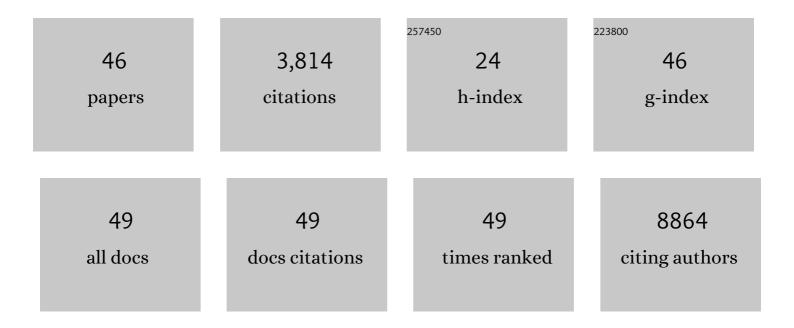
## Albert M Levin

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Neonatal gut microbiota associates with childhood multisensitized atopy and T cell differentiation. Nature Medicine, 2016, 22, 1187-1191.	30.7	844
2	Multi-ancestry genome-wide association study of 21,000 cases and 95,000 controls identifies new risk loci for atopic dermatitis. Nature Genetics, 2015, 47, 1449-1456.	21.4	529
3	Multiancestry association study identifies new asthma risk loci that colocalize with immune-cell enhancer marks. Nature Genetics, 2018, 50, 42-53.	21.4	426
4	Assembly of a pan-genome from deep sequencing of 910 humans of African descent. Nature Genetics, 2019, 51, 30-35.	21.4	276
5	Variation in the glucose transporter gene SLC2A2 is associated with glycemic response to metformin. Nature Genetics, 2016, 48, 1055-1059.	21.4	165
6	A continuum of admixture in the Western Hemisphere revealed by the African Diaspora genome. Nature Communications, 2016, 7, 12522.	12.8	136
7	Joint effects of pregnancy, sociocultural, and environmental factors on early life gut microbiome structure and diversity. Scientific Reports, 2016, 6, 31775.	3.3	122
8	Genome-wide association and HLA fine-mapping studies identify risk loci and genetic pathways underlying allergic rhinitis. Nature Genetics, 2018, 50, 1072-1080.	21.4	106
9	Genome-Wide Association Study of African and European Americans Implicates Multiple Shared and Ethnic Specific Loci in Sarcoidosis Susceptibility. PLoS ONE, 2012, 7, e43907.	2.5	105
10	Discovery and fine-mapping of adiposity loci using high density imputation of genome-wide association studies in individuals of African ancestry: African Ancestry Anthropometry Genetics Consortium. PLoS Genetics, 2017, 13, e1006719.	3.5	98
11	Multiethnic meta-analysis identifies ancestry-specific and cross-ancestry loci for pulmonary function. Nature Communications, 2018, 9, 2976.	12.8	85
12	Association study in African-admixed populations across the Americas recapitulates asthma risk loci in non-African populations. Nature Communications, 2019, 10, 880.	12.8	71
13	Challenges and disparities in the application of personalized genomic medicine to populations with African ancestry. Nature Communications, 2016, 7, 12521.	12.8	68
14	Ethnic-specific associations of rare and low-frequency DNA sequence variants with asthma. Nature Communications, 2015, 6, 5965.	12.8	66
15	Mutational Landscape of Aggressive Prostate Tumors in African American Men. Cancer Research, 2016, 76, 1860-1868.	0.9	61
16	A meta-analysis of genome-wide association studies for serum total IgE in diverse study populations. Journal of Allergy and Clinical Immunology, 2013, 131, 1176-1184.	2.9	58
17	A genome-wide association and admixture mapping study of bronchodilator drug response in African Americans with asthma. Pharmacogenomics Journal, 2019, 19, 249-259.	2.0	54
18	Genome-wide association study and admixture mapping reveal new loci associated with total IgE levels in Latinos. Journal of Allergy and Clinical Immunology, 2015, 135, 1502-1510.	2.9	52

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19	Expression quantitative trait locus fine mapping of the 17q12–21 asthma locus in African American children: a genetic association and gene expression study. Lancet Respiratory Medicine,the, 2020, 8, 482-492.	10.7	47
20	Association of <i>HLA</i> - <i>DRB1</i> with Sarcoidosis Susceptibility and Progression in African Americans. American Journal of Respiratory Cell and Molecular Biology, 2015, 53, 206-216.	2.9	42
21	Integrative approach identifies corticosteroid response variant in diverse populations with asthma. Journal of Allergy and Clinical Immunology, 2019, 143, 1791-1802.	2.9	33
22	Admixture Fine-Mapping in African Americans Implicates XAF1 as a Possible Sarcoidosis Risk Gene. PLoS ONE, 2014, 9, e92646.	2.5	31
23	Identification of miRNomes associated with adult neurogenesis after stroke using Argonaute 2-based RNA sequencing. RNA Biology, 2017, 14, 488-499.	3.1	30
24	Genetic determinants of telomere length from 109,122 ancestrally diverse whole-genome sequences in TOPMed. Cell Genomics, 2022, 2, 100084.	6.5	29
25	Real-Time Feedback of Histotripsy Thrombolysis Using Bubble-Induced Color Doppler. Ultrasound in Medicine and Biology, 2015, 41, 1386-1401.	1.5	26
26	Performance of HLA allele prediction methods in African Americans for class II genes HLA-DRB1, â^'DQB1, and –DPB1. BMC Genetics, 2014, 15, 72.	2.7	24
27	Single Cell Transcriptomics Implicate Novel Monocyte and T Cell Immune Dysregulation in Sarcoidosis. Frontiers in Immunology, 2020, 11, 567342.	4.8	21
28	Genome-Wide Association Study of Ocular Sarcoidosis Confirms HLA Associations and Implicates Barrier Function and Autoimmunity in African Americans. Ocular Immunology and Inflammation, 2021, 29, 244-249.	1.8	21
29	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. American Journal of Human Genetics, 2021, 108, 564-582.	6.2	18
30	Fine mapping of chromosome 15q25 implicates <scp>ZNF</scp> 592 in neurosarcoidosis patients. Annals of Clinical and Translational Neurology, 2015, 2, 972-977.	3.7	17
31	Association between cesarean delivery types and obesity in preadolescence. International Journal of Obesity, 2020, 44, 2023-2034.	3.4	17
32	Breast and prostate cancers harbor common somatic copy number alterations that consistently differ by race and are associated with survival. BMC Medical Genomics, 2020, 13, 116.	1.5	17
33	Mapping the 17q12–21.1 Locus for Variants Associated with Early-Onset Asthma in African Americans. American Journal of Respiratory and Critical Care Medicine, 2021, 203, 424-436.	5.6	16
34	Asthma and its relationship to mitochondrial copy number: Results from the Asthma Translational Genomics Collaborative (ATGC) of the Trans-Omics for Precision Medicine (TOPMed) program. PLoS ONE, 2020, 15, e0242364.	2.5	16
35	Haplotype and diplotype analyses of variation in <i>ERCC5</i> transcription <i>cis</i> -regulation in normal bronchial epithelial cells. Physiological Genomics, 2016, 48, 537-543.	2.3	13
36	Association between cadmium and androgen receptor protein expression differs in prostate tumors of African American and European American men. Journal of Trace Elements in Medicine and Biology, 2018, 48, 233-238.	3.0	13

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37	Infant gut bacterial community composition and foodâ€related manifestation of atopy in early childhood. Pediatric Allergy and Immunology, 2022, 33, .	2.6	13
38	Extended methods for gene–environmentâ€wide interaction scans in studies of admixed individuals with varying degrees of relationships. Genetic Epidemiology, 2019, 43, 414-426.	1.3	10
39	How does race and ethnicity effect the precision treatment of asthma?. Expert Review of Precision Medicine and Drug Development, 2019, 4, 337-356.	0.7	7
40	Performance of the Genomic Evaluators of Metastatic Prostate Cancer (GEMCaP) Tumor Biomarker for Identifying Recurrent Disease in African American Patients. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 1677-1682.	2.5	6
41	Whole Genome Sequencing Identifies CRISPLD2 as a Lung Function Gene in Children With Asthma. Chest, 2019, 156, 1068-1079.	0.8	5
42	Novel HLA associations with outcomes of <i>Mycobacterium tuberculosis</i> exposure and sarcoidosis in individuals of African ancestry using nearestâ€neighbor feature selection. Genetic Epidemiology, 2022, 46, 463-474.	1.3	5
43	MicroRNA Profile Differentiates Head and Neck Keloid and Adjacent Normal Skin Tissue. Facial Plastic Surgery and Aesthetic Medicine, 2021, , .	0.9	2
44	Exploring latent classes to identify prenatal and early-life sources of racial disparities in allergic disease. Annals of Allergy, Asthma and Immunology, 2019, 122, 650-652.e1.	1.0	0
45	Abstract P751: Global Analyses of Protein and Microrna Cargo in Neural Stem Cell Derived-Exosomes After Brain Ischemia. Stroke, 2021, 52, .	2.0	0
46	Powerful Tests for Multi-Marker Association Analysis Using Ensemble Learning. PLoS ONE, 2015, 10, e0143489.	2.5	0