Daryl J Thomas

List of Publications by Year in descending order

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#	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.	13.7	4,709
2	A second generation human haplotype map of over 3.1 million SNPs. Nature, 2007, 449, 851-861.	13.7	4,137
3	Genome-wide detection and characterization of positive selection in human populations. Nature, 2007, 449, 913-918.	13.7	1,788
4	NF-AT components define a family of transcription factors targeted in T-cell activation. Nature, 1994, 369, 497-502.	13.7	572
5	Conserved noncoding sequences are selectively constrained and not mutation cold spots. Nature Genetics, 2006, 38, 223-227.	9.4	214
6	Analyses of deep mammalian sequence alignments and constraint predictions for 1% of the human genome. Genome Research, 2007, 17, 760-774.	2.4	184
7	Fast-evolving noncoding sequences in the human genome. Genome Biology, 2007, 8, R118.	13.9	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.	1.6	159
9	The ENCODE Project at UC Santa Cruz. Nucleic Acids Research, 2007, 35, D663-D667.	6.5	92
10	Sequence features in regions of weak and strong linkage disequilibrium. Genome Research, 2005, 15, 1519-1534.	2.4	89
11	Variation resources at UC Santa Cruz. Nucleic Acids Research, 2007, 35, D716-D720.	6.5	19
12	Presymptomatic Risk Assessment for Chronic Non-Communicable Diseases. PLoS ONE, 2010, 5, e14338.	1.1	15