Tony R Merriman

List of Publications by Year
in descending order

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	Global and regional mortality from 235 causes of death for 20 age groups in 1990 and 2010: a systematic analysis for the Global Burden of Disease Study 2010. Lancet, The, 2012, 380, 2095-2128.	13.7	11,038
2	A comparative risk assessment of burden of disease and injury attributable to 67 risk factors and risk factor clusters in 21 regions, 1990â€"2010: a systematic analysis for the Global Burden of Disease Study 2010. Lancet, The, 2012, 380, 2224-2260.	13.7	9,397
3	Global, regional, and national prevalence of overweight and obesity in children and adults during 1980â€"2013: a systematic analysis for the Global Burden of Disease Study 2013. Lancet, The, 2014, 384, 766-781.	13.7	9,122
4	Disability-adjusted life years (DALYs) for 291 diseases and injuries in 21 regions, 1990â€"2010: a systematic analysis for the Global Burden of Disease Study 2010. Lancet, The, 2012, 380, 2197-2223.	13.7	7,061
5	Years lived with disability (YLDs) for 1160 sequelae of 289 diseases and injuries 1990â€"2010: a systematic analysis for the Global Burden of Disease Study 2010. Lancet, The, 2012, 380, 2163-2196.	13.7	6,376
6	Gout. Lancet, The, 2016, 388, 2039-2052.	13.7	774
7	Genome-wide association study identifies new multiple sclerosis susceptibility loci on chromosomes 12 and 20. Nature Genetics, 2009, 41, 824-828.	21.4	501
8	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
9	The global burden of gout: estimates from the Global Burden of Disease 2010 study. Annals of the Rheumatic Diseases, 2014, 73, 1470-1476.	0.9	206
10	The genetically isolated populations of Finland and Sardinia may not be a panacea for linkage disequilibrium mapping of common disease genes. Nature Genetics, 2000, 25, 320-323.	21.4	186
11	An update on the genetics of hyperuricaemia and gout. Nature Reviews Rheumatology, 2018, 14, 341-353.	8.0	186
12	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
13	The genetic basis of hyperuricaemia and gout. Joint Bone Spine, 2011, 78, 35-40.	1.6	143
14	Confirmation of association of IRGM and NCF4 with ileal Crohn's disease in a population-based cohort. Genes and Immunity, 2008, 9, 561-565.	4.1	142
15	Evidence for an influence of chemokine ligand 3-like 1 (CCL3L1) gene copy number on susceptibility to rheumatoid arthritis. Annals of the Rheumatic Diseases, 2007, 67, 409-413.	0.9	139
16	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
17	Association analysis of the interleukin 17A gene in Caucasian rheumatoid arthritis patients from Norway and New Zealand. Rheumatology, 2009, 48, 367-370.	1.9	133
18	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

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19	An update on the genetic architecture of hyperuricemia and gout. Arthritis Research and Therapy, 2015, 17, 98.	3.5	123
20	Hypodontia: An Update on Its Etiology, Classification, and Clinical Management. BioMed Research International, 2017, 2017, 1-9.	1.9	121
21	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
22	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
23	IL23R R381Q and ATG16L1 T300A Are Strongly Associated With Crohn's Disease in a Study of New Zealand Caucasians With Inflammatory Bowel Disease. American Journal of Gastroenterology, 2007, 102, 2754-2761.	0.4	109
24	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
25	Mouse models for human hyperuricaemia: a critical review. Nature Reviews Rheumatology, 2019, 15, 413-426.	8.0	99
26	Role of the urate transporter <i>SLC2A9</i> gene in susceptibility to gout in New Zealand MÄori, Pacific Island, and Caucasian caseâ€“control sample sets. Arthritis and Rheumatism, 2009, 60, 3485-3492.	6.7	98
27	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
28	Risk factors for cryptorchidism. Nature Reviews Urology, 2017, 14, 534-548.	3.8	93
29	Evidence of interaction of CARD8 rs2043211 with NALP3 rs35829419 in Crohn's disease. Genes and Immunity, 2010, 11, 351-356.	4.1	92
30	Association of CDH1 haplotypes with susceptibility to sporadic diffuse gastric cancer. Oncogene, 2002, 21, 8192-8195.	5.9	91
31	A sequence variant associated with sortilin-1 (SORT1) on 1p13.3 is independently associated with abdominal aortic aneurysm. Human Molecular Genetics, 2013, 22, 2941-2947.	2.9	88
32	Prevalence of airway and parenchymal abnormalities in newly diagnosed rheumatoid arthritis. Respiratory Medicine, 2012, 106, 1441-1446.	2.9	87
33	The molecular basis of the Kidd blood group polymorphism and its lack of association with type 1 diabetes susceptibility. Human Molecular Genetics, 1997, 6, 1017-1020.	2.9	85
34	Associations of autozygosity with a broad range of human phenotypes. Nature Communications, 2019, 10, 4957.	12.8	84
35	Analysis of the Fc Receptor-Like-3 (FCRL3) Locus in Caucasians with Autoimmune Disorders Suggests a Complex Pattern of Disease Association. Journal of Clinical Endocrinology and Metabolism, 2007, 92, 1106-1111.	3.6	83
36	Association of Higher DEFB4 Genomic Copy Number With Crohn's Disease. American Journal of Gastroenterology, 2010, 105, 354-359.	0.4	83

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37	ABCG2 loss-of-function polymorphism predicts poor response to allopurinol in patients with gout. Pharmacogenomics Journal, 2017, 17, 201-203.	2.0	82
38	Evidence by allelic association-dependent methods for a type 1 diabetes polygene (IDDM6) on chromosome 18q21. Human Molecular Genetics, 1997, 6, 1003-1010.	2.9	81
39	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
40	Sugar-sweetened beverage consumption: a risk factor for prevalent gout with<i>SLC2A9</i>genotype-specific effects on serum urate and risk of gout. Annals of the Rheumatic Diseases, 2014, 73, 2101-2106.	0.9	77
41	Association of thePTPN22 locus with rheumatoid arthritis in a New Zealand Caucasian cohort. Arthritis and Rheumatism, 2005, 52, 2222-2225.	6.7	75
42	Differential association of two PTPN22 coding variants with CrohnÊ¼s disease and ulcerative colitis. Inflammatory Bowel Diseases, 2011, 17, 2287-2294.	1.9	73
43	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
44	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
45	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
46	Genomic DNA pooling for whole-genome association scans in complex disease: empirical demonstration of efficacy in rheumatoid arthritis. Genes and Immunity, 2007, 8, 57-68.	4.1	71
47	<i>TLR2</i>, <i>TLR4</i> and <i>TLR9</i> polymorphisms and Crohn's disease in a New Zealand Caucasian cohort. Journal of Gastroenterology and Hepatology (Australia), 2007, 22, 1760-1766.	2.8	71
48	The ABCG2 Q141K hyperuricemia and gout associated variant illuminates the physiology of human urate excretion. Nature Communications, 2020, 11, 2767.	12.8	71
49	Genetic progress towards the molecular basis of autoimmunity. Trends in Molecular Medicine, 2006, 12, 90-98.	6.7	69
50	Hospital admissions associated with gout and their comorbidities in New Zealand and England 1999-2009. Rheumatology, 2013, 52, 118-126.	1.9	66
51	The <i>PTPN22</i> R263Q polymorphism is a risk factor for rheumatoid arthritis in Caucasian caseâ€"control samples. Arthritis and Rheumatism, 2011, 63, 365-372.	6.7	64
52	Modulation of Genetic Associations with Serum Urate Levels by Body-Mass-Index in Humans. PLoS ONE, 2015, 10, e0119752.	2.5	64
53	Novel germlineCDH1mutations in hereditary diffuse gastric cancer families. Human Mutation, 2002, 19, 518-525.	2.5	63
54	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

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55	ABCG2 polymorphisms in gout: insights into disease susceptibility and treatment approaches. Pharmacogenomics and Personalized Medicine, 2017, Volume 10, 129-142.	0.7	63
56	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
57	Functional non-synonymous variants of ABCG2 and gout risk. Rheumatology, 2017, 56, 1982-1992.	1.9	62
58	Population-specific influence of<i>SLC2A9</i>genotype on the acute hyperuricaemic response to a fructose load. Annals of the Rheumatic Diseases, 2013, 72, 1868-1873.	0.9	61
59	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
60	Construction and use of a self-cloning promoter probe vector for Gram-negative bacteria. Gene, 1993, 126, 17-23.	2.2	60
61	Are Liquid Sugars Different from Solid Sugar in Their Ability to Cause Metabolic Syndrome?. Obesity, 2019, 27, 879-887.	3.0	60
62	KCNN4 Gene Variant Is Associated With Ileal Crohn's Disease in the Australian and New Zealand Population. American Journal of Gastroenterology, 2010, 105, 2209-2217.	0.4	59
63	Myeloidâ€Related Proteins 8 and 14 Contribute to Monosodium Urate Monohydrate Crystalâ€“Induced Inflammation in Gout. Arthritis and Rheumatology, 2014, 66, 1327-1339.	5.6	58
64	Mutations in the Zinc Finger Protein Gene,<i>ZNF469</i>, Contribute to the Pathogenesis of Keratoconus. , 2014, 55, 5629.		57
65	Fine Mapping of the Diabetes-Susceptibility Locus, IDDM4, on Chromosome 11q13. American Journal of Human Genetics, 1998, 63, 547-556.	6.2	56
66	Multiplicative interaction of functional inflammasome genetic variants in determining the risk of gout. Arthritis Research and Therapy, 2015, 17, 288.	3.5	54
67	Evidence for a type 1 diabetes susceptibility locus (IDDM10) on human chromosome 10p11-q11. Human Molecular Genetics, 1997, 6, 1011-1016.	2.9	53
68	The renal urate transporter SLC17A1 locus: confirmation of association with gout. Arthritis Research and Therapy, 2012, 14, R92.	3.5	53
69	A Genetic Association Study of Serum Acute-Phase C-Reactive Protein Levels in Rheumatoid Arthritis: Implications for Clinical Interpretation. PLoS Medicine, 2010, 7, e1000341.	8.4	52
70	Interaction of the inflammasome genes CARD8 and NLRP3 in abdominal aortic aneurysms. Atherosclerosis, 2011, 218, 123-126.	0.8	52
71	A bioinformatics workflow for detecting signatures of selection in genomic data. Frontiers in Genetics, 2014, 5, 293.	2.3	51
72	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

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73	Genetics of autoimmune disease. Current Opinion in Immunology, 1995, 7, 786-792.	5.5	50
74	Polymorphisms in the organic cation transporter genes SLC22A4 and SLC22A5 and Crohn's disease in a New Zealand Caucasian cohort. Immunology and Cell Biology, 2006, 84, 233-236.	2.3	50
75	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
76	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
77	Serum Metabolomics Identifies Dysregulated Pathways and Potential Metabolic Biomarkers for Hyperuricemia and Gout. Arthritis and Rheumatology, 2021, 73, 1738-1748.	5.6	49
78	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
79	Association of thymidylate synthase polymorphisms with gastric cancer susceptibility. International Journal of Cancer, 2004, 112, 1010-1014.	5.1	46
80	Predicting allopurinol response in patients with gout. British Journal of Clinical Pharmacology, 2016, 81, 277-289.	2.4	46
81	Insight into rheumatological cause and effect through the use of Mendelian randomization. Nature Reviews Rheumatology, 2016, 12, 486-496.	8.0	46
82	Brief Report: <i>IRF4</i> 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
83	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
84	Multiple common and rare variants of <i>ABCG2</i> cause gout. RMD Open, 2017, 3, e000464.	3.8	46
85	Shared Genetic Risk Factors of Intracranial, Abdominal, and Thoracic Aneurysms. Journal of the American Heart Association, 2016, 5, .	3.7	45
86	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
87	The genetics of gout: towards personalised medicine?. BMC Medicine, 2017, 15, 108.	5.5	44
88	Performance of gout definitions for genetic epidemiological studies: analysis of UK Biobank. Arthritis Research and Therapy, 2017, 19, 181.	3.5	44
89	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
90	Meta-analysis confirms a role for deletion in FCGR3B in autoimmune phenotypes. Human Molecular Genetics, 2012, 21, 2370-2376.	2.9	43

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91	IL-23R rs11209026 polymorphism modulates IL-17A expression in patients with rheumatoid arthritis. Genes and Immunity, 2012, 13, 282-287.	4.1	43
92	Impaired response or insufficient dosage?â€"Examining the potential causes of â€œinadequate responseâ€•to allopurinol in the treatment of gout. Seminars in Arthritis and Rheumatism, 2014, 44, 170-174.	3.4	43
93	Transmission of haplotypes of microsatellite markers rather than single marker alleles in the mapping of a putative type 1 diabetes susceptibility gene (IDDM6). Human Molecular Genetics, 1998, 7, 517-524.	2.9	42
94	Evidence for association of an interleukin 23 receptor variant independent of the R381Q variant with rheumatoid arthritis. Annals of the Rheumatic Diseases, 2009, 68, 1340-1344.	0.9	41
95	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
96	Hyperuricaemia in the Pacific: why the elevated serum urate levels?. Rheumatology International, 2014, 34, 743-757.	3.0	37
97	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
98	Mutation of the glucagon receptor gene and diabetes mellitus in the UK: association or founder effect?. Human Molecular Genetics, 1995, 4, 1609-1612.	2.9	36
99	The SLC2A9 nonsynonymous Arg265His variant and gout: evidence for a population-specific effect on severity. Arthritis Research and Therapy, 2011, 13, R85.	3.5	36
100	Causal or Noncausal Relationship of Uric Acid With Diabetes. Diabetes, 2015, 64, 2720-2722.	0.6	36
101	Only one independent genetic association with rheumatoid arthritis within the KIAA1109-TENR-IL2-IL21 locus in Caucasian sample sets: confirmation of association of rs6822844 with rheumatoid arthritis at a genome-wide level of significance. Arthritis Research and Therapy, 2010, 12, R116.	3.5	35
102	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
103	The Genetic Basis of Gout. Rheumatic Disease Clinics of North America, 2014, 40, 279-290.	1.9	35
104	Pacific Populations, Metabolic Disease and â€˜Justâ€Šo Storiesâ€™: A Critique of the â€˜Thrifty Genotypeâ€™ Hypothesis in Oceania. Annals of Human Genetics, 2015, 79, 470-480.	0.8	35
105	Advances in our understanding of gout as an auto-inflammatory disease. Seminars in Arthritis and Rheumatism, 2020, 50, 1089-1100.	3.4	35
106	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
107	Association between ABCG2 rs2231142 and poor response to allopurinol: replication and meta-analysis. Rheumatology, 2018, 57, 656-660.	1.9	34
108	Mendelian Randomization Provides No Evidence for a Causal Role of Serum Urate in Increasing Serum Triglyceride Levels. Circulation: Cardiovascular Genetics, 2014, 7, 830-837.	5.1	33

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109	Risk-taking: behind the warrior gene story. New Zealand Medical Journal, 2007, 120, U2440.	0.5	32
110	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
111	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
112	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
113	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
114	Systematic genetic analysis of early-onset gout: ABCG2 is the only associated locus. Rheumatology, 2020, 59, 2544-2549.	1.9	30
115	Population Heterogeneity in the Genetic Control of Serum Urate. Seminars in Nephrology, 2011, 31, 420-425.	1.6	29
116	Evidence that deletion at FCGR3B is a risk factor for systemic sclerosis. Genes and Immunity, 2012, 13, 458-460.	4.1	29
117	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
118	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
119	Testing the Validity of Taxonic Schizotypy Using Genetic and Environmental Risk Variables. Schizophrenia Bulletin, 2017, 43, sbw108.	4.3	28
120	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
121	Association of Autoimmune Addison's Disease with Alleles of STAT4 and GATA3 in European Cohorts. PLoS ONE, 2014, 9, e88991.	2.5	27
122	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
123	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
124	An association of smoking with serum urate and gout: A health paradox. Seminars in Arthritis and Rheumatism, 2018, 47, 825-842.	3.4	27
125	Maternal Psychological Reaction to Newborn Genetic Screening for Type 1 Diabetes. Pediatrics, 2007, 120, e324-e335.	2.1	26
126	Type 1 diabetes, the A1 milk hypothesis and vitamin D deficiency. Diabetes Research and Clinical Practice, 2009, 83, 149-156.	2.8	26

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127	The population pharmacokinetics of allopurinol and oxypurinol in patients with gout. European Journal of Clinical Pharmacology, 2013, 69, 1411-1421.	1.9	26
128	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
129	Population-specific association between ABCG2 variants and tophaceous disease in people with gout. Arthritis Research and Therapy, 2017, 19, 43.	3.5	25
130	Genetics of Type 1 Diabetes and Autoimmune Thyroid Disease. Endocrinology and Metabolism Clinics of North America, 2009, 38, 289-301.	3.2	24
131	Prevalence of HLA-B27 in the New Zealand population: effect of age and ethnicity. Arthritis Research and Therapy, 2013, 15, R158.	3.5	24
132	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
133	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
134	The relationship between ferritin and urate levels and risk of gout. Arthritis Research and Therapy, 2018, 20, 179.	3.5	23
135	Comorbidities in gout and hyperuricemia: causality or epiphenomena?. Current Opinion in Rheumatology, 2020, 32, 126-133.	4.3	23
136	Polymorphisms of CARD15/NOD2 and CD14 genes in New Zealand Crohn's disease patients. Immunology and Cell Biology, 2005, 83, 498-503.	2.3	22
137	Colocalization of Mouse Autoimmune Diabetes Loci Idd21.1 and Idd21.2 With IDDM6 (Human) and Iddm3 (Rat). Diabetes, 2005, 54, 2820-2825.	0.6	22
138	The CNVrd2 package: measurement of copy number at complex loci using high-throughput sequencing data. Frontiers in Genetics, 2014, 5, 248.	2.3	22
139	The relationship of apolipoprotein B and very low density lipoprotein triglyceride with hyperuricemia and gout. Arthritis Research and Therapy, 2014, 16, 495.	3.5	22
140	Expert opinion on emerging urate-lowering therapies. Expert Opinion on Emerging Drugs, 2018, 23, 201-209.	2.4	22
141	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
142	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
143	Vitamin D receptor gene polymorphism associated with inflammatory bowel disease in New Zealand males. Alimentary Pharmacology and Therapeutics, 2011, 33, 855-856.	3.7	20
144	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

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145	Association of CD247 Polymorphisms with Rheumatoid Arthritis: A Replication Study and a Meta-Analysis. PLoS ONE, 2013, 8, e68295.	2.5	19
146	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
147	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
148	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
149	Gout in MÂori. Rheumatology, 2014, 53, 773-774.	1.9	18
150	Smoking behaviour modifies <i>IL23r</i>â€associated disease risk in patients with Crohn's disease. Journal of Gastroenterology and Hepatology (Australia), 2015, 30, 299-307.	2.8	18
151	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
152	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
153	Functional Urate-Associated Genetic Variants Influence Expression of lincRNAs LINC01229 and MAFTRR. Frontiers in Genetics, 2018, 9, 733.	2.3	18
154	Multiplexed Nanopore Sequencing of HLA-B Locus in MÄori and Pacific Island Samples. Frontiers in Genetics, 2018, 9, 152.	2.3	17
155	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
156	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
157	Genetic and Physiological Effects of Insulin on Human Urate Homeostasis. Frontiers in Physiology, 2021, 12, 713710.	2.8	17
158	Macrophage migration inhibitory factor gene polymorphisms in inflammatory bowel disease: An association study in New Zealand Caucasians and meta-analysis. World Journal of Gastroenterology, 2013, 19, 6656.	3.3	17
159	Association of SLC2A9 genotype with phenotypic variability of serum urate in pre-menopausal women. Frontiers in Genetics, 2015, 6, 313.	2.3	16
160	Replication of association of the apolipoprotein A1-C3-A4 gene cluster with the risk of gout. Rheumatology, 2016, 55, 1421-1430.	1.9	16
161	Genomic Influences on Hyperuricemia and Gout. Rheumatic Disease Clinics of North America, 2017, 43, 389-399.	1.9	16
162	Mediation analysis to understand genetic relationships between habitual coffee intake and gout. Arthritis Research and Therapy, 2018, 20, 135.	3.5	16

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163	Untangling the complex relationships between incident gout risk, serum urate, and its comorbidities. Arthritis Research and Therapy, 2018, 20, 90.	3.5	16
164	Association of FcgR2a, but not FcgR3a, with inflammatory bowel diseases across three Caucasian populationsâ€. Inflammatory Bowel Diseases, 2010, 16, 2080-2089.	1.9	15
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