

# Michael W Smith

## List of Publications by Year in descending order

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33  
papers

6,837  
citations

201385

27  
h-index

395343

33  
g-index

33  
all docs

33  
docs citations

33  
times ranked

8516  
citing authors

#	ARTICLE	IF	CITATIONS
1	Selecting SNPs informative for African, American Indian and European Ancestry: application to the Family Investigation of Nephropathy and Diabetes (FIND). <i>BMC Genomics</i> , 2016, 17, 325.	1.2	1
2	Genome-Wide Association and Trans-ethnic Meta-Analysis for Advanced Diabetic Kidney Disease: Family Investigation of Nephropathy and Diabetes (FIND). <i>PLoS Genetics</i> , 2015, 11, e1005352.	1.5	118
3	Genomewide Linkage Scan for Diabetic Renal Failure and Albuminuria: The FIND Study. <i>American Journal of Nephrology</i> , 2011, 33, 381-389.	1.4	52
4	Genome-wide scans for footprints of natural selection. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2010, 365, 185-205.	1.8	343
5	Admixture Mapping Comes of Age. <i>Annual Review of Genomics and Human Genetics</i> , 2010, 11, 65-89.	2.5	192
6	Genome and gene alterations by insertions and deletions in the evolution of human and chimpanzee chromosome 22. <i>BMC Genomics</i> , 2009, 10, 51.	1.2	17
7	Association of Single-Nucleotide Polymorphisms in JAK3, STAT4, and STAT6 With New Cardiovascular Events in Incident Dialysis Patients. <i>American Journal of Kidney Diseases</i> , 2009, 53, 845-855.	2.1	19
8	The Genetic Structure and History of Africans and African Americans. <i>Science</i> , 2009, 324, 1035-1044.	6.0	1,267
9	MYH9 is a major-effect risk gene for focal segmental glomerulosclerosis. <i>Nature Genetics</i> , 2008, 40, 1175-1184.	9.4	636
10	MYH9 is associated with nondiabetic end-stage renal disease in African Americans. <i>Nature Genetics</i> , 2008, 40, 1185-1192.	9.4	587
11	Identifying Selected Regions from Heterozygosity and Divergence Using a Light-Coverage Genomic Dataset from Two Human Populations. <i>PLoS ONE</i> , 2008, 3, e1712.	1.1	50
12	Reliability of high-throughput genotyping of whole genome amplified DNA in SNP genotyping studies. <i>Electrophoresis</i> , 2007, 28, 2812-2817.	1.3	27
13	C-Reactive Protein Haplotype Predicts Serum C-Reactive Protein Levels But Not Cardiovascular Disease Risk in a Dialysis Cohort. <i>American Journal of Kidney Diseases</i> , 2007, 49, 118-126.	2.1	36
14	Elevated male European and female African contributions to the genomes of African American individuals. <i>Human Genetics</i> , 2006, 120, 713-722.	1.8	84
15	Behavioral Risk Exposure and Host Genetics of Susceptibility to HIV-1 Infection. <i>Journal of Infectious Diseases</i> , 2006, 193, 16-26.	1.9	49
16	Haplotype of Signal Transducer and Activator of Transcription 3 Gene Predicts Cardiovascular Disease in Dialysis Patients. <i>Journal of the American Society of Nephrology: JASN</i> , 2006, 17, 2285-2292.	3.0	10
17	Mapping by admixture linkage disequilibrium: advances, limitations and guidelines. <i>Nature Reviews Genetics</i> , 2005, 6, 623-632.	7.7	197
18	A High-Density Admixture Map for Disease Gene Discovery in African Americans. <i>American Journal of Human Genetics</i> , 2004, 74, 1001-1013.	2.6	416

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19	Methods for High-Density Admixture Mapping of Disease Genes. <i>American Journal of Human Genetics</i> , 2004, 74, 979-1000.	2.6	437
20	Dominant Effects of CCR2-CCR5 Haplotypes in HIV-1 Disease Progression. <i>Journal of Acquired Immune Deficiency Syndromes</i> (1999), 2004, 37, 1534-1538.	0.9	32
21	Markers for Mapping by Admixture Linkage Disequilibrium in African American and Hispanic Populations. <i>American Journal of Human Genetics</i> , 2001, 69, 1080-1094.	2.6	130
22	Influence of CCR5 promoter haplotypes on AIDS progression in African-Americans. <i>Aids</i> , 2000, 14, 2117-2122.	1.0	70
23	Polygenic and Multifactorial Disease Gene Association in Man: Lessons from AIDS. <i>Annual Review of Genetics</i> , 2000, 34, 563-591.	3.2	80
24	Significant Admixture Linkage Disequilibrium across 30 cM around the FY Locus in African Americans. <i>American Journal of Human Genetics</i> , 2000, 66, 969-978.	2.6	93
25	A Scan for Linkage Disequilibrium Across the Human Genome. <i>Genetics</i> , 1999, 152, 1711-1722.	1.2	158
26	Dating the Origin of the CCR5-Δ32 AIDS-Resistance Allele by the Coalescence of Haplotypes. <i>American Journal of Human Genetics</i> , 1998, 62, 1507-1515.	2.6	507
27	Genetic Restriction of AIDS Pathogenesis by an SDF-1 Chemokine Gene Variant. <i>Science</i> , 1998, 279, 389-393.	6.0	674
28	CCR2 chemokine receptor and AIDS progression. <i>Nature Medicine</i> , 1997, 3, 1052-1053.	15.2	96
29	Primary Structure and Phylogenetic Relationships of Glyceraldehyde-3-Phosphate Dehydrogenase Genes of Free-Living and Parasitic Diplomonad Flagellates. <i>Journal of Eukaryotic Microbiology</i> , 1996, 43, 330-340.	0.8	30
30	Evidence for the Heterolobosea from Phylogenetic Analysis of Genes Encoding Glyceraldehyde-3-Phosphate Dehydrogenase. <i>Journal of Eukaryotic Microbiology</i> , 1996, 43, 475-485.	0.8	53
31	Genomic sequence sampling: a strategy for high resolution sequence-based physical mapping of complex genomes. <i>Nature Genetics</i> , 1994, 7, 40-47.	9.4	63
32	Evolution by acquisition: the case for horizontal gene transfers. <i>Trends in Biochemical Sciences</i> , 1992, 17, 489-493.	3.7	292
33	Anomalous phylogeny involving the enzyme glucose-6-phosphate isomerase. <i>Journal of Molecular Evolution</i> , 1992, 34, 544-545.	0.8	21