

# Amanda Jane Phipps-Green

## List of Publications by Year in descending order

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Version: 2024-02-01

51  
papers

1,583  
citations

279798

23  
h-index

315739

38  
g-index

54  
all docs

54  
docs citations

54  
times ranked

2159  
citing authors

#	ARTICLE	IF	CITATIONS
1	Relationship between serum urate concentration and clinically evident incident gout: an individual participant data analysis. <i>Annals of the Rheumatic Diseases</i> , 2018, 77, 1048-1052.	0.9	131
2	Twenty-eight loci that influence serum urate levels: analysis of association with gout. <i>Annals of the Rheumatic Diseases</i> , 2016, 75, 124-130.	0.9	116
3	A strong role for the ABCG2 gene in susceptibility to gout in New Zealand Pacific Island and Caucasian, but not Māori, case and control sample sets. <i>Human Molecular Genetics</i> , 2010, 19, 4813-4819.	2.9	100
4	Evidence of interaction of CARD8 rs2043211 with NALP3 rs35829419 in Crohn's disease. <i>Genes and Immunity</i> , 2010, 11, 351-356.	4.1	92
5	ABCG2 loss-of-function polymorphism predicts poor response to allopurinol in patients with gout. <i>Pharmacogenomics Journal</i> , 2017, 17, 201-203.	2.0	82
6	Sugar-sweetened beverage consumption: a risk factor for prevalent gout with <i>SLC2A9</i> genotype-specific effects on serum urate and risk of gout. <i>Annals of the Rheumatic Diseases</i> , 2014, 73, 2101-2106.	0.9	77
7	The ABCG2 Q141K hyperuricemia and gout associated variant illuminates the physiology of human urate excretion. <i>Nature Communications</i> , 2020, 11, 2767.	12.8	71
8	Modulation of Genetic Associations with Serum Urate Levels by Body-Mass-Index in Humans. <i>PLoS ONE</i> , 2015, 10, e0119752.	2.5	64
9	Association of variation in Fcγ receptor 3B gene copy number with rheumatoid arthritis in Caucasian samples. <i>Annals of the Rheumatic Diseases</i> , 2010, 69, 1711-1716.	0.9	63
10	Population-specific influence of <i>SLC2A9</i> genotype on the acute hyperuricaemic response to a fructose load. <i>Annals of the Rheumatic Diseases</i> , 2013, 72, 1868-1873.	0.9	61
11	The renal urate transporter SLC17A1 locus: confirmation of association with gout. <i>Arthritis Research and Therapy</i> , 2012, 14, R92.	3.5	53
12	Gout Is a Chronic Inflammatory Disease in Which High Levels of Interleukin-8 (CXCL8), Myeloid-Related Protein 8/Myeloid-Related Protein 14 Complex, and an Altered Proteome Are Associated With Diabetes Mellitus and Cardiovascular Disease. <i>Arthritis and Rheumatology</i> , 2015, 67, 3303-3313.	5.6	51
13	Population-specific Resequencing Associates the ATP-binding Cassette Subfamily C Member 4 Gene With Gout in New Zealand Māori and Pacific Men. <i>Arthritis and Rheumatology</i> , 2017, 69, 1461-1469.	5.6	46
14	mTOR inhibition by metformin impacts monosodium urate crystal-induced inflammation and cell death in gout: a prelude to a new add-on therapy?. <i>Annals of the Rheumatic Diseases</i> , 2019, 78, 663-671.	0.9	45
15	Gout, Rheumatoid Arthritis, and the Risk of Death Related to Coronavirus Disease 2019: An Analysis of the UK Biobank. <i>ACR Open Rheumatology</i> , 2021, 3, 333-340.	2.1	37
16	Association analysis of the SLC22A11 (organic anion transporter 4) and SLC22A12 (urate transporter 1) urate transporter locus with gout in New Zealand case-control sample sets reveals multiple ancestral-specific effects. <i>Arthritis Research and Therapy</i> , 2013, 15, R220.	3.5	35
17	Association of the lipoprotein receptor-related protein 2 gene with gout and non-additive interaction with alcohol consumption. <i>Arthritis Research and Therapy</i> , 2013, 15, R177.	3.5	34
18	Hyperuricaemia: contributions of urate transporter ABCG2 and the fractional renal clearance of urate. <i>Annals of the Rheumatic Diseases</i> , 2016, 75, 1363-1366.	0.9	30

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19	Differential DNA Methylation of Networked Signaling, Transcriptional, Innate and Adaptive Immunity, and Osteoclastogenesis Genes and Pathways in Gout. <i>Arthritis and Rheumatology</i> , 2020, 72, 802-814.	5.6	30
20	Systematic genetic analysis of early-onset gout: ABCG2 is the only associated locus. <i>Rheumatology</i> , 2020, 59, 2544-2549.	1.9	30
21	Pleiotropic effect of the ABCG2 gene in gout: involvement in serum urate levels and progression from hyperuricemia to gout. <i>Arthritis Research and Therapy</i> , 2020, 22, 45.	3.5	28
22	Immunostimulatory biodegradable implants containing the adjuvant Quil-A™ Part II: In vivo evaluation. <i>Journal of Drug Targeting</i> , 2008, 16, 224-232.	4.4	27
23	Association of Autoimmune Addison's Disease with Alleles of STAT4 and GATA3 in European Cohorts. <i>PLoS ONE</i> , 2014, 9, e88991.	2.5	27
24	Influence of the ABCG2 gout risk 141K allele on urate metabolism during a fructose challenge. <i>Arthritis Research and Therapy</i> , 2014, 16, R34.	3.5	27
25	A non-coding genetic variant maximally associated with serum urate levels is functionally linked to HNF4A-dependent PDZK1 expression. <i>Human Molecular Genetics</i> , 2018, 27, 3964-3973.	2.9	26
26	Population-specific association between ABCG2 variants and tophaceous disease in people with gout. <i>Arthritis Research and Therapy</i> , 2017, 19, 43.	3.5	25
27	Body mass index modulates the relationship of sugar-sweetened beverage intake with serum urate concentrations and gout. <i>Arthritis Research and Therapy</i> , 2015, 17, 263.	3.5	24
28	Replication of association of the apolipoprotein A1-C3-A4 gene cluster with the risk of gout. <i>Rheumatology</i> , 2016, 55, 1421-1430.	1.9	16
29	The ITGAV rs3738919 variant and susceptibility to rheumatoid arthritis in four Caucasian sample sets. <i>Arthritis Research and Therapy</i> , 2009, 11, R152.	3.5	14
30	Replication of association of the interleukin 23 receptor rs1343151 variant with rheumatoid arthritis in Caucasian sample sets. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 155-157.	0.9	13
31	The impact of diuretic use and ABCG2 genotype on the predictive performance of a published allopurinol dosing tool. <i>British Journal of Clinical Pharmacology</i> , 2018, 84, 937-943.	2.4	11
32	Population-specific factors associated with fractional excretion of uric acid. <i>Arthritis Research and Therapy</i> , 2019, 21, 234.	3.5	11
33	Association analysis of the beta-3 adrenergic receptor Trp64Arg (rs4994) polymorphism with urate and gout. <i>Rheumatology International</i> , 2016, 36, 255-261.	3.0	10
34	The Oxytocin Receptor Gene (OXTR) Variant rs53576 Is Not Related to Emotional Traits or States in Young Adults. <i>Frontiers in Psychology</i> , 2018, 9, 2548.	2.1	9
35	Aotearoa New Zealand Māori and Pacific Population-amplified Gout Risk Variants: CLNK Is a Separate Risk Gene at the SLC2A9 Locus. <i>Journal of Rheumatology</i> , 2021, 48, 1736-1744.	2.0	8
36	No evidence for association of the systemic lupus erythematosus-associated ITGAM variant, R77H, with rheumatoid arthritis in the Caucasian population. <i>Rheumatology</i> , 2009, 48, 1614-1615.	1.9	7

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37	Population-specific effects of <i>SLC17A1</i> genotype on serum urate concentrations and renal excretion of uric acid during a fructose load. <i>Annals of the Rheumatic Diseases</i> , 2014, 73, 313-314.	0.9	7
38	Potential <i>PINK1</i> Founder Effect in Polynesia Causing Early-Onset Parkinson's Disease. <i>Movement Disorders</i> , 2021, 36, 2199-2200.	3.9	7
39	Trans-ancestral dissection of urate- and gout-associated major loci <i>SLC2A9</i> and <i>ABCG2</i> reveals primate-specific regulatory effects. <i>Journal of Human Genetics</i> , 2021, 66, 161-169.	2.3	6
40	Clinical and genetic features of diuretic-associated gout: a case-control study. <i>Rheumatology</i> , 2016, 55, 1172-1176.	1.9	5
41	Mid-pass whole genome sequencing enables biomedical genetic studies of diverse populations. <i>BMC Genomics</i> , 2021, 22, 666.	2.8	5
42	<i>Smad2</i> : A Candidate Gene for the Murine Autoimmune Diabetes Locus <i>Idd21.1</i> . <i>Journal of Clinical Endocrinology and Metabolism</i> , 2011, 96, E2072-E2077.	3.6	4
43	Genetic Polymorphisms on <i>OPRM1</i> , <i>DRD2</i> , <i>DRD4</i> , and <i>COMT</i> in Young Adults: Lack of Association With Alcohol Consumption. <i>Frontiers in Psychiatry</i> , 2020, 11, 549429.	2.6	4
44	Influence of genetic variants on renal uric acid handling in response to frusemide: an acute intervention study. <i>RMD Open</i> , 2017, 3, e000424.	3.8	3
45	Correspondence on Variants in urate transporters, <i>ADH1B</i> , <i>GCKR</i> and <i>MEPE</i> genes associated with transition from asymptomatic hyperuricaemia to gout: results of the first gout versus asymptomatic hyperuricaemia GWAS in Caucasians using data from the UK Biobank™. <i>Annals of the Rheumatic Diseases</i> , 2023, 82, e174-e174.	0.9	3
46	A Polynesian-specific copy number variant encompassing the <i>MICA</i> gene associates with gout. <i>Human Molecular Genetics</i> , 2022, 31, 3757-3768.	2.9	3
47	Analysis of association of <i>DNASE2</i> promoter variation with rheumatoid arthritis in European Caucasians. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 1512-1514.	0.9	2
48	Do Serum Urate-associated Genetic Variants Influence Gout Risk in People Taking Diuretics? Analysis of the UK Biobank. <i>Journal of Rheumatology</i> , 2020, 47, 1704-1711.	2.0	2
49	Response to: "The reference levels of serum urate for clinically evident incident gout"™ by Chen and Ding. <i>Annals of the Rheumatic Diseases</i> , 2019, 78, e42-e42.	0.9	0
50	Longitudinal development of incident gout from low-normal baseline serum urate concentrations: individual participant data analysis. <i>BMC Rheumatology</i> , 2021, 5, 33.	1.6	0
51	Is repeat serum urate testing superior to a single test to predict incident gout over time?. <i>PLoS ONE</i> , 2022, 17, e0263175.	2.5	0