

# Daryl J Thomas

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10477457/publications.pdf>

Version: 2024-02-01

12  
papers

12,141  
citations

759233

12  
h-index

1199594

12  
g-index

12  
all docs

12  
docs citations

12  
times ranked

20222  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification and analysis of functional elements in 1% of the human genome by the ENCODE pilot project. Nature, 2007, 447, 799-816.	27.8	4,709
2	A second generation human haplotype map of over 3.1 million SNPs. Nature, 2007, 449, 851-861.	27.8	4,137
3	Genome-wide detection and characterization of positive selection in human populations. Nature, 2007, 449, 913-918.	27.8	1,788
4	NF-AT components define a family of transcription factors targeted in T-cell activation. Nature, 1994, 369, 497-502.	27.8	572
5	Conserved noncoding sequences are selectively constrained and not mutation cold spots. Nature Genetics, 2006, 38, 223-227.	21.4	214
6	Analyses of deep mammalian sequence alignments and constraint predictions for 1% of the human genome. Genome Research, 2007, 17, 760-774.	5.5	184
7	Fast-evolving noncoding sequences in the human genome. Genome Biology, 2007, 8, R118.	9.6	163
8	NFATc3, a Lymphoid-specific NFATc Family Member That Is Calcium-regulated and Exhibits Distinct DNA Binding Specificity. Journal of Biological Chemistry, 1995, 270, 19898-19907.	3.4	159
9	The ENCODE Project at UC Santa Cruz. Nucleic Acids Research, 2007, 35, D663-D667.	14.5	92
10	Sequence features in regions of weak and strong linkage disequilibrium. Genome Research, 2005, 15, 1519-1534.	5.5	89
11	Variation resources at UC Santa Cruz. Nucleic Acids Research, 2007, 35, D716-D720.	14.5	19
12	Presymptomatic Risk Assessment for Chronic Non-Communicable Diseases. PLoS ONE, 2010, 5, e14338.	2.5	15