

Duncan M Baird

List of Publications by Year in descending order

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Version: 2024-02-01

106
papers

5,263
citations

76196

40
h-index

91712

69
g-index

111
all docs

111
docs citations

111
times ranked

7899
citing authors

#	ARTICLE	IF	CITATIONS
1	Extensive allelic variation and ultrashort telomeres in senescent human cells. <i>Nature Genetics</i> , 2003, 33, 203-207.	9.4	469
2	Identification and Clonal Characterisation of a Progenitor Cell Sub-Population in Normal Human Articular Cartilage. <i>PLoS ONE</i> , 2010, 5, e13246.	1.1	338
3	The nature of telomere fusion and a definition of the critical telomere length in human cells. <i>Genes and Development</i> , 2007, 21, 2495-2508.	2.7	237
4	Two subsets of stem-like CD8+ memory T cell progenitors with distinct fate commitments in humans. <i>Nature Immunology</i> , 2020, 21, 1552-1562.	7.0	167
5	Fibroblast Dysfunction Is a Key Factor in the Non-Healing of Chronic Venous Leg Ulcers. <i>Journal of Investigative Dermatology</i> , 2008, 128, 2526-2540.	0.3	166
6	Measuring telomere length and telomere dynamics in evolutionary biology and ecology. <i>Methods in Ecology and Evolution</i> , 2014, 5, 299-310.	2.2	158
7	Telomere dysfunction and fusion during the progression of chronic lymphocytic leukemia: evidence for a telomere crisis. <i>Blood</i> , 2010, 116, 1899-1907.	0.6	148
8	Telomere instability in the male germline. <i>Human Molecular Genetics</i> , 2006, 15, 45-51.	1.4	141
9	Reproducibility of telomere length assessment: an international collaborative study. <i>International Journal of Epidemiology</i> , 2015, 44, 1673-1683.	0.9	133
10	Telomerase Reverse Transcriptase Locus Polymorphisms and Cancer Risk: A Field Synopsis and Meta-Analysis. <i>Journal of the National Cancer Institute</i> , 2012, 104, 840-854.	3.0	119
11	Signalling of DNA damage and cytokines across cell barriers exposed to nanoparticles depends on barrier thickness. <i>Nature Nanotechnology</i> , 2011, 6, 824-833.	15.6	114
12	Blood Cell Telomere Length Is a Dynamic Feature. <i>PLoS ONE</i> , 2011, 6, e21485.	1.1	111
13	Structural stability and chromosome-specific telomere length is governed by cis-acting determinants in humans. <i>Human Molecular Genetics</i> , 2006, 15, 725-733.	1.4	110
14	Variation at the <i>TERT</i> locus and predisposition for cancer. <i>Expert Reviews in Molecular Medicine</i> , 2010, 12, e16.	1.6	101
15	Mechanisms underlying telomere repeat turnover, revealed by hypervariable variant repeat distribution patterns in the human Xp/Yp telomere. <i>EMBO Journal</i> , 1995, 14, 5433-5443.	3.5	92
16	New developments in telomere length analysis. <i>Experimental Gerontology</i> , 2005, 40, 363-368.	1.2	85
17	Prevention of Accelerated Cell Aging in Werner Syndrome Using a p38 Mitogen-Activated Protein Kinase Inhibitor. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2005, 60, 1386-1393.	1.7	84
18	Structural and functional analysis of the human POT1-TPP1 telomeric complex. <i>Nature Communications</i> , 2017, 8, 14928.	5.8	84

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19	Extensive Telomere Erosion in the Initiation of Colorectal Adenomas and Its Association With Chromosomal Instability. <i>Journal of the National Cancer Institute</i> , 2013, 105, 1202-1211.	3.0	81
20	Fusion of short telomeres in human cells is characterized by extensive deletion and microhomology, and can result in complex rearrangements. <i>Nucleic Acids Research</i> , 2010, 38, 1841-1852.	6.5	78
21	Characterisation of a divergent progenitor cell sub-populations in human osteoarthritic cartilage: the role of telomere erosion and replicative senescence. <i>Scientific Reports</i> , 2017, 7, 41421.	1.6	78
22	High Levels of Sequence Polymorphism and Linkage Disequilibrium at the Telomere of 12q: Implications for Telomere Biology and Human Evolution. <i>American Journal of Human Genetics</i> , 2000, 66, 235-250.	2.6	77
23	Telomere dysfunction accurately predicts clinical outcome in chronic lymphocytic leukaemia, even in patients with early stage disease. <i>British Journal of Haematology</i> , 2014, 167, 214-223.	1.2	73
24	CCR8 Expression Defines Tissue-Resident Memory T Cells in Human Skin. <i>Journal of Immunology</i> , 2018, 200, 1639-1650.	0.4	71
25	A subterminal satellite located adjacent to telomeres in chimpanzees is absent from the human genome. <i>Nature Genetics</i> , 1994, 6, 52-56.	9.4	69
26	Genotoxic effects of particles of surgical cobalt chrome alloy on human cells of different age in vitro. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2007, 619, 45-58.	0.4	69
27	Human Stem Cell-like Memory T Cells Are Maintained in a State of Dynamic Flux. <i>Cell Reports</i> , 2016, 17, 2811-2818.	2.9	67
28	Escape from Telomere-Driven Crisis Is DNA Ligase III Dependent. <i>Cell Reports</i> , 2014, 8, 1063-1076.	2.9	65
29	Mapping H4K20me3 onto the chromatin landscape of senescent cells indicates a function in control of cell senescence and tumor suppression through preservation of genetic and epigenetic stability. <i>Genome Biology</i> , 2016, 17, 158.	3.8	65
30	Telomere dynamics in human cells. <i>Biochimie</i> , 2008, 90, 116-121.	1.3	54
31	The Plasticity of Human Telomeres Demonstrated by a Hypervariable Telomere Repeat Array That Is Located on Some Copies of 16p and 16q. <i>Human Molecular Genetics</i> , 1999, 8, 1637-1646.	1.4	52
32	Telomere dysfunction and its role in haematological cancer. <i>British Journal of Haematology</i> , 2012, 156, 573-587.	1.2	51
33	Short telomeres are preferentially elongated by telomerase in human cells. <i>FEBS Letters</i> , 2009, 583, 3076-3080.	1.3	49
34	TERT promoter mutation in adult granulosa cell tumor of the ovary. <i>Modern Pathology</i> , 2018, 31, 1107-1115.	2.9	49
35	Telomere Length Dynamics and the Evolution of Cancer Genome Architecture. <i>International Journal of Molecular Sciences</i> , 2018, 19, 482.	1.8	48
36	Normal telomere erosion rates at the single cell level in Werner syndrome fibroblast cells. <i>Human Molecular Genetics</i> , 2004, 13, 1515-1524.	1.4	47

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37	Chromothripsis during telomere crisis is independent of NHEJ, and consistent with a replicative origin. <i>Genome Research</i> , 2019, 29, 737-749.	2.4	47
38	Human TSCM cell dynamics in vivo are compatible with long-lived immunological memory and stemness. <i>PLoS Biology</i> , 2018, 16, e2005523.	2.6	46
39	The Extent and Significance of Telomere Loss with Age. <i>Annals of the New York Academy of Sciences</i> , 2004, 1019, 265-268.	1.8	44
40	Telomere length predicts progression and overall survival in chronic lymphocytic leukemia: data from the UK LRF CLL4 trial. <i>Leukemia</i> , 2015, 29, 2411-2414.	3.3	42
41	Mechanisms of telomeric instability. <i>Cytogenetic and Genome Research</i> , 2008, 122, 308-314.	0.6	41
42	Extreme telomere erosion in ATM-mutated and 11q-deleted CLL patients is independent of disease stage. <i>Leukemia</i> , 2012, 26, 826-830.	3.3	39
43	Salivary Gland Stem Cells Age Prematurely in Primary Sjögren's Syndrome. <i>Arthritis and Rheumatology</i> , 2019, 71, 133-142.	2.9	39
44	Sister chromatid telomere fusions, but not NHEJ-mediated inter-chromosomal telomere fusions, occur independently of DNA ligases 3 and 4. <i>Genome Research</i> , 2016, 26, 588-600.	2.4	38
45	Telomeres and genomic evolution. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2018, 373, 20160437.	1.8	35
46	Telomere fusion threshold identifies a poor prognostic subset of breast cancer patients. <i>Molecular Oncology</i> , 2015, 9, 1186-1193.	2.1	34
47	Catastrophic Endgames: Emerging Mechanisms of Telomere-Driven Genomic Instability. <i>Trends in Genetics</i> , 2020, 36, 347-359.	2.9	34
48	Effects of hTERT on metal ion-induced genomic instability. <i>Oncogene</i> , 2006, 25, 3424-3435.	2.6	32
49	UPF1 promotes the formation of R loops to stimulate DNA double-strand break repair. <i>Nature Communications</i> , 2021, 12, 3849.	5.8	32
50	Telomeres. <i>Experimental Gerontology</i> , 2006, 41, 1223-1227.	1.2	31
51	Telomere length maintenance in stem cell populations. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2009, 1792, 324-328.	1.8	31
52	Functional Effector Memory T Cells Enrich the Peritoneal Cavity of Patients Treated with Peritoneal Dialysis. <i>Journal of the American Society of Nephrology: JASN</i> , 2009, 20, 1895-1900.	3.0	29
53	Strain-specific telomere length revealed by single telomere length analysis in <i>Caenorhabditis elegans</i> . <i>Nucleic Acids Research</i> , 2004, 32, 3383-3391.	6.5	27
54	DNA damaging bystander signalling from stem cells, cancer cells and fibroblasts after Cr(VI) exposure and its dependence on telomerase. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2010, 683, 1-8.	0.4	26

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55	Differential TERT promoter methylation and response to 5-azadeoxycytidine in acute myeloid leukemia cell lines: TERT expression, telomerase activity, telomere length, and cell death. <i>Genes Chromosomes and Cancer</i> , 2012, 51, 768-780.	1.5	26
56	Telomere length is a critical determinant for survival in multiple myeloma. <i>British Journal of Haematology</i> , 2017, 178, 94-98.	1.2	26
57	Increased Oral Fibroblast Lifespan Is Telomerase-independent. <i>Journal of Dental Research</i> , 2009, 88, 916-921.	2.5	25
58	Chromosomal instability and telomere lengths of each chromosomal arm measured by Q-FISH in human fibroblast strains prior to replicative senescence. <i>Mechanisms of Ageing and Development</i> , 2010, 131, 614-624.	2.2	25
59	The C-Terminal Extension Unique to the Long Isoform of the Shelterin Component TIN2 Enhances Its Interaction with TRF2 in a Phosphorylation- and Dyskeratosis Congenita Cluster-Dependent Fashion. <i>Molecular and Cellular Biology</i> , 2018, 38, .	1.1	25
60	Sequences from higher primates orthologous to the human Xp/Yp telomere junction region reveal gross rearrangements and high levels of divergence. <i>Human Molecular Genetics</i> , 1997, 6, 2291-2299.	1.4	22
61	Telomere dynamics during replicative senescence are not directly modulated by conditions of oxidative stress in IMR90 fibroblast cells. <i>Biogerontology</i> , 2009, 10, 683-693.	2.0	22
62	Telomere length is an independent prognostic marker in <sc>MDS</sc> but not in <i>de novo</i> <sc>AML</sc>. <i>British Journal of Haematology</i> , 2017, 178, 240-249.	1.2	21
63	Telomere instability detected in sporadic colon cancers, some showing mutations in a mismatch repair gene. <i>Oncogene</i> , 2004, 23, 3434-3443.	2.6	20
64	Telomere length heterogeneity in placenta revealed with high-resolution telomere length analysis. <i>Placenta</i> , 2017, 59, 61-68.	0.7	20
65	Mre11 modulates the fidelity of fusion between short telomeres in human cells. <i>Nucleic Acids Research</i> , 2012, 40, 2518-2526.	6.5	19
66	DNA Ligase 1 is an essential mediator of sister chromatid telomere fusions in G2 cell cycle phase. <i>Nucleic Acids Research</i> , 2019, 47, 2402-2424.	6.5	19
67	CD57+ Memory T Cells Proliferate In Vivo. <i>Cell Reports</i> , 2020, 33, 108501.	2.9	18
68	Integrative analysis of spontaneous CLL regression highlights genetic and microenvironmental interdependency in CLL. <i>Blood</i> , 2020, 135, 411-428.	0.6	17
69	Highly purified CD38 ⁺ subpopulations show no evidence of preferential clonal evolution despite having increased proliferative activity when compared with CD38 ⁺ subpopulations derived from the same chronic lymphocytic leukaemia patient. <i>British Journal of Haematology</i> , 2008, 142, 595-605.	1.2	15
70	Metformin and insulin treatment prevent placental telomere attrition in boys exposed to maternal diabetes. <i>PLoS ONE</i> , 2018, 13, e0208533.	1.1	15
71	PARP1 is required for preserving telomeric integrity but is dispensable for A-NHEJ. <i>Oncotarget</i> , 2018, 9, 34821-34837.	0.8	14
72	CGGBP1 phosphorylation constitutes a telomere-protection signal. <i>Cell Cycle</i> , 2014, 13, 96-105.	1.3	13

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73	Dysgu: efficient structural variant calling using short or long reads. <i>Nucleic Acids Research</i> , 2022, 50, e53-e53.	6.5	13
74	Telomere length predicts for outcome to FCR chemotherapy in CLL. <i>Leukemia</i> , 2019, 33, 1953-1963.	3.3	12
75	High-throughput STELA provides a rapid test for the diagnosis of telomere biology disorders. <i>Human Genetics</i> , 2021, 140, 945-955.	1.8	12
76	Telomeres II. <i>Experimental Gerontology</i> , 2007, 43, 15-9.	1.2	10
77	Human Endometrial Carcinogenesis Is Associated with Significant Reduction in Long Non-Coding RNA, TERRA. <i>International Journal of Molecular Sciences</i> , 2020, 21, 8686.	1.8	10
78	Telomere fusions associate with coding sequence and copy number alterations in CLL. <i>Leukemia</i> , 2019, 33, 2093-2097.	3.3	9
79	DNA damage signalling from the placenta to foetal blood as a potential mechanism for childhood leukaemia initiation. <i>Scientific Reports</i> , 2019, 9, 4370.	1.6	9
80	Reproducibility of telomere length assessment: Authors'™ Response to Damjan Krstajic and Ljubomir Buturovic. <i>International Journal of Epidemiology</i> , 2015, 44, 1739-1741.	0.9	8
81	Is Southern blotting necessary to measure telomere length reproducibly? Authors'™ Response to: Commentary: The reliability of telomere length measurements. <i>International Journal of Epidemiology</i> , 2015, 44, 1686-1687.	0.9	8
82	Telomere erosion in NF1 tumorigenesis. <i>Oncotarget</i> , 2017, 8, 40132-40139.	0.8	8
83	Intra-allelic mutation at human telomeres. <i>Biochemical Society Transactions</i> , 2006, 34, 581-582.	1.6	7
84	Telomere length profiles in primary human peritoneal mesothelial cells are consistent with senescence. <i>Mechanisms of Ageing and Development</i> , 2017, 164, 37-40.	2.2	7
85	Extensive telomere erosion is consistent with localised clonal expansions in Barrett's™ metaplasia. <i>PLoS ONE</i> , 2017, 12, e0174833.	1.1	6
86	Shortened telomeres: a driving force behind leukemia?. <i>Future Oncology</i> , 2010, 6, 1681-1686.	1.1	5
87	Telomeres and Chromosomal Translocations. <i>Advances in Experimental Medicine and Biology</i> , 2018, 1044, 89-112.	0.8	5
88	Tracking telomere fusions through crisis reveals conflict between DNA transcription and the DNA damage response. <i>NAR Cancer</i> , 2021, 3, zcaa044.	1.6	5
89	Symptoms of Prenatal Depression Associated with Shorter Telomeres in Female Placenta. <i>International Journal of Molecular Sciences</i> , 2021, 22, 7458.	1.8	5
90	Telomere analysis to predict chronic lymphocytic leukemia outcome: a STELA test to change clinical practice?. <i>Expert Review of Hematology</i> , 2014, 7, 701-703.	1.0	4

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91	Combined analysis of IGHV mutations, telomere length and CD49d identifies long-term progression-free survivors in TP53 wild-type CLL treated with FCR-based therapies. <i>Leukemia</i> , 2022, 36, 271-274.	3.3	4
92	PARP inhibition prevents escape from a telomere-driven crisis and inhibits cell immortalisation. <i>Oncotarget</i> , 2018, 9, 37549-37563.	0.8	4
93	BCL-3 loss sensitises colorectal cancer cells to DNA damage by targeting homologous recombination. <i>DNA Repair</i> , 2022, 115, 103331.	1.3	3
94	POLQ suppresses genome instability and alterations in DNA repeat tract lengths. <i>NAR Cancer</i> , 2022, 4, .	1.6	3
95	Alternative end joining, clonal evolution, and escape from a telomere-driven crisis. <i>Molecular and Cellular Oncology</i> , 2015, 2, e975623.	0.3	2
96	In Vitro Co-Culture of CLL-B Cells Reveals Long-Term Survival, Proliferation, and Maintenance of Telomere Length. <i>Blood</i> , 2016, 128, 350-350.	0.6	1
97	Everything your supervisor should have told you but you were too afraid to ask!. <i>Trends in Biotechnology</i> , 1999, 17, 174.	4.9	0
98	Translating cancer genetics into mechanism-based drug discovery. <i>Toxicology</i> , 2006, 226, 15-16.	2.0	0
99	4.24 Telomere Dysfunction is the Critical Determinant of Clinical Outcome in CLL. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2011, 11, S234-S235.	0.2	0
100	CBMT-21. TERT PROMOTER-MUTANT GLIOBLASTOMAS EXHIBIT DEPENDENCY ON TELOMERASE. <i>Neuro-Oncology</i> , 2019, 21, vi37-vi37.	0.6	0
101	Editorial overview: The instability of the cancer genome: it starts at the end. <i>Current Opinion in Genetics and Development</i> , 2020, 60, iii-vi.	1.5	0
102	Longitudinal Analysis Reveals Telomere Length Maintenance In CLL B-Cells But Marked Erosion In CLL Patient T-Cells. <i>Blood</i> , 2013, 122, 1617-1617.	0.6	0
103	CLL Is Associated with Development of Subclinical Cytomegalovirus Viraemia and Accumulation of Large Populations of "exhausted" CMV-Specific CD4+ T Cells. <i>Blood</i> , 2014, 124, 3290-3290.	0.6	0
104	Telomere Length Is Associated with Epigenetic Programming in CLL and Is a Superior Predictor of Clinical Outcome with the Ability to Bifurcate Patients with the Same CLL-IPI Score. <i>Blood</i> , 2018, 132, 1833-1833.	0.6	0
105	Telomere Length Predicts for Outcome to FCR Chemoimmunotherapy in CLL. <i>Blood</i> , 2018, 132, 1854-1854.	0.6	0
106	Telomere Length and CD49d Cooperate with IGHV Gene Status As Predictors of Long-Term Progression-Free Survival in CLL Patients Treated with FCR-Based Regimens. <i>Blood</i> , 2020, 136, 46-47.	0.6	0