 1443-1450.	3.8	82
15	Genomic and evolutionary classification of lung cancer in never smokers. <i>Nature Genetics</i> , 2021, 53, 1348-1359.	21.4	81
16	Neurofibromatosis Type 1—Associated MPNST State of the Science: Outlining a Research Agenda for the Future. <i>Journal of the National Cancer Institute</i> , 2017, 109, .	6.3	80
17	The chromosome 9q subtelomere deletion syndrome. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics</i> , 2007, 145C, 383-392.	1.6	79
18	Temporal order of RNase IIIb and loss-of-function mutations during development determines phenotype in pleuropulmonary blastoma / <i>DICER1</i> syndrome: a unique variant of the two-hit tumor suppression model. <i>F1000Research</i> , 2015, 4, 214.	1.6	78

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19	The prevalence of <i>DICER1</i> pathogenic variation in population databases. <i>International Journal of Cancer</i> , 2017, 141, 2030-2036.	5.1	75
20	Low mutation burden and frequent loss of CDKN2A/B and SMARCA2, but not PRC2, define premalignant neurofibromatosis type 1-associated atypical neurofibromas. <i>Neuro-Oncology</i> , 2019, 21, 981-992.	1.2	69
21	Variable population prevalence estimates of germline <i>TP53</i> variants: A gnomAD-based analysis. <i>Human Mutation</i> , 2019, 40, 97-105.	2.5	66
22	DICER1 tumor predisposition syndrome: an evolving story initiated with the pleuropulmonary blastoma. <i>Modern Pathology</i> , 2022, 35, 4-22.	5.5	65
23	Diagnosis, management, and complications of glomus tumours of the digits in neurofibromatosis type 1. <i>Journal of Medical Genetics</i> , 2010, 47, 525-532.	3.2	61
24	Mitotic recombination as evidence of alternative pathogenesis of gastrointestinal stromal tumours in neurofibromatosis type 1. <i>Journal of Medical Genetics</i> , 2006, 44, e61-e61.	3.2	59
25	Points to consider for reporting of germline variation in patients undergoing tumor testing: a statement of the American College of Medical Genetics and Genomics (ACMG). <i>Genetics in Medicine</i> , 2020, 22, 1142-1148.	2.4	59
26	Is Pulmonary Arterial Hypertension in Neurofibromatosis Type 1 Secondary to a Plexogenic Arteriopathy?. <i>Chest</i> , 2007, 132, 798-808.	0.8	56
27	The Cyclic AMP Pathway Is a Sex-Specific Modifier of Glioma Risk in Type I Neurofibromatosis Patients. <i>Cancer Research</i> , 2015, 75, 16-21.	0.9	56
28	Germline Cancer Predisposition Variants in Pediatric Rhabdomyosarcoma: A Report From the Children's Oncology Group. <i>Journal of the National Cancer Institute</i> , 2021, 113, 875-883.	6.3	55
29	Germline mutations in <i>Protection of Telomeres 1</i> in two families with Hodgkin lymphoma. <i>British Journal of Haematology</i> , 2018, 181, 372-377.	2.5	48
30	Evidence of polyclonality in neurofibromatosis type 2-associated multilobulated vestibular schwannomas. <i>Neuro-Oncology</i> , 2015, 17, 566-573.	1.2	45
31	Association of Inherited Pathogenic Variants in Checkpoint Kinase 2 ( <i>CHEK2</i> ) With Susceptibility to Testicular Germ Cell Tumors. <i>JAMA Oncology</i> , 2019, 5, 514.	7.1	43
32	Whole exome sequencing reveals a C-terminal germline variant in CEBPA-associated acute myeloid leukemia: 45-year follow up of a large family. <i>Haematologica</i> , 2016, 101, 846-852.	3.5	42
33	Macrocephaly associated with the DICER1 syndrome. <i>Genetics in Medicine</i> , 2017, 19, 244-248.	2.4	42
34	Higher-than-expected population prevalence of potentially pathogenic germline <i>TP53</i> variants in individuals unselected for cancer history. <i>Human Mutation</i> , 2017, 38, 1723-1730.	2.5	40
35	Advancing RAS/RASopathy therapies: An NCI-sponsored intramural and extramural collaboration for the study of RASopathies. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 866-876.	1.2	40
36	Mosaic chromosome Y loss and testicular germ cell tumor risk. <i>Journal of Human Genetics</i> , 2017, 62, 637-640.	2.3	34

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37	Management of individuals with germline variants in PALB2: a clinical practice resource of the American College of Medical Genetics and Genomics (ACMG). <i>Genetics in Medicine</i> , 2021, 23, 1416-1423.	2.4	34
38	Jaffeâ€œCampanacci syndrome, revisited: detailed clinical and molecular analyses determine whether patients have neurofibromatosis type 1, coincidental manifestations, or a distinct disorder. <i>Genetics in Medicine</i> , 2014, 16, 448-459.	2.4	33
39	Genetic Modifiers of Neurofibromatosis Type 1-Associated CafÃ©-au-Lait Macule Count Identified Using Multi-platform Analysis. <i>PLoS Genetics</i> , 2014, 10, e1004575.	3.5	31
40	Dubowitz Syndrome Is a Complex Comprised of Multiple, Genetically Distinct and Phenotypically Overlapping Disorders. <i>PLoS ONE</i> , 2014, 9, e98686.	2.5	29
41	Cephalometry in adults and children with neurofibromatosis type 1: Implications for the pathogenesis of sphenoid wing dysplasia and the â€œNF1 faciesâ€œ. <i>European Journal of Medical Genetics</i> , 2015, 58, 584-590.	1.3	29
42	Mitotic recombination of chromosome arm 17q as a cause of loss of heterozygosity of <i>NF1</i> in neurofibromatosis type 1-associated glomus tumors. <i>Genes Chromosomes and Cancer</i> , 2012, 51, 429-437.	2.8	27
43	Pathogenic Germline Variants in Cancer Susceptibility Genes in Children and Young Adults With Rhabdomyosarcoma. <i>JCO Precision Oncology</i> , 2021, 5, 75-87.	3.0	27
44	A Genome-First Approach to Characterize <i>DICER1</i> Pathogenic Variant Prevalence, Penetrance, and Phenotype. <i>JAMA Network Open</i> , 2021, 4, e210112.	5.9	25
45	Rare inactivating PDE11A variants associated with testicular germ cell tumors. <i>Endocrine-Related Cancer</i> , 2015, 22, 909-917.	3.1	24
46	Research participant interest in primary, secondary, and incidental genomic findings. <i>Genetics in Medicine</i> , 2016, 18, 1218-1225.	2.4	24
47	The prevalence of germline <i>DICER1</i> pathogenic variation in cancer populations. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2019, 7, e555.	1.2	24
48	Pleuropulmonary blastoma-like peritoneal sarcoma: a newly described malignancy associated with biallelic <i>DICER1</i> pathogenic variation. <i>Modern Pathology</i> , 2020, 33, 1922-1929.	5.5	24
49	Presacral malignant teratoid neoplasm in association with pathogenic <i>DICER1</i> variation. <i>Modern Pathology</i> , 2019, 32, 1744-1750.	5.5	22
50	Juvenile myelomonocytic leukemia due to a germline CBL Y371C mutation: 35-year follow-up of a large family. <i>Human Genetics</i> , 2015, 134, 775-787.	3.8	21
51	<i>DICER1</i> Syndrome. <i>Ophthalmology</i> , 2019, 126, 296-304.	5.2	20
52	Comparative clinical and genomic analysis of neurofibromatosis type 2-associated cranial and spinal meningiomas. <i>Scientific Reports</i> , 2020, 10, 12563.	3.3	16
53	A genotype-first approach to exploring Mendelian cardiovascular traits with clear external manifestations. <i>Genetics in Medicine</i> , 2021, 23, 94-102.	2.4	16
54	Prevalence of pathogenic/likely pathogenic variants in the 24 cancer genes of the ACMG Secondary Findings v2.0 list in a large cancer cohort and ethnicity-matched controls. <i>Genome Medicine</i> , 2018, 10, 99.	8.2	15

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55	Structural renal abnormalities in the DICER1 syndrome: a family-based cohort study. <i>Pediatric Nephrology</i> , 2018, 33, 2281-2288.	1.7	14
56	First insight into the somatic mutation burden of neurofibromatosis type 2-associated grade I and grade II meningiomas: a case report comprehensive genomic study of two cranial meningiomas with vastly different clinical presentation. <i>BMC Cancer</i> , 2017, 17, 127.	2.6	13
57	Frequency of Pathogenic Germline Variants in Cancer-Susceptibility Genes in the Childhood Cancer Survivor Study. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab007.	2.9	11
58	Incidence of Benign Meningiomas in the United States: Current and Future Trends. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab035.	2.9	11
59	Gynecologic and reproductive health in patients with pathogenic germline variants in DICER1. <i>Gynecologic Oncology</i> , 2020, 156, 647-653.	1.4	10
60	Hematologic indices in individuals with pathogenic germline <i>DICER1</i> variants. <i>Blood Advances</i> , 2021, 5, 216-223.	5.2	6
61	Endemic Burkitt Lymphoma in second-degree relatives in Northern Uganda: in-depth genome-wide analysis suggests clues about genetic susceptibility. <i>Leukemia</i> , 2021, 35, 1209-1213.	7.2	5
62	Unusual phenotypes in patients with a pathogenic germline variant in DICER1. <i>Familial Cancer</i> , 2021, , 1.	1.9	5
63	Nasal chondromesenchymal hamartomas in a cohort with pathogenic germline variation in DICER1. <i>Rhinology</i> , 2020, 3, 15-24.	0.3	5
64	Prospectively Identified Incident Testicular Cancer Risk in a Familial Testicular Cancer Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1614-1621.	2.5	4
65	Dental abnormalities in individuals with pathogenic germline variation in <i>DICER1</i> . <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 1820-1825.	1.2	4
66	Stomaching Multigene Panel Testing: What to Do About CDH1?. <i>Journal of the National Cancer Institute</i> , 2020, 112, 325-326.	6.3	4
67	Genome-wide association study of café-au-lait macule number in neurofibromatosis type 1. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2020, 8, e1400.	1.2	4
68	Testicular Sertoli cell tumour and potentially testicular Leydig cell tumour are features of <i>DICER1</i> syndrome. <i>Journal of Medical Genetics</i> , 2022, 59, 346-350.	3.2	4
69	In search of genetic factors predisposing to familial hairy cell leukemia (HCL): exome-sequencing of four multiplex HCL pedigrees. <i>Leukemia</i> , 2020, 34, 1934-1938.	7.2	3
70	Quantification of Discordant Variant Interpretations in a Large Family-Based Study of Li-Fraumeni Syndrome. <i>JCO Precision Oncology</i> , 2021, 5, 1727-1737.	3.0	3
71	Identification of Genetic Risk Factors for Familial Urinary Bladder Cancer: An Exome Sequencing Study. <i>JCO Precision Oncology</i> , 2021, 5, 1830-1839.	3.0	3
72	Comment on: <i>DICER1</i> -Negative Pleuropulmonary Blastoma in a Patient With Selective IgA Deficiency. <i>Pediatric Blood and Cancer</i> , 2016, 63, 1869-1870.	1.5	2

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73	Response to: Concern regarding classification of germline TP53 variants as likely pathogenic. Human Mutation, 2019, 40, 832-833.	2.5	1
74	Novel MAPK/AKT-impairing germline NRAS variant identified in a melanoma-prone family. Familial Cancer, 2022, 21, 347-355.	1.9	1
75	Another Weak Link in the Electron Transport Chain: Variants in SUCLG2 and Risk for Pheochromocytoma and Paraganglioma. Journal of the National Cancer Institute, 2021, , .	6.3	1
76	Response to Hannah-Shmouni and Stratakis. Genetics in Medicine, 2019, 21, 1256.	2.4	0
77	Lack of pathogenic germline DICER1 variants in males with testicular germ-cell tumors. Cancer Genetics, 2020, 248-249, 49-56.	0.4	0
78	Reply: pleuropulmonary blastoma-like peritoneal sarcoma and DICER1-associated sarcomas: toward a unified nomenclature. Modern Pathology, 2021, 34, 1229-1230.	5.5	0
79	Whole exome sequencing and copy-number variation analysis of 20 NF2-associated spinal and cranial meningiomas.. Journal of Clinical Oncology, 2017, 35, 2051-2051.	1.6	0
80	Insights Into Immune-Mediated Disease and Cancer Risk”Delivering on the Promise of UK Biobank Big Data. JAMA Oncology, 2021, , .	7.1	0