# Amanda Jane Phipps-Green

List of Publications by Year
in descending order



| 51 | 1,583 | 279798 23 | 315739 38 |
|---|---|---|---|
| papers | citations | h-index | g-index |

| 54 | 54 | 54 | 2159 |
|---|---|---|---|
| all docs | docs citations | times ranked | citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | Correspondence on â€˜Variants in urate transporters,ADH1B,GCKRandMEPEgenes associated with transition from asymptomatic hyperuricaemia to gout: results of the first gout versus asymptomatic hyperuricaemia GWAS in Caucasians using data from the UK Biobankâ€™. Annals of the Rheumatic Diseases, 2023, 82, e174-e174. | 0.9 | 3 |
| 2 | Is repeat serum urate testing superior to a single test to predict incident gout over time?. PLoS ONE, 2022, 17, e0263175. | 2.5 | 0 |
| 3 | A Polynesian-specific copy number variant encompassing the MICA gene associates with gout. Human Molecular Genetics, 2022, 31, 3757-3768. | 2.9 | 3 |
| 4 | Trans-ancestral dissection of urate- and gout-associated major loci SLC2A9 and ABCG2 reveals primate-specific regulatory effects. Journal of Human Genetics, 2021, 66, 161-169. | 2.3 | 6 |
| 5 | Gout, Rheumatoid Arthritis, and the Risk of Death Related to Coronavirus Disease 2019: An Analysis of the UK Biobank. ACR Open Rheumatology, 2021, 3, 333-340. | 2.1 | 37 |
| 6 | Potential <scp>PINK1</scp> Founder Effect in Polynesia Causing Earlyâ€Onset Parkinson's Disease. Movement Disorders, 2021, 36, 2199-2200. | 3.9 | 7 |
| 7 | Aotearoa New Zealand MÄori and Pacific Population-amplified Gout Risk Variants: CLNK Is a Separate Risk Gene at the SLC2A9 Locus. Journal of Rheumatology, 2021, 48, 1736-1744. | 2.0 | 8 |
| 8 | Longitudinal development of incident gout from low-normal baseline serum urate concentrations: individual participant data analysis. BMC Rheumatology, 2021, 5, 33. | 1.6 | 0 |
| 9 | Mid-pass whole genome sequencing enables biomedical genetic studies of diverse populations. BMC Genomics, 2021, 22, 666. | 2.8 | 5 |
| 10 | Differential <scp>DNA</scp> Methylation of Networked Signaling, Transcriptional, Innate and Adaptive Immunity, and Osteoclastogenesis Genes and Pathways in Gout. Arthritis and Rheumatology, 2020, 72, 802-814. | 5.6 | 30 |
| 11 | Genetic Polymorphisms on OPRM1, DRD2, DRD4, and COMT in Young Adults: Lack of Association With Alcohol Consumption. Frontiers in Psychiatry, 2020, 11, 549429. | 2.6 | 4 |
| 12 | The ABCG2 Q141K hyperuricemia and gout associated variant illuminates the physiology of human urate excretion. Nature Communications, 2020, 11, 2767. | 12.8 | 71 |
| 13 | Pleiotropic effect of the ABCG2 gene in gout: involvement in serum urate levels and progression from hyperuricemia to gout. Arthritis Research and Therapy, 2020, 22, 45. | 3.5 | 28 |
| 14 | Systematic genetic analysis of early-onset gout: ABCG2 is the only associated locus. Rheumatology, 2020, 59, 2544-2549. | 1.9 | 30 |
| 15 | Do Serum Urateâ€“associated Genetic Variants Influence Gout Risk in People Taking Diuretics? Analysis of the UK Biobank. Journal of Rheumatology, 2020, 47, 1704-1711. | 2.0 | 2 |
| 16 | Population-specific factors associated with fractional excretion of uric acid. Arthritis Research and Therapy, 2019, 21, 234. | 3.5 | 11 |
| 17 | mTOR inhibition by metformin impacts monosodium urate crystalâ€“induced inflammation and cell death in gout: a prelude to a new add-on therapy?. Annals of the Rheumatic Diseases, 2019, 78, 663-671. | 0.9 | 45 |
| 18 | Response to: â€˜The reference levels of serum urate for clinically evident incident goutâ€™ by Chen and Ding. Annals of the Rheumatic Diseases, 2019, 78, e42-e42. | 0.9 | 0 |

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|---|---|---|---|
| 19 | Relationship between serum urate concentration and clinically evident incident gout: an individual participant data analysis. Annals of the Rheumatic Diseases, 2018, 77, 1048-1052. | 0.9 | 131 |
| 20 | The impact of diuretic use and ABCG2 genotype on the predictive performance of a published allopurinol dosing tool. British Journal of Clinical Pharmacology, 2018, 84, 937-943. | 2.4 | 11 |
| 21 | The Oxytocin Receptor Gene (OXTR) Variant rs53576 Is Not Related to Emotional Traits or States in Young Adults. Frontiers in Psychology, 2018, 9, 2548. | 2.1 | 9 |
| 22 | A non-coding genetic variant maximally associated with serum urate levels is functionally linked to HNF4A-dependent PDZK1 expression. Human Molecular Genetics, 2018, 27, 3964-3973. | 2.9 | 26 |
| 23 | ABCG2 loss-of-function polymorphism predicts poor response to allopurinol in patients with gout. Pharmacogenomics Journal, 2017, 17, 201-203. | 2.0 | 82 |
| 24 | Influence of genetic variants on renal uric acid handling in response to frusemide: an acute intervention study. RMD Open, 2017, 3, e000424. | 3.8 | 3 |
| 25 | Populationâ€Specific Resequencing Associates the ATPâ€Binding Cassette Subfamily C Member 4 Gene With Gout in New Zealand MÄori and Pacific Men. Arthritis and Rheumatology, 2017, 69, 1461-1469. | 5.6 | 46 |
| 26 | Population-specific association between ABCG2 variants and tophaceous disease in people with gout. Arthritis Research and Therapy, 2017, 19, 43. | 3.5 | 25 |
| 27 | Clinical and genetic features of diuretic-associated gout: a case-control study. Rheumatology, 2016, 55, 1172-1176. | 1.9 | 5 |
| 28 | Replication of association of the apolipoprotein A1-C3-A4 gene cluster with the risk of gout. Rheumatology, 2016, 55, 1421-1430. | 1.9 | 16 |
| 29 | Association analysis of the beta-3 adrenergic receptor Trp64Arg (rs4994) polymorphism with urate and gout. Rheumatology International, 2016, 36, 255-261. | 3.0 | 10 |
| 30 | Twenty-eight loci that influence serum urate levels: analysis of association with gout. Annals of the Rheumatic Diseases, 2016, 75, 124-130. | 0.9 | 116 |
| 31 | Hyperuricaemia: contributions of urate transporter ABCG2 and the fractional renal clearance of urate. Annals of the Rheumatic Diseases, 2016, 75, 1363-1366. | 0.9 | 30 |
| 32 | Gout Is a Chronic Inflammatory Disease in Which High Levels of Interleukinâ€8 (CXCL8), Myeloidâ€Related Protein 8/Myeloidâ€Related Protein 14 Complex, and an Altered Proteome Are Associated With Diabetes Mellitus and Cardiovascular Disease. Arthritis and Rheumatology, 2015, 67, 3303-3313. | 5.6 | 51 |
| 33 | Body mass index modulates the relationship of sugar-sweetened beverage intake with serum urate concentrations and gout. Arthritis Research and Therapy, 2015, 17, 263. | 3.5 | 24 |
| 34 | Modulation of Genetic Associations with Serum Urate Levels by Body-Mass-Index in Humans. PLoS ONE, 2015, 10, e0119752. | 2.5 | 64 |
| 35 | Association of Autoimmune Addison's Disease with Alleles of STAT4 and GATA3 in European Cohorts. PLoS ONE, 2014, 9, e88991. | 2.5 | 27 |
| 36 | Influence of the ABCG2 gout risk 141ÂK allele on urate metabolism during a fructose challenge. Arthritis Research and Therapy, 2014, 16, R34. | 3.5 | 27 |

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|---|---|---|---|
| 37 | Sugar-sweetened beverage consumption: a risk factor for prevalent gout withSLC2A9genotype-specific effects on serum urate and risk of gout. Annals of the Rheumatic Diseases, 2014, 73, 2101-2106. | 0.9 | 77 |
| 38 | Population-specific effects ofSLC17A1genotype on serum urate concentrations and renal excretion of uric acid during a fructose load. Annals of the Rheumatic Diseases, 2014, 73, 313-314. | 0.9 | 7 |
| 39 | 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. Arthritis Research and Therapy, 2013, 15, R220. | 3.5 | 35 |
| 40 | Population-specific influence ofSLC2A9genotype on the acute hyperuricaemic response to a fructose load. Annals of the Rheumatic Diseases, 2013, 72, 1868-1873. | 0.9 | 61 |
| 41 | Association of the lipoprotein receptor-related protein 2 gene with gout and non-additive interaction with alcohol consumption. Arthritis Research and Therapy, 2013, 15, R177. | 3.5 | 34 |
| 42 | Replication of association of the interleukin 23 receptor rs1343151 variant with rheumatoid arthritis in Caucasian sample sets. Annals of the Rheumatic Diseases, 2012, 71, 155-157. | 0.9 | 13 |
| 43 | The renal urate transporter SLC17A1 locus: confirmation of association with gout. Arthritis Research and Therapy, 2012, 14, R92. | 3.5 | 53 |
| 44 | Smad2:A Candidate Gene for the Murine Autoimmune Diabetes LocusIdd21.1. Journal of Clinical Endocrinology and Metabolism, 2011, 96, E2072-E2077. | 3.6 | 4 |
| 45 | Analysis of association of DNASE2 promoter variation with rheumatoid arthritis in European Caucasians. Annals of the Rheumatic Diseases, 2011, 70, 1512-1514. | 0.9 | 2 |
| 46 | Evidence of interaction of CARD8 rs2043211 with NALP3 rs35829419 in Crohn's disease. Genes and Immunity, 2010, 11, 351-356. | 4.1 | 92 |
| 47 | Association of variation in FcÂ receptor 3B gene copy number with rheumatoid arthritis in Caucasian samples. Annals of the Rheumatic Diseases, 2010, 69, 1711-1716. | 0.9 | 63 |
| 48 | A strong role for the ABCG2 gene in susceptibility to gout in New Zealand Pacific Island and Caucasian, but not MÄori, case and control sample sets. Human Molecular Genetics, 2010, 19, 4813-4819. | 2.9 | 100 |
| 49 | No evidence for association of the systemic lupus erythematosus-associated ITGAM variant, R77H, with rheumatoid arthritis in the Caucasian population. Rheumatology, 2009, 48, 1614-1615. | 1.9 | 7 |
| 50 | The ITGAV rs3738919 variant and susceptibility to rheumatoid arthritis in four Caucasian sample sets. Arthritis Research and Therapy, 2009, 11, R152. | 3.5 | 14 |
| 51 | Immunostimulatory biodegradable implants containing the adjuvant Quil-Aâ€"Part II:In vivoevaluation. Journal of Drug Targeting, 2008, 16, 224-232. | 4.4 | 27 |