

# Michael Pheasant

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2289861/publications.pdf>

Version: 2024-02-01

15  
papers

3,742  
citations

687363

13  
h-index

1058476

14  
g-index

15  
all docs

15  
docs citations

15  
times ranked

6771  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomics Virtual Laboratory: A Practical Bioinformatics Workbench for the Cloud. PLoS ONE, 2015, 10, e0140829.	2.5	119
2	Quality assurance study of real-time targeted massive parallel sequencing (MPS) comprehensive cancer panel (CCP) and network pathway analysis in early breast cancer (EBC).. Journal of Clinical Oncology, 2014, 32, 11092-11092.	1.6	0
3	Comparison of Ultra-Conserved Elements in Drosophilids and Vertebrates. PLoS ONE, 2013, 8, e82362.	2.5	14
4	The UCSC Genome Browser database: update 2010. Nucleic Acids Research, 2010, 38, D613-D619.	14.5	537
5	ENCODE whole-genome data in the UCSC Genome Browser. Nucleic Acids Research, 2010, 38, D620-D625.	14.5	218
6	Large-Scale Appearance of Ultraconserved Elements in Tetrapod Genomes and Slowdown of the Molecular Clock. Molecular Biology and Evolution, 2008, 25, 402-408.	8.9	103
7	Raising the estimate of functional human sequences: Figure 1.. Genome Research, 2007, 17, 1245-1253.	5.5	217
8	Orthologous MicroRNA Genes Are Located in Cancer-Associated Genomic Regions in Human and Mouse. PLoS ONE, 2007, 2, e1133.	2.5	34
9	The relationship between non-protein-coding DNA and eukaryotic complexity. BioEssays, 2007, 29, 288-299.	2.5	578
10	Maintenance of transposon-free regions throughout vertebrate evolution. BMC Genomics, 2007, 8, 470.	2.8	28
11	Evidence for Control of Splicing by Alternative RNA Secondary Structures in Dipteran Homothorax Pre-mRNA. RNA Biology, 2006, 3, 36-39.	3.1	13
12	Genomics: The amazing complexity of the human transcriptome. European Journal of Human Genetics, 2005, 13, 894-897.	2.8	171
13	Transposon-free regions in mammalian genomes. Genome Research, 2005, 16, 164-172.	5.5	102
14	Ultraconserved elements in insect genomes: A highly conserved intronic sequence implicated in the control of homothorax mRNA splicing. Genome Research, 2005, 15, 800-808.	5.5	112
15	Ultraconserved Elements in the Human Genome. Science, 2004, 304, 1321-1325.	12.6	1,496