Kees Albers

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/9151609/publications.pdf

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13 papers	17,235 citations	840119 11 h-index	1125271 13 g-index
13 all docs	13 docs citations	13 times ranked	35943 citing authors

#	Article	IF	Citations
1	New insights into the genetic basis of TAR (thrombocytopenia-absent radii) syndrome. Current Opinion in Genetics and Development, 2013, 23, 316-323.	1.5	74
2	SMIM1 underlies the Vel blood group and influences red blood cell traits. Nature Genetics, 2013, 45, 542-545.	9.4	96
3	The origin, evolution, and functional impact of short insertionâ€“deletion variants identified in 179 human genomes. Genome Research, 2013, 23, 749-761.	2.4	206
4	Maps of open chromatin highlight cell typeâ€“restricted patterns of regulatory sequence variation at hematological trait loci. Genome Research, 2013, 23, 1130-1141.	2.4	34
5	Seventy-five genetic loci influencing the human red blood cell. Nature, 2012, 492, 369-375.	13.7	320
6	An integrated map of genetic variation from 1,092 human genomes. Nature, 2012, 491, 56-65.	13.7	7,199
7	Compound inheritance of a low-frequency regulatory SNP and a rare null mutation in exon-junction complex subunit RBM8A causes TAR syndrome. Nature Genetics, 2012, 44, 435-439.	9.4	355
8	A Systematic Survey of Loss-of-Function Variants in Human Protein-Coding Genes. Science, 2012, 335, 823-828.	6.0	1,095
9	Dindel: Accurate indel calls from short-read data. Genome Research, 2011, 21, 961-973.	2.4	383
10	Exome sequencing identifies NBEAL2 as the causative gene for gray platelet syndrome. Nature Genetics, 2011, 43, 735-737.	9.4	245
11	A map of human genome variation from population-scale sequencing. Nature, 2010, 467, 1061-1073.	13.7	7,209
12	Multipoint Approximations of Identity-by-Descent Probabilities for Accurate Linkage Analysis of Distantly Related Individuals. American Journal of Human Genetics, 2008, 82, 607-622.	2.6	9
13	Haplotype Inference in General Pedigrees Using the Cluster Variation Method. Genetics, 2007, 177, 1101-1116.	1.2	10