

# Adam Woolfe

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12069129/publications.pdf>

Version: 2024-02-01

10  
papers

1,775  
citations

1040056

9  
h-index

1372567

10  
g-index

10  
all docs

10  
docs citations

10  
times ranked

2973  
citing authors

#	ARTICLE	IF	CITATIONS
1	Mislocalization of the Centromeric Histone Variant CenH3/CENP-A in Human Cells Depends on the Chaperone DAXX. <i>Molecular Cell</i> , 2014, 53, 631-644.	9.7	214
2	Dynamics of Histone H3 Deposition In Vivo Reveal a Nucleosome Gap-Filling Mechanism for H3.3 to Maintain Chromatin Integrity. <i>Molecular Cell</i> , 2011, 44, 928-941.	9.7	329
3	Early Evolution of Conserved Regulatory Sequences Associated with Development in Vertebrates. <i>PLoS Genetics</i> , 2009, 5, e1000762.	3.5	82
4	Chapter 12 Organization of Conserved Elements Near Key Developmental Regulators in Vertebrate Genomes. <i>Advances in Genetics</i> , 2008, 61, 307-338.	1.8	29
5	Comparative genomics using Fugu reveals insights into regulatory subfunctionalization. <i>Genome Biology</i> , 2007, 8, R53.	9.6	66
6	CONDOR: a database resource of developmentally associated conserved non-coding elements. <i>BMC Developmental Biology</i> , 2007, 7, 100.	2.1	60
7	Characterisation of conserved non-coding sequences in vertebrate genomes using bioinformatics, statistics and functional studies. <i>Comparative Biochemistry and Physiology Part D: Genomics and Proteomics</i> , 2006, 1, 46-58.	1.0	5
8	Defining a genomic radius for long-range enhancer action: duplicated conserved non-coding elements hold the key. <i>Trends in Genetics</i> , 2006, 22, 5-10.	6.7	69
9	Ancient duplicated conserved noncoding elements in vertebrates: A genomic and functional analysis. <i>Genome Research</i> , 2006, 16, 451-465.	5.5	88
10	Highly Conserved Non-Coding Sequences Are Associated with Vertebrate Development. <i>PLoS Biology</i> , 2004, 3, e7.	5.6	833