Martijn Vermaat

List of Publications by Year in descending order

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Version: 2024-02-01

1040056 1474206 1,338 9 9 9 citations h-index g-index papers 9 9 9 5665 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Disease variants alter transcription factor levels and methylation of their binding sites. Nature Genetics, 2017, 49, 131-138.	21.4	390
2	Identification of context-dependent expression quantitative trait loci in whole blood. Nature Genetics, 2017, 49, 139-145.	21.4	363
3	The Genome of the Netherlands: design, and project goals. European Journal of Human Genetics, 2014, 22, 221-227.	2.8	246
4	Full-length mRNA sequencing uncovers a widespread coupling between transcription initiation and mRNA processing. Genome Biology, 2018, 19, 46.	8.8	106
5	Transmission of human mtDNA heteroplasmy in the Genome of the Netherlands families: support for a variable-size bottleneck. Genome Research, 2016, 26, 417-426.	5.5	84
6	Next-generation sequencing-based genome diagnostics across clinical genetics centers: implementation choices and their effects. European Journal of Human Genetics, 2015, 23, 1142-1150.	2.8	56
7	A SNP panel for identification of DNA and RNA specimens. BMC Genomics, 2018, 19, 90.	2.8	47
8	An efficient algorithm for the extraction of HGVS variant descriptions from sequences. Bioinformatics, 2015, 31, 3751-3757.	4.1	25
9	Integrated Whole Genome and Transcriptome Analysis Identified a Therapeutic Minor Histocompatibility Antigen in a Splice Variant of <i>ITGB2</i> . Clinical Cancer Research, 2016, 22, 4185-4196.	7.0	21