Tony R Merriman

List of Publications by Year
in descending order

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273 papers	53,747 citations	25034 57 h-index	1284 225 g-index
298 all docs	298 docs citations	298 times ranked	81628 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	Association of the Quantity, Duration, and Type of Alcohol Consumption on the Development of Gouty Tophi. Arthritis Care and Research, 2023, 75, 1079-1087.	3.4	2
3	Genetic association studies of the progression from hyperuricaemia to gout. Rheumatology, 2022, , .	1.9	2
4	Gout and the risk of COVID-19 diagnosis and death in the UK Biobank: a population-based study. Lancet Rheumatology, The, 2022, 4, e274-e281.	3.9	19
5	Is repeat serum urate testing superior to a single test to predict incident gout over time?. PLoS ONE, 2022, 17, e0263175.	2.5	0
6	CREBRF missense variant rs373863828 has both direct and indirect effects on type 2 diabetes and fasting glucose in Polynesian peoples living in Samoa and Aotearoa New Zealand. BMJ Open Diabetes Research and Care, 2022, 10, e002275.	2.8	2
7	A machine learning-assisted model for renal urate underexcretion with genetic and clinical variables among Chinese men with gout. Arthritis Research and Therapy, 2022, 24, 67.	3.5	4
8	Racial Differences in XO (Xanthine Oxidase) and Mitochondrial DNA Damage-Associated Molecular Patterns in Resistant Hypertension. Hypertension, 2022, 79, 775-784.	2.7	4
9	The minor allele of the CREBRF rs373863828 p.R457Q coding variant is associated with reduced levels of myostatin in males: Implications for body composition. Molecular Metabolism, 2022, 59, 101464.	6.5	2
10	A Polynesian-specific copy number variant encompassing the MICA gene associates with gout. Human Molecular Genetics, 2022, 31, 3757-3768.	2.9	3
11	Is Rheumatoid Arthritis a Causal Factor in Cardiovascular Disease?. Arthritis and Rheumatology, 2022, 74, 1612-1614.	5.6	1
12	Risk factors for acute rheumatic fever: A case-control study. The Lancet Regional Health - Western Pacific, 2022, 26, 100508.	2.9	9
13	Change in serum urate level with urate-lowering therapy initiation associates in the immediate term with patient-reported outcomes in people with gout. Seminars in Arthritis and Rheumatism, 2022, 56, 152057.	3.4	2
14	Trends in the manifestations of 9754 gout patients in a Chinese clinical center: A 10-year observational study. Joint Bone Spine, 2021, 88, 105078.	1.6	15
15	The efficacy and safety of citrate mixture vs sodium bicarbonate on urine alkalization in Chinese primary gout patients with benzbromarone: a prospective, randomized controlled study. Rheumatology, 2021, 60, 2661-2671.	1.9	6
16	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
17	Common variants of EDA are associated with nonâ€syndromic hypodontia. Orthodontics and Craniofacial Research, 2021, 24, 155-163.	2.8	5
18	The comparative effect of exposure to various risk factors on the risk of hyperuricaemia: diet has a weak causal effect. Arthritis Research and Therapy, 2021, 23, 75.	3.5	19

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19	Variable expression quantitative trait loci analysis of breast cancer risk variants. Scientific Reports, 2021, 11, 7192.	3.3	6
20	Effects of fenofibrate therapy on renal function in primary gout patients. Rheumatology, 2021, 60, 5020-5027.	1.9	2
21	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
22	Mapping pleiotropic loci using a fast-sequential testing algorithm. European Journal of Human Genetics, 2021, 29, 1762-1773.	2.8	1
23	Potential <scp>PINK1</scp> Founder Effect in Polynesia Causing Earlyâ€Onset Parkinson's Disease. Movement Disorders, 2021, 36, 2199-2200.	3.9	7
24	Higher Serum Urate Levels Are Associated With an Increased Risk for Sudden Cardiac Death. Journal of Rheumatology, 2021, 48, 1745-1753.	2.0	3
25	Elevated Urate Levels Do Not Alter Bone Turnover Markers: Randomized Controlled Trial of Inosine Supplementation in Postmenopausal Women. Arthritis and Rheumatology, 2021, 73, 1758-1764.	5.6	5
26	Manifestations de la goutte chez 9Â754Âpatients dâ€™un centre clinique chinoisÂ: Ã©tude observationnelle sur 10Âans. Revue Du Rhumatisme (Edition Francaise), 2021, 89, 65-65.	0.0	0
27	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
28	Longitudinal development of incident gout from low-normal baseline serum urate concentrations: individual participant data analysis. BMC Rheumatology, 2021, 5, 33.	1.6	0
29	Assessing the Relationship Between Serum Urate and Urolithiasis Using Mendelian Randomization: An Analysis of the UK Biobank. American Journal of Kidney Diseases, 2021, 78, 210-218.	1.9	8
30	The CREBRF diabetes-protective rs373863828-A allele is associated with enhanced early insulin release in men of MÄori and Pacific ancestry. Diabetologia, 2021, 64, 2779-2789.	6.3	7
31	Factors associated with orthodontic pain. Journal of Oral Rehabilitation, 2021, 48, 1135-1143.	3.0	11
32	Genetic and Physiological Effects of Insulin on Human Urate Homeostasis. Frontiers in Physiology, 2021, 12, 713710.	2.8	17
33	Serum Metabolomics Identifies Dysregulated Pathways and Potential Metabolic Biomarkers for Hyperuricemia and Gout. Arthritis and Rheumatology, 2021, 73, 1738-1748.	5.6	49
34	Assessing the Causal Relationships Between Insulin Resistance and Hyperuricemia and Gout Using Bidirectional Mendelian Randomization. Arthritis and Rheumatology, 2021, 73, 2096-2104.	5.6	49
35	Association of low-level environmental exposure to cadmium and lead with gout flare using a cohort study design. Chemosphere, 2021, 280, 130648.	8.2	3
36	Genetic correlations between traits associated with hyperuricemia, gout, and comorbidities. European Journal of Human Genetics, 2021, 29, 1438-1445.	2.8	11

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37	Mid-pass whole genome sequencing enables biomedical genetic studies of diverse populations. BMC Genomics, 2021, 22, 666.	2.8	5
38	The genetic basis of urate control and gout: Insights into molecular pathogenesis from follow-up study of genome-wide association study loci. Best Practice and Research in Clinical Rheumatology, 2021, 35, 101721.	3.3	8
39	Effect of Clinical Typing on Serum Urate Targets of Benzbromarone in Chinese Gout Patients: A Prospective Cohort Study. Frontiers in Medicine, 2021, 8, 806710.	2.6	6
40	The MÄori and Pacific specific CREBRF variant and adult height. International Journal of Obesity, 2020, 44, 748-752.	3.4	15
41	Relationships Between Allopurinol Dose, Oxypurinol Concentration and Urateâ€Lowering Responseâ€"In Search of a Minimum Effective Oxypurinol Concentration. Clinical and Translational Science, 2020, 13, 110-115.	3.1	6
42	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
43	Nonsynonymous SNPs in LPA homologous to plasminogen deficiency mutants represent novel null apo(a) alleles. Journal of Lipid Research, 2020, 61, 432-444.	4.2	17
44	Comorbidities in gout and hyperuricemia: causality or epiphenomena?. Current Opinion in Rheumatology, 2020, 32, 126-133.	4.3	23
45	Genetic testing in Polynesian long QT syndrome probands reveals a lower diagnostic yield and an increased prevalence of rare variants. Heart Rhythm, 2020, 17, 1304-1311.	0.7	3
46	Do Serum Urateâ€"Associated Genetic Variants Differentially Contribute to Gout Risk According to Body Mass Index? Analysis of the UK Biobank. Arthritis and Rheumatology, 2020, 72, 1184-1191.	5.6	10
47	Urate in fingernail represents the deposition of urate burden in gout patients. Scientific Reports, 2020, 10, 15575.	3.3	6
48	Randomised cross-over trial of vildagliptin and pioglitazone as add-on therapy in patients with type 2 diabetes: predicting Which One is Right Here (WORTH) study protocol. BMJ Open, 2020, 10, e036518.	1.9	2
49	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
50	Effect of body mass index on serum urate and renal uric acid handling responses to an oral inosine load: experimental intervention study in healthy volunteers. Arthritis Research and Therapy, 2020, 22, 259.	3.5	11
51	The ABCG2 Q141K hyperuricemia and gout associated variant illuminates the physiology of human urate excretion. Nature Communications, 2020, 11, 2767.	12.8	71
52	Rare genetic variants in interleukin-37 link this anti-inflammatory cytokine to the pathogenesis and treatment of gout. Annals of the Rheumatic Diseases, 2020, 79, 536-544.	0.9	44
53	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
54	Advances in our understanding of gout as an auto-inflammatory disease. Seminars in Arthritis and Rheumatism, 2020, 50, 1089-1100.	3.4	35

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55	The Pacific-specific CREBRF rs373863828 allele protects against gestational diabetes mellitus in MÄori and Pacific women with obesity. Diabetologia, 2020, 63, 2169-2176.	6.3	14
56	Genomic dissection of 43 serum urate-associated loci provides multiple insights into molecular mechanisms of urate control. Human Molecular Genetics, 2020, 29, 923-943.	2.9	40
57	Systematic genetic analysis of early-onset gout: ABCG2 is the only associated locus. Rheumatology, 2020, 59, 2544-2549.	1.9	30
58	Subtype-specific gout susceptibility loci and enrichment of selection pressure on ABCG2 and ALDH2 identified by subtype genome-wide meta-analyses of clinically defined gout patients. Annals of the Rheumatic Diseases, 2020, 79, 657-665.	0.9	24
59	The Shared Genetic Basis of Hyperuricemia, Gout, and Kidney Function. Seminars in Nephrology, 2020, 40, 586-599.	1.6	10
60	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
61	Gout, Hyperuricemia, and Crystalâ€Associated Disease Network Consensus Statement Regarding Labels and Definitions for Disease Elements in Gout. Arthritis Care and Research, 2019, 71, 427-434.	3.4	73
62	Genome-wide association study revealed novel loci which aggravate asymptomatic hyperuricaemia into gout. Annals of the Rheumatic Diseases, 2019, 78, 1430-1437.	0.9	73
63	Associations of autozygosity with a broad range of human phenotypes. Nature Communications, 2019, 10, 4957.	12.8	84
64	Population-specific factors associated with fractional excretion of uric acid. Arthritis Research and Therapy, 2019, 21, 234.	3.5	11
65	Gout, Hyperuricaemia and Crystal-Associated Disease Network (G-CAN) consensus statement regarding labels and definitions of disease states of gout. Annals of the Rheumatic Diseases, 2019, 78, 1592-1600.	0.9	72
66	Urateâ€lowering therapy alleviates atherosclerosis inflammatory response factors and neointimal lesions in a mouse model of induced carotid atherosclerosis. FEBS Journal, 2019, 286, 1346-1359.	4.7	22
67	Genetic advances in gout: potential applications in clinical practice. Current Opinion in Rheumatology, 2019, 31, 144-151.	4.3	15
68	Mouse models for human hyperuricaemia: a critical review. Nature Reviews Rheumatology, 2019, 15, 413-426.	8.0	99
69	Are Liquid Sugars Different from Solid Sugar in Their Ability to Cause Metabolic Syndrome?. Obesity, 2019, 27, 879-887.	3.0	60
70	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
71	What predicts regression from pre-diabetes to normal glucose regulation following a primary care nurse-delivered dietary intervention? A study protocol for a prospective cohort study. BMJ Open, 2019, 9, e033358.	1.9	4
72	Risk Factors for Acute Rheumatic Fever: Literature Review and Protocol for a Case-Control Study in New Zealand. International Journal of Environmental Research and Public Health, 2019, 16, 4515.	2.6	49

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73	No association between ATP-binding cassette transporter G2 rs2231142 (Q141K) and urate-lowering response to febuxostat. Rheumatology, 2019, 58, 547-548.	1.9	6
74	No causal effects of serum urate levels on the risk of chronic kidney disease: A Mendelian randomization study. PLoS Medicine, 2019, 16, e1002725.	8.4	97
75	Interactions between serum urate-associated genetic variants and sex on gout risk: analysis of the UK Biobank. Arthritis Research and Therapy, 2019, 21, 13.	3.5	19
76	Greater insulin response to acute fructose ingestion among MÄori and Pacific people compared to European people living in Aotearoa New Zealand. Internal Medicine Journal, 2019, 49, 196-202.	0.8	3
77	Genetics of Hyperuricemia and Gout. , 2019, , 9-27.		0
78	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
79	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
80	Association between ABCG2 rs2231142 and poor response to allopurinol: replication and meta-analysis. Rheumatology, 2018, 57, 656-660.	1.9	34
81	Hyperuricemia, Acute and Chronic Kidney Disease, Hypertension, and Cardiovascular Disease: Report of a Scientific Workshop Organized by the National Kidney Foundation. American Journal of Kidney Diseases, 2018, 71, 851-865.	1.9	362
82	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
83	Mitochondrial genetic variation and gout in MÄori and Pacific people living in Aotearoa New Zealand. Annals of the Rheumatic Diseases, 2018, 77, 571-578.	0.9	30
84	Elevated serum uric acid levels are associated with endothelial dysfunction in HIV patients receiving highly-active antiretroviral therapy. Atherosclerosis, 2018, 272, 101-107.	0.8	11
85	Plasma oxypurinol as a measure of adherence in clinical trials. Annals of the Rheumatic Diseases, 2018, 77, 313-314.	0.9	9
86	An association of smoking with serum urate and gout: A health paradox. Seminars in Arthritis and Rheumatism, 2018, 47, 825-842.	3.4	27
87	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
88	ABCG2 rs2231142 (Q141K) and oxypurinol concentrations in people with gout receiving allopurinol. Drug Metabolism and Pharmacokinetics, 2018, 33, 241-242.	2.2	7
89	Expert opinion on emerging urate-lowering therapies. Expert Opinion on Emerging Drugs, 2018, 23, 201-209.	2.4	22
90	Evaluation of the diet wide contribution to serum urate levels: meta-analysis of population based cohorts. BMJ: British Medical Journal, 2018, 363, k3951.	2.3	139

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91	Copy number variants implicate cardiac function and development pathways in earthquake-induced stress cardiomyopathy. Scientific Reports, 2018, 8, 7548.	3.3	8
92	Mediation analysis to understand genetic relationships between habitual coffee intake and gout. Arthritis Research and Therapy, 2018, 20, 135.	3.5	16
93	Pharmaceutical interventions for weight-loss maintenance: no effect from cabergoline. International Journal of Obesity, 2018, 42, 1871-1879.	3.4	7
94	Re: â€œWidespread prevalence of a CREBRF variant among MÄori and Pacific children is associated with weight and height in early childhoodâ€. International Journal of Obesity, 2018, 42, 1389-1391.	3.4	5
95	Multiplexed Nanopore Sequencing of HLA-B Locus in MÄori and Pacific Island Samples. Frontiers in Genetics, 2018, 9, 152.	2.3	17
96	An update on the genetics of hyperuricaemia and gout. Nature Reviews Rheumatology, 2018, 14, 341-353.	8.0	186
97	Ecological momentary assessment of pain in adolescents undergoing orthodontic treatment using a smartphone app. Seminars in Orthodontics, 2018, 24, 209-216.	1.4	4
98	Application of Genetic Epidemiology to CETP (Cholesteryl Ester Transfer Protein) Concentration and Risk of Cardiovascular Disease. Circulation Genomic and Precision Medicine, 2018, 11, e002138.	3.6	4
99	Untangling the complex relationships between incident gout risk, serum urate, and its comorbidities. Arthritis Research and Therapy, 2018, 20, 90.	3.5	16
100	Discordant association of the CREBRF rs373863828 A allele with increased BMI and protection from type 2 diabetes in MÄori and Pacific (Polynesian) people living in Aotearoa/New Zealand. Diabetologia, 2018, 61, 1603-1613.	6.3	61
101	The relationship between ferritin and urate levels and risk of gout. Arthritis Research and Therapy, 2018, 20, 179.	3.5	23
102	Testicular Cancer in New Zealand (TCNZ) study: protocol for a national caseâ€“control study. BMJ Open, 2018, 8, e025212.	1.9	1
103	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
104	Cardio-metabolic disease genetic risk factors among MÄori and Pacific Island people in Aotearoa New Zealand: current state of knowledge and future directions. Annals of Human Biology, 2018, 45, 202-214.	1.0	17
105	Functional Urate-Associated Genetic Variants Influence Expression of lincRNAs LINC01229 and MAFTRR. Frontiers in Genetics, 2018, 9, 733.	2.3	18
106	Association of Crohnâ€™s disease-related chromosome 1q32 with ankylosing spondylitis is independent of bowel symptoms and faecal calprotectin. PeerJ, 2018, 6, e5088.	2.0	4
107	Genomic medicine must reduce, not compound, health inequities: the case for hauora-enhancing genomic resources for New Zealand. New Zealand Medical Journal, 2018, 131, 81-89.	0.5	13
108	PPARGC1B: insight into the expression of the gouty inflammation phenotype. Rheumatology, 2017, 56, kew453.	1.9	3

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109	Testing the Validity of Taxonic Schizotypy Using Genetic and Environmental Risk Variables. Schizophrenia Bulletin, 2017, 43, sbw108.	4.3	28
110	ABCG2 loss-of-function polymorphism predicts poor response to allopurinol in patients with gout. Pharmacogenomics Journal, 2017, 17, 201-203.	2.0	82
111	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
112	Meta-Analysis of Genome-Wide Association Studies for Abdominal Aortic Aneurysm Identifies Four New Disease-Specific Risk Loci. Circulation Research, 2017, 120, 341-353.	4.5	166
113	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
114	Population-specific association between ABCG2 variants and tophaceous disease in people with gout. Arthritis Research and Therapy, 2017, 19, 43.	3.5	25
115	GWAS of clinically defined gout and subtypes identifies multiple susceptibility loci that include urate transporter genes. Annals of the Rheumatic Diseases, 2017, 76, 869-877.	0.9	114
116	Dietary Sodium Modifies Serum Uric Acid Concentrations in Humans. American Journal of Hypertension, 2017, 30, 1196-1202.	2.0	11
117	Multiple common and rare variants of ABCG2 cause gout. RMD Open, 2017, 3, e000464.	3.8	46
118	Geo-epidemiology of temporal artery biopsy-positive giant cell arteritis in Australia and New Zealand: is there a seasonal influence?. RMD Open, 2017, 3, e000531.	3.8	18
119	Functional non-synonymous variants of ABCG2 and gout risk. Rheumatology, 2017, 56, 1982-1992.	1.9	62
120	Genomic Influences on Hyperuricemia and Gout. Rheumatic Disease Clinics of North America, 2017, 43, 389-399.	1.9	16
121	Genotypic variability based association identifies novel non-additive loci DHCR7 and IRF4 in sero-negative rheumatoid arthritis. Scientific Reports, 2017, 7, 5261.	3.3	20
122	Risk factors for cryptorchidism. Nature Reviews Urology, 2017, 14, 534-548.	3.8	93
123	The genetics of gout: towards personalised medicine?. BMC Medicine, 2017, 15, 108.	5.5	44
124	Hypodontia: An Update on Its Etiology, Classification, and Clinical Management. BioMed Research International, 2017, 2017, 1-9.	1.9	121
125	ABCG2 polymorphisms in gout: insights into disease susceptibility and treatment approaches. Pharmacogenomics and Personalized Medicine, 2017, Volume 10, 129-142.	0.7	63
126	Interaction of the GCKR and A1CF loci with alcohol consumption to influence the risk of gout. Arthritis Research and Therapy, 2017, 19, 161.	3.5	29

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127	Performance of gout definitions for genetic epidemiological studies: analysis of UK Biobank. Arthritis Research and Therapy, 2017, 19, 181.	3.5	44
128	SRBreak: A Read-Depth and Split-Read Framework to Identify Breakpoints of Different Events Inside Simple Copy-Number Variable Regions. Frontiers in Genetics, 2016, 7, 160.	2.3	7
129	Association study involving polymorphisms in IL-6, IL-1RA, and CTLA4 genes and rheumatic heart disease in New Zealand population of MÄori and Pacific ancestry. Cytokine, 2016, 85, 201-206.	3.2	13
130	Predicting allopurinol response in patients with gout. British Journal of Clinical Pharmacology, 2016, 81, 277-289.	2.4	46
131	PTPN22R620W minor allele is a genetic risk factor for giant cell arteritis. RMD Open, 2016, 2, e000246.	3.8	9
132	Gout. Lancet, The, 2016, 388, 2039-2052.	13.7	774
133	Clinical and genetic features of diuretic-associated gout: a case-control study. Rheumatology, 2016, 55, 1172-1176.	1.9	5
134	Replication of association of the apolipoprotein A1-C3-A4 gene cluster with the risk of gout. Rheumatology, 2016, 55, 1421-1430.	1.9	16
135	Insight into rheumatological cause and effect through the use of Mendelian randomization. Nature Reviews Rheumatology, 2016, 12, 486-496.	8.0	46
136	Shared Genetic Risk Factors of Intracranial, Abdominal, and Thoracic Aneurysms. Journal of the American Heart Association, 2016, 5, .	3.7	45
137	Lack of direct evidence for natural selection at the candidate thrifty gene locus, PPARGC1A. BMC Medical Genetics, 2016, 17, 80.	2.1	10
138	Association analysis of the beta-3 adrenergic receptor Trp64Arg (rs4994) polymorphism with urate and gout. Rheumatology International, 2016, 36, 255-261.	3.0	10
139	Brief Report: IRF4 Newly Identified as a Common Susceptibility Locus for Systemic Sclerosis and Rheumatoid Arthritis in a Crossâ€Disease Metaâ€Analysis of Genomeâ€Wide Association Studies. Arthritis and Rheumatology, 2016, 68, 2338-2344.	5.6	46
140	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
141	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
142	The Toll-Like Receptor 4 (TLR4) Variant rs2149356 and Risk of Gout in European and Polynesian Sample Sets. PLoS ONE, 2016, 11, e0147939.	2.5	31
143	Mendelian Randomization Analysis to Examine for a Causal Effect of Urate on Bone Mineral Density. Journal of Bone and Mineral Research, 2015, 30, 985-991.	2.8	50
144	The distribution and impact of common copy-number variation in the genome of the domesticated apple, Malus x domestica Borkh. BMC Genomics, 2015, 16, 848.	2.8	21

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145	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
146	Pacific Populations, Metabolic Disease and â€˜Justâ€So Storiesâ€™: A Critique of the â€˜Thrifty Genotypeâ€™ Hypothesis in Oceania. Annals of Human Genetics, 2015, 79, 470-480.	0.8	35
147	A human leukocyte antigen locus haplotype confers risk for allopurinol-related adverse effects in Caucasian patients with gout. Pharmacogenetics and Genomics, 2015, 25, 412-415.	1.5	7
148	Association of SLC2A9 genotype with phenotypic variability of serum urate in pre-menopausal women. Frontiers in Genetics, 2015, 6, 313.	2.3	16
149	Sugar Sweetened Beverage Consumption among Adults with Gout or Type 2 Diabetes. PLoS ONE, 2015, 10, e0125543.	2.5	10
150	A genome-wide association study of rheumatoid arthritis without antibodies against citrullinated peptides. Annals of the Rheumatic Diseases, 2015, 74, e15-e15.	0.9	62
151	Does midlife obesity really lower dementia risk?. Lancet Diabetes and Endocrinology,the, 2015, 3, 501.	11.4	2
152	An update on the genetic architecture of hyperuricemia and gout. Arthritis Research and Therapy, 2015, 17, 98.	3.5	123
153	Positive association of tomato consumption with serum urate: support for tomato consumption as an anecdotal trigger of gout flares. BMC Musculoskeletal Disorders, 2015, 16, 196.	1.9	27
154	Causal or Noncausal Relationship of Uric Acid With Diabetes. Diabetes, 2015, 64, 2720-2722.	0.6	36
155	Smoking behaviour modifies IL23râ€associated disease risk in patients with Crohn's disease. Journal of Gastroenterology and Hepatology (Australia), 2015, 30, 299-307.	2.8	18
156	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
157	Lack of geneâ€“diuretic interactions on the risk of incident gout: the Nursesâ€™ Health Study and Health Professionals Follow-up Study. Annals of the Rheumatic Diseases, 2015, 74, 1394-1398.	0.9	18
158	Multiplicative interaction of functional inflammasome genetic variants in determining the risk of gout. Arthritis Research and Therapy, 2015, 17, 288.	3.5	54
159	Modulation of Genetic Associations with Serum Urate Levels by Body-Mass-Index in Humans. PLoS ONE, 2015, 10, e0119752.	2.5	64
160	Association of Autoimmune Addison's Disease with Alleles of STAT4 and GATA3 in European Cohorts. PLoS ONE, 2014, 9, e88991.	2.5	27
161	Mendelian randomization analysis associates increased serum urate, due to genetic variation in uric acid transporters, with improved renal function. Kidney International, 2014, 85, 344-351.	5.2	78
162	The CNVrd2 package: measurement of copy number at complex loci using high-throughput sequencing data. Frontiers in Genetics, 2014, 5, 248.	2.3	22

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164	Abundant local interactions in the 4p16.1 region suggest functional mechanisms underlying SLC2A9 associations with human serum uric acid. Human Molecular Genetics, 2014, 23, 5061-5068.	2.9	29
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