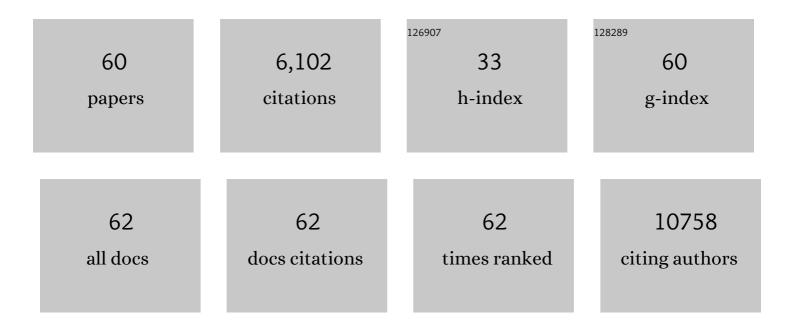
Hilda A Pickett

List of Publications by Year in descending order

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HILDA A DICKETT

#	Article	IF	CITATIONS
1	Whole-genome landscapes of major melanoma subtypes. Nature, 2017, 545, 175-180.	27.8	1,068
2	Whole-genome landscape of pancreatic neuroendocrine tumours. Nature, 2017, 543, 65-71.	27.8	716
3	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. Nature Genetics, 2013, 45, 371-384.	21.4	493
4	DNA C-circles are specific and quantifiable markers of alternative-lengthening-of-telomeres activity. Nature Biotechnology, 2009, 27, 1181-1185.	17.5	406
5	The Transcriptional and Functional Properties of Mouse Epiblast Stem Cells Resemble the Anterior Primitive Streak. Cell Stem Cell, 2014, 14, 107-120.	11.1	255
6	Spontaneous occurrence of telomeric DNA damage response in the absence of chromosome fusions. Nature Structural and Molecular Biology, 2009, 16, 1244-1251.	8.2	220
7	Control of telomere length by a trimming mechanism that involves generation of t-circles. EMBO Journal, 2009, 28, 799-809.	7.8	178
8	Molecular mechanisms of activity and derepression of alternative lengthening of telomeres. Nature Structural and Molecular Biology, 2015, 22, 875-880.	8.2	174
9	Telomerase activity is associated with an increase in DNA methylation at the proximal subtelomere and a reduction in telomeric transcription. Nucleic Acids Research, 2009, 37, 1152-1159.	14.5	137
10	Alternative Lengthening of Telomeres: DNA Repair Pathways Converge. Trends in Genetics, 2017, 33, 921-932.	6.7	132
11	Variant repeats are interspersed throughout the telomeres and recruit nuclear receptors in ALT cells. Journal of Cell Biology, 2012, 199, 893-906.	5.2	129
12	BLM and SLX4 play opposing roles in recombinationâ€dependent replication at human telomeres. EMBO Journal, 2017, 36, 2907-2919.	7.8	127
13	The FANCM-BLM-TOP3A-RMI complex suppresses alternative lengthening of telomeres (ALT). Nature Communications, 2019, 10, 2252.	12.8	125
14	Molecular characterization of inter-telomere and intra-telomere mutations in human ALT cells. Nature Genetics, 2002, 30, 301-305.	21.4	117
15	Alternative lengthening of telomeres in normal mammalian somatic cells. Genes and Development, 2013, 27, 18-23.	5.9	111
16	Telomere Loop Dynamics in Chromosome End Protection. Molecular Cell, 2018, 71, 510-525.e6.	9.7	102
17	Telomerase Recruitment Requires both TCAB1 and Cajal Bodies Independently. Molecular and Cellular Biology, 2012, 32, 2384-2395.	2.3	101
18	NuRD–ZNF827 recruitment to telomeres creates a molecular scaffold for homologous recombination. Nature Structural and Molecular Biology, 2014, 21, 760-770.	8.2	101

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19	Inherited bone marrow failure associated with germline mutation of ACD, the gene encoding telomere protein TPP1. Blood, 2014, 124, 2767-2774.	1.4	97
20	Telomere extension by telomerase and ALT generates variant repeats by mechanistically distinct processes. Nucleic Acids Research, 2014, 42, 1733-1746.	14.5	92
21	Normal mammalian cells negatively regulate telomere length by telomere trimming. Human Molecular Genetics, 2011, 20, 4684-4692.	2.9	87
22	Histone variant H3.3 provides the heterochromatic H3 lysine 9 tri-methylation mark at telomeres. Nucleic Acids Research, 2015, 43, gkv847.	14.5	79
23	Alternative lengthening of telomeres: remodeling the telomere architecture. Frontiers in Oncology, 2013, 3, 27.	2.8	78
24	Extensive Proliferation of Human Cancer Cells with Ever-Shorter Telomeres. Cell Reports, 2017, 19, 2544-2556.	6.4	75
25	FANCM suppresses DNA replication stress at ALT telomeres by disrupting TERRA R-loops. Scientific Reports, 2019, 9, 19110.	3.3	73
26	Targeting telomeres: advances in telomere maintenance mechanism-specific cancer therapies. Nature Reviews Cancer, 2022, 22, 515-532.	28.4	68
27	Folate Deficiency Induces Dysfunctional Long and Short Telomeres; Both States Are Associated with Hypomethylation and DNA Damage in Human WIL2-NS Cells. Cancer Prevention Research, 2014, 7, 128-138.	1.5	59
28	The role of telomere trimming in normal telomere length dynamics. Cell Cycle, 2012, 11, 1309-1315.	2.6	53
29	Functional Polymorphisms in the TERT Promoter Are Associated with Risk of Serous Epithelial Ovarian and Breast Cancers. PLoS ONE, 2011, 6, e24987.	2.5	48
30	The Medical Genome Reference Bank contains whole genome and phenotype data of 2570 healthy elderly. Nature Communications, 2020, 11, 435.	12.8	47
31	Mechanisms that drive telomere maintenance and recombination in human cancers. Current Opinion in Genetics and Development, 2020, 60, 25-30.	3.3	45
32	Comparative analysis of whole genome sequencing-based telomere length measurement techniques. Methods, 2017, 114, 4-15.	3.8	43
33	Telomere Length Measurement by Molecular Combing. Frontiers in Cell and Developmental Biology, 2020, 8, 493.	3.7	41
34	Telomere sequence content can be used to determine ALT activity in tumours. Nucleic Acids Research, 2018, 46, 4903-4918.	14.5	40
35	<i>MYC</i> -Driven Neuroblastomas Are Addicted to a Telomerase-Independent Function of Dyskerin. Cancer Research, 2016, 76, 3604-3617.	0.9	38
36	Human telomeres that contain (CTAGGG)n repeats show replication dependent instability in somatic cells and the male germline. Nucleic Acids Research, 2009, 37, 6225-6238.	14.5	37

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37	A Common Cancer Risk-Associated Allele in the hTERT Locus Encodes a Dominant Negative Inhibitor of Telomerase. PLoS Genetics, 2015, 11, e1005286.	3.5	35
38	Telomerase promotes formation of a telomere protective complex in cancer cells. Science Advances, 2019, 5, eaav4409.	10.3	27
39	Functional dissection of breast cancer risk-associated <i>TERT</i> promoter variants. Oncotarget, 2017, 8, 67203-67217.	1.8	21
40	Enhanced cardiac repair by telomerase reverse transcriptase over-expression in human cardiac mesenchymal stromal cells. Scientific Reports, 2019, 9, 10579.	3.3	21
41	Targeted Therapy of <i>TERT</i> -Rearranged Neuroblastoma with BET Bromodomain Inhibitor and Proteasome Inhibitor Combination Therapy. Clinical Cancer Research, 2021, 27, 1438-1451.	7.0	20
42	Telomeric replication stress: the beginning and the end for alternative lengthening of telomeres cancers. Open Biology, 2022, 12, 220011.	3.6	20
43	Amplification of Telomerase Reverse Transcriptase Gene in Human Mammary Epithelial Cells with Limiting Telomerase RNA Expression Levels. Cancer Research, 2008, 68, 3115-3123.	0.9	18
44	Aurora Kinase B, a novel regulator of TERF1 binding and telomeric integrity. Nucleic Acids Research, 2017, 45, 12340-12353.	14.5	18
45	Progerin impairs 3D genome organization and induces fragile telomeres by limiting the dNTP pools. Scientific Reports, 2021, 11, 13195.	3.3	18
46	Efficient inÂvivo editing of OTC-deficient patient-derived primary human hepatocytes. JHEP Reports, 2020, 2, 100065.	4.9	18
47	<i>DKC1</i> is a transcriptional target of GATA1 and drives upregulation of telomerase activity in normal human erythroblasts. Haematologica, 2020, 105, 1517-1526.	3.5	15
48	Anti-recombination function of MutSα restricts telomere extension by ALT-associated homology-directed repair. Cell Reports, 2021, 37, 110088.	6.4	15
49	Structural and functional characterization of the RBBP4–ZNF827 interaction and its role in NuRD recruitment to telomeres. Biochemical Journal, 2018, 475, 2667-2679.	3.7	13
50	TELO-SCOPE study: a randomised, double-blind, placebo-controlled, phase 2 trial of danazol for short telomere related pulmonary fibrosis. BMJ Open Respiratory Research, 2021, 8, e001127.	3.0	13
51	Platelet-Derived Growth Factor Receptor-Alpha Expressing Cardiac Progenitor Cells Can Be Derived from Previously Cryopreserved Human Heart Samples. Stem Cells and Development, 2018, 27, 184-198.	2.1	12
52	Telomeres and stress: Promising avenues for research in psycho-oncology. Asia-Pacific Journal of Oncology Nursing, 2016, 3, 137-147.	1.6	10
53	qmotif: determination of telomere content from whole-genome sequence data. Bioinformatics Advances, 2022, 2, .	2.4	5
54	Loss-of-function variants in <i>POT1</i> predispose to uveal melanoma. Journal of Medical Genetics, 2021, 58, 234-236.	3.2	3

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#	Article	IF	CITATIONS
55	Functional interaction between compound heterozygous <i>TERT</i> mutations causes severe telomere biology disorder. Blood Advances, 2022, 6, 3779-3791.	5.2	2
56	A novel cause of DKC1 â€related bone marrow failure: Partial deletion of the 3′ untranslated region. EJHaem, 2021, 2, 157-166.	1.0	1
57	Age Adjusted Telomere Length Decreases Following Treatment for Pediatric Acute Lymphoblastic Leukemia, but Does Not Predict Toxicity. Blood, 2015, 126, 4984-4984.	1.4	1
58	Age-Adjusted Telomere Length May Predict Treatment-Related Mortality in Children Undergoing Hematopoietic Stem Cell Transplantation. Blood, 2015, 126, 5445-5445.	1.4	1
59	A DNA-fiber protocol for single molecule analysis of telomere (SMAT) length and extension events in cancer cells. STAR Protocols, 2022, 3, 101212.	1.2	1
60	Dyskerin Is Upregulated During Erythroid Differentiation of Human Hematopoietic Progenitor Cells and Hyperactivates Telomerase in Erythroid Precursor Cells. Blood, 2012, 120, 980-980.	1.4	0