Digna R Velez Edwards

List of Publications by Year in descending order

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107 papers

6,599 citations

36 h-index 79698 73 g-index

118 all docs

118 docs citations

118 times ranked

13795 citing authors

#	Article	IF	CITATIONS
1	Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits. Nature Genetics, 2018 , 50 , 1412 - 1425 .	21.4	924
2	Rare and low-frequency coding variants alter human adult height. Nature, 2017, 542, 186-190.	27.8	544
3	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. Nature Genetics, 2017, 49, 834-841.	21.4	426
4	Trans-ethnic association study of blood pressure determinants in over 750,000 individuals. Nature Genetics, 2019, 51, 51-62.	21.4	328
5	Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. Nature Genetics, 2018, 50, 26-41.	21.4	286
6	A meta-analysis identifies new loci associated with body mass index in individuals of African ancestry. Nature Genetics, 2013, 45, 690-696.	21.4	232
7	Meta-analysis identifies five novel loci associated with endometriosis highlighting key genes involved in hormone metabolism. Nature Communications, 2017, 8, 15539.	12.8	230
8	Meta-analysis identifies common and rare variants influencing blood pressure and overlapping with metabolic trait loci. Nature Genetics, 2016, 48, 1162-1170.	21.4	223
9	Genome-wide Association Analysis of Blood-Pressure Traits in African-Ancestry Individuals Reveals Common Associated Genes in African and Non-African Populations. American Journal of Human Genetics, 2013, 93, 545-554.	6.2	189
10	Phenotype risk scores identify patients with unrecognized Mendelian disease patterns. Science, 2018, 359, 1233-1239.	12.6	164
11	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. Cancer Discovery, 2016, 6, 1052-1067.	9.4	157
12	Toll-like Receptor 1 Polymorphisms Increase Susceptibility to Candidemia. Journal of Infectious Diseases, 2012, 205, 934-943.	4.0	116
13	Population Stratification in Genetic Association Studies. Current Protocols in Human Genetics, 2017, 95, 1.22.1-1.22.23.	3.5	108
14	Risk of Miscarriage Among Black Women and White Women in a US Prospective Cohort Study. American Journal of Epidemiology, 2013, 177, 1271-1278.	3.4	102
15	Discovery of rare variants associated with blood pressure regulation through meta-analysis of 1.3 million individuals. Nature Genetics, 2020, 52, $1314-1332$.	21.4	91
16	Shared heritability and functional enrichment across six solid cancers. Nature Communications, 2019, 10, 431.	12.8	88
17	Single-trait and multi-trait genome-wide association analyses identify novel loci for blood pressure in African-ancestry populations. PLoS Genetics, 2017, 13, e1006728.	3.5	88
18	Platelet-Related Variants Identified by Exomechip Meta-analysis in 157,293 Individuals. American Journal of Human Genetics, 2016, 99, 40-55.	6.2	82

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19	Obesity and pelvic organ prolapse: a systematic review and meta-analysis of observational studies. American Journal of Obstetrics and Gynecology, 2017, 217, 11-26.e3.	1.3	81
20	Genetic Variation in the Dectin-1/CARD9 Recognition Pathway and Susceptibility to Candidemia. Journal of Infectious Diseases, 2011, 204, 1138-1145.	4.0	80
21	A genetic association study of maternal and fetal candidate genes that predispose to preterm prelabor rupture of membranes (PROM). American Journal of Obstetrics and Gynecology, 2010, 203, 361.e1-361.e30.	1.3	78
22	Identification of fetal and maternal single nucleotide polymorphisms in candidate genes that predispose to spontaneous preterm labor with intact membranes. American Journal of Obstetrics and Gynecology, 2010, 202, 431.e1-431.e34.	1.3	77
23	Cytokine Gene Polymorphisms and the Outcome of Invasive Candidiasis: A Prospective Cohort Study. Clinical Infectious Diseases, 2012, 54, 502-510.	5.8	68
24	Genetic epidemiology of pelvic organ prolapse: a systematic review. American Journal of Obstetrics and Gynecology, 2014, 211, 326-335.	1.3	62
25	Exome Genotyping Identifies Pleiotropic Variants Associated with Red Blood Cell Traits. American Journal of Human Genetics, 2016, 99, 8-21.	6.2	60
26	Alcohol Use in Pregnancy and Miscarriage: A Systematic Review and Metaâ€Analysis. Alcoholism: Clinical and Experimental Research, 2019, 43, 1606-1616.	2.4	55
27	A Transcriptome-Wide Association Study Among 97,898 Women to Identify Candidate Susceptibility Genes for Epithelial Ovarian Cancer Risk. Cancer Research, 2018, 78, 5419-5430.	0.9	54
28	Large-Scale Exome-wide Association Analysis Identifies Loci for White Blood Cell Traits and Pleiotropy with Immune-Mediated Diseases. American Journal of Human Genetics, 2016, 99, 22-39.	6.2	50
29	Genetic Data from Nearly 63,000 Women of European Descent Predicts DNA Methylation Biomarkers and Epithelial Ovarian Cancer Risk. Cancer Research, 2019, 79, 505-517.	0.9	49
30	Deep learning predicts extreme preterm birth from electronic health records. Journal of Biomedical Informatics, 2019, 100, 103334.	4.3	49
31	A genome-wide association study of polycystic ovary syndrome identified from electronic health records. American Journal of Obstetrics and Gynecology, 2020, 223, 559.e1-559.e21.	1.3	49
32	Admixture mapping identifies a locus at 15q21.2–22.3 associated with keloid formation in African Americans. Human Genetics, 2014, 133, 1513-1523.	3.8	47
33	Periconceptional Over-the-Counter Nonsteroidal Anti-inflammatory Drug Exposure and Risk for Spontaneous Abortion. Obstetrics and Gynecology, 2012, 120, 113-122.	2.4	46
34	Association of Age at Menarche With Increasing Number of Fibroids in a Cohort of Women Who Underwent Standardized Ultrasound Assessment. American Journal of Epidemiology, 2013, 178, 426-433.	3.4	43
35	HTR1B, ADIPOR1, PPARGC1A, and CYP19A1 and Obesity in a Cohort of Caucasians and African Americans: An Evaluation of Gene-Environment Interactions and Candidate Genes. American Journal of Epidemiology, 2012, 175, 11-21.	3.4	42
36	Gene-environment interactions and obesity traits among postmenopausal African-American and Hispanic women in the Women's Health Initiative SHARe Study. Human Genetics, 2013, 132, 323-336.	3.8	41

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37	Association of Genetic Variants, Ethnicity and Preterm Birth with Amniotic Fluid Cytokine Concentrations. Annals of Human Genetics, 2010, 74, 165-183.	0.8	40
38	Genomeâ€Wide Association and Linkage Study in the Amish Detects a Novel Candidate Lateâ€Onset Alzheimer Disease Gene. Annals of Human Genetics, 2012, 76, 342-351.	0.8	40
39	A Polygenic and Phenotypic Risk Prediction for Polycystic Ovary Syndrome Evaluated by Phenome-Wide Association Studies. Journal of Clinical Endocrinology and Metabolism, 2020, 105, 1918-1936.	3.6	40
40	A multi-stage genome-wide association study of uterine fibroids in African Americans. Human Genetics, 2017, 136, 1363-1373.	3.8	39
41	Leiomyomas in Pregnancy and Spontaneous Abortion. Obstetrics and Gynecology, 2017, 130, 1065-1072.	2.4	37
42	Expression Quantitative Trait Locus Mapping Studies in Mid-secretory Phase Endometrial Cells Identifies HLA-F and TAP2 as Fecundability-Associated Genes. PLoS Genetics, 2016, 12, e1005858.	3.5	36
43	Inverse Association of Female Hormone Replacement Therapy with Age-Related Macular Degeneration and Interactions with <i>ARMS2 </i> Polymorphisms., 2010, 51, 1873.		33
44	Interleukin 12B (IL12B) Genetic Variation and Pulmonary Tuberculosis: A Study of Cohorts from The Gambia, Guinea-Bissau, United States and Argentina. PLoS ONE, 2011, 6, e16656.	2.5	33
45	BET1L and TNRC6B associate with uterine fibroid risk among European Americans. Human Genetics, 2013, 132, 943-953.	3.8	33
46	Uterine leiomyomata and cesarean birth risk: a prospective cohort with standardized imaging. Annals of Epidemiology, 2014, 24, 122-126.	1.9	32
47	A Trans-Ethnic Genome-Wide Association Study of Uterine Fibroids. Frontiers in Genetics, 2019, 10, 511.	2.3	32
48	Progestogens for Preterm Birth Prevention. Obstetrics and Gynecology, 2012, 120, 897-907.	2.4	31
49	Variants in BET1L and TNRC6B associate with increasing fibroid volume and fibroid type among European Americans. Human Genetics, 2013, 132, 1361-1369.	3.8	28
50	Successful Aging Shows Linkage to Chromosomes 6, 7, and 14 in the Amish. Annals of Human Genetics, 2011, 75, 516-528.	0.8	27
51	Interpregnancy Interval After Pregnancy Loss and Risk of Repeat Miscarriage. Obstetrics and Gynecology, 2017, 130, 1312-1318.	2.4	26
52	Linkage and association of successful aging to the 6q25 region in large Amish kindreds. Age, 2013, 35, 1467-1477.	3.0	25
53	African genetic ancestry interacts with body mass index to modify risk for uterine fibroids. PLoS Genetics, 2017, 13, e1006871.	3.5	25
54	Week-by-week alcohol consumption in early pregnancy and spontaneous abortion risk: a prospective cohort study. American Journal of Obstetrics and Gynecology, 2021, 224, 97.e1-97.e16.	1.3	24

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55	Race/Ethnicity and the Pharmacogenetics of Reported Suicidality With Efavirenz Among Clinical Trials Participants. Journal of Infectious Diseases, 2017, 216, 554-564.	4.0	23
56	Association of Thyroid Function Genetic Predictors With Atrial Fibrillation. JAMA Cardiology, 2019, 4, 136.	6.1	23
57	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. European Journal of Human Genetics, 2022, 30, 349-362.	2.8	23
58	Relationship between very low low-density lipoprotein cholesterol concentrations not due to statin therapy and risk of type 2 diabetes: A US-based cross-sectional observational study using electronic health records. PLoS Medicine, 2018, 15, e1002642.	8.4	22
59	Heritability and genome-wide association study of benign prostatic hyperplasia (BPH) in the eMERGE network. Scientific Reports, 2019, 9, 6077.	3.3	21
60	Enhancing Uterine Fibroid Research Through Utilization of Biorepositories Linked to Electronic Medical Record Data. Journal of Women's Health, 2014, 23, 1027-1032.	3.3	20
61	A genome-wide association study meta-analysis of clinical fracture in 10,012 African American women. Bone Reports, 2016, 5, 233-242.	0.4	20
62	Validation of maternal recall of early pregnancy medication exposure using prospective diary data. Annals of Epidemiology, 2017, 27, 135-139.e2.	1.9	19
63	Maternal Western-style diet affects offspring islet composition and function in a non-human primate model of maternal over-nutrition. Molecular Metabolism, 2019, 25, 73-82.	6.5	19
64	Genome-Wide Association Study of Serum Creatinine Levels during Vancomycin Therapy. PLoS ONE, 2015, 10, e0127791.	2.5	19
65	Polymorphisms in maternal and fetal genes encoding for proteins involved in extracellular matrix metabolism alter the risk for small-for-gestational-age. Journal of Maternal-Fetal and Neonatal Medicine, 2011, 24, 362-380.	1.5	17
66	Genetic Determinants of Pelvic Organ Prolapse among African American and Hispanic Women in the Women's Health Initiative. PLoS ONE, 2015, 10, e0141647.	2.5	17
67	Evidence of selection as a cause for racial disparities in fibroproliferative disease. PLoS ONE, 2017, 12, e0182791.	2.5	17
68	MCP1 SNPs and Pulmonary Tuberculosis in Cohorts from West Africa, the USA and Argentina: Lack of Association or Epistasis with IL12B Polymorphisms. PLoS ONE, 2012, 7, e32275.	2.5	16
69	Analysis of potential protein-modifying variants in 9000 endometriosis patients and 150000 controls of European ancestry. Scientific Reports, 2017, 7, 11380.	3.3	16
70	Characterizing the Clinical and Genetic Spectrum of Polycystic Ovary Syndrome in Electronic Health Records. Journal of Clinical Endocrinology and Metabolism, 2021, 106, 153-167.	3.6	16
71	Gene-based evaluation of low-frequency variation and genetically-predicted gene expression impacting risk of keloid formation. Annals of Human Genetics, 2018, 82, 206-215.	0.8	15
72	Equity in Health: Consideration of Race and Ethnicity in Precision Medicine. Trends in Genetics, 2020, 36, 807-809.	6.7	14

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73	Transethnic and race-stratified genome-wide association study of fibroid characteristics in African American and European American women. Fertility and Sterility, 2018, 110, 737-745.e34.	1.0	12
74	Cross-Cancer Genome-Wide Association Study of Endometrial Cancer and Epithelial Ovarian Cancer Identifies Genetic Risk Regions Associated with Risk of Both Cancers. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 217-228.	2.5	12
75	Progestogens for preterm birth prevention: a systematic review and meta-analysis by drug route. Archives of Gynecology and Obstetrics, 2013, 287, 1059-1066.	1.7	11
76	Admixture mapping of uterine fibroid size and number in African American women. Fertility and Sterility, 2017, 108, 1034-1042.e26.	1.0	11
77	Western-style diet consumption impairs maternal insulin sensitivity and glucose metabolism during pregnancy in a Japanese macaque model. Scientific Reports, 2021, 11, 12977.	3.3	11
78	Phenotyping clinical disorders: lessons learned from pelvic organ prolapse. American Journal of Obstetrics and Gynecology, 2013, 208, 360-365.	1.3	10
79	Racial differences in risk of spontaneous abortions associated with periconceptional over-the-counter nonsteroidal anti-inflammatory drug exposure. Annals of Epidemiology, 2014, 24, 111-115.e1.	1.9	10
80	What Results Should Be Returned from Opportunistic Screening in Translational Research?. Journal of Personalized Medicine, 2020, 10, 13.	2.5	10
81	<i>PTX3</i> Genetic Variation and Dizygotic Twinning in The Gambia: Could Pleiotropy with Innate Immunity Explain Common Dizygotic Twinning in Africa?. Annals of Human Genetics, 2012, 76, 454-463.	0.8	9
82	First-trimester antihistamine exposure and risk of spontaneous abortion or preterm birth. Pharmacoepidemiology and Drug Safety, 2014, 23, 1043-1050.	1.9	9
83	Gestational Age at Arrest of Development: An Alternative Approach for Assigning Time at Risk in Studies of Time-Varying Exposures and Miscarriage. American Journal of Epidemiology, 2019, 188, 570-578.	3.4	9
84	Evidence that geographic variation in genetic ancestry associates with uterine fibroids. Human Genetics, 2021, 140, 1433-1440.	3.8	9
85	Evaluating risk factors for differences in fibroid size and number using a large electronic health record population. Maturitas, 2018, 114, 9-13.	2.4	8
86	Evaluating the role of race and medication in protection of uterine fibroids by type 2 diabetes exposure. BMC Women's Health, 2017, 17, 28.	2.0	7
87	Uterine fibroids and risk of preterm birth by clinical subtypes: a prospective cohort study. BMC Pregnancy and Childbirth, 2021, 21, 560.	2.4	7
88	Analyses of germline variants associated with ovarian cancer survival identify functional candidates at the 1q22 and 19p12 outcome loci. Oncotarget, 2017, 8, 64670-64684.	1.8	7
89	Association of gene coding variation and resting metabolic rate in a multi-ethnic sample of children and adults. BMC Obesity, 2017, 4, 12.	3.1	6
90	Association between First Trimester Antidepressant Use and Risk of Spontaneous Abortion. Pharmacotherapy, 2019, 39, 889-898.	2.6	6

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91	Evaluation of vitamin D biosynthesis and pathway target genes reveals UGT2A1/2 and EGFR polymorphisms associated with epithelial ovarian cancer in African American Women. Cancer Medicine, 2019, 8, 2503-2513.	2.8	6
92	Estimating Uterine Fibroid SNP-Based Heritability in European American Women with Imaging-Confirmed Fibroids. Human Heredity, 2019, 84, 73-81.	0.8	5
93	Determinants of stage at diagnosis of HPV-related cancer including area deprivation and clinical factors. Journal of Public Health, 2022, 44, 18-27.	1.8	5
94	Admixture mapping of pelvic organ prolapse in African Americans from the Women's Health Initiative Hormone Therapy trial. PLoS ONE, 2017, 12, e0178839.	2.5	4
95	Association of uterine fibroids with birthweight and gestational age. Annals of Epidemiology, 2020, 50, 35-40.e2.	1.9	4
96	Learning to Identify Severe Maternal Morbidity from Electronic Health Records. Studies in Health Technology and Informatics, 2019, 264, 143-147.	0.3	4
97	Uterine fibroid polygenic risk score (PRS) associates and predicts risk for uterine fibroid. Human Genetics, 2022, 141, 1739-1748.	3.8	4
98	The Relationship between Total Fibroid Burden and First Trimester Bleeding and Pain. Paediatric and Perinatal Epidemiology, 2016, 30, 115-123.	1.7	3
99	Racial and Ethnic Variation in Genetic Susceptibility: Are Disparities in Infertility Prevalence and Outcomes more than Black and White?. Reproductive Sciences, 2022, 29, 2081-2083.	2.5	3
100	Association of Apparent Treatment-Resistant Hypertension With Differential Risk of End-Stage Kidney Disease Across Racial Groups in the Million Veteran Program. Hypertension, 2021, 78, 376-386.	2.7	2
101	Nonsteroidal Anti-inflammatory Drug Interaction with Prostacyclin Synthase Protects from Miscarriage. Scientific Reports, 2017, 7, 9874.	3.3	1
102	THE AUTHORS REPLY. American Journal of Epidemiology, 2018, 187, 1133-1134.	3.4	1
103	Learning the impact of acute and chronic diseases on forecasting neonatal encephalopathy. Computer Methods and Programs in Biomedicine, 2021, 211, 106397.	4.7	1
104	Associations of biogeographic ancestry with hypertension traits. Journal of Hypertension, 2021, 39, 633-642.	0.5	1
105	Leveraging Electronic Health Records to Learn Progression Path for Severe Maternal Morbidity. Studies in Health Technology and Informatics, 2019, 264, 148-152.	0.3	1
106	Genetic Sex Effects of Polycystic Ovary Syndrome Reveal Distinct Metabolic Etiology. Journal of the Endocrine Society, 2021, 5, A766-A766.	0.2	0
107	SAT-024 Investigating Racial and Ethnic Comorbidity Patterns of Polycystic Ovary Syndrome. Journal of the Endocrine Society, 2020, 4, .	0.2	0