

# Keith R Bradnam

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4254092/publications.pdf>

Version: 2024-02-01

16  
papers

5,181  
citations

643344

15  
h-index

1113639

15  
g-index

17  
all docs

17  
docs citations

17  
times ranked

12808  
citing authors

#	ARTICLE	IF	CITATIONS
1	Catastrophic chromosomal restructuring during genome elimination in plants. <i>ELife</i> , 2015, 4, .	2.8	104
2	Comparative analysis of tandem repeats from hundreds of species reveals unique insights into centromere evolution. <i>Genome Biology</i> , 2013, 14, R10.	13.9	397
3	Assemblathon 2: evaluating de novo methods of genome assembly in three vertebrate species. <i>GigaScience</i> , 2013, 2, 10.	3.3	582
4	Evidence for a DNA-Based Mechanism of Intron-Mediated Enhancement. <i>Frontiers in Plant Science</i> , 2011, 2, 98.	1.7	60
5	Myc and Miz-1 have coordinate genomic functions including targeting Hox genes in human embryonic stem cells. <i>Epigenetics and Chromatin</i> , 2011, 4, 20.	1.8	30
6	Assemblathon 1: A competitive assessment of de novo short read assembly methods. <i>Genome Research</i> , 2011, 21, 2224-2241.	2.4	443
7	Comparative and functional analysis of intron-mediated enhancement signals reveals conserved features among plants. <i>Nucleic Acids Research</i> , 2011, 39, 5328-5337.	6.5	136
8	Assessing the gene space in draft genomes. <i>Nucleic Acids Research</i> , 2009, 37, 289-297.	6.5	395
9	Longer First Introns Are a General Property of Eukaryotic Gene Structure. <i>PLoS ONE</i> , 2008, 3, e3093.	1.1	117
10	CEGMA: a pipeline to accurately annotate core genes in eukaryotic genomes. <i>Bioinformatics</i> , 2007, 23, 1061-1067.	1.8	1,979
11	WormBase: a comprehensive data resource for <i>Caenorhabditis</i> biology and genomics. <i>Nucleic Acids Research</i> , 2004, 33, D383-D389.	6.5	155
12	WormBase as an Integrated Platform for the <i>C. elegans</i> ORFeome. <i>Genome Research</i> , 2004, 14, 2155-2161.	2.4	19
13	WormBase: a multi-species resource for nematode biology and genomics. <i>Nucleic Acids Research</i> , 2004, 32, 411D-417.	6.5	610
14	WormBase: a cross-species database for comparative genomics. <i>Nucleic Acids Research</i> , 2003, 31, 133-137.	6.5	107
15	The UK Crop Plant Bioinformatics Network (UK CropNet). <i>Yeast</i> , 2000, 1, 335-338.	0.8	0
16	G+C content variation along and among <i>Saccharomyces cerevisiae</i> chromosomes. <i>Molecular Biology and Evolution</i> , 1999, 16, 666-675.	3.5	45