

# Ana M Valdes

## List of Publications by Year in descending order

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Version: 2024-02-01

287  
papers

31,966  
citations

4641

85  
h-index

5519

163  
g-index

301  
all docs

301  
docs citations

301  
times ranked

42802  
citing authors

#	ARTICLE	IF	CITATIONS
1	Upregulated expression of <i>FFAR2</i> and <i>SOC3</i> genes is associated with gout. <i>Rheumatology</i> , 2023, 62, 977-983.	0.9	4
2	Role of the gut microbiome in chronic diseases: a narrative review. <i>European Journal of Clinical Nutrition</i> , 2022, 76, 489-501.	1.3	168
3	Clinical and Preclinical Evidence for Roles of Soluble Epoxide Hydrolase in Osteoarthritis Knee Pain. <i>Arthritis and Rheumatology</i> , 2022, 74, 623-633.	2.9	10
4	Body mass index mediates the effect of the DASH diet on hypertension: Common metabolites underlying the association. <i>Journal of Human Nutrition and Dietetics</i> , 2022, 35, 214-222.	1.3	6
5	Impact of insufficient sleep on dysregulated blood glucose control under standardised meal conditions. <i>Diabetologia</i> , 2022, 65, 356-365.	2.9	29
6	Pre-existing polymerase-specific T cells expand in abortive seronegative SARS-CoV-2. <i>Nature</i> , 2022, 601, 110-117.	13.7	280
7	Acceptability of a nurse-led non-pharmacological complex intervention for knee pain: Nurse and patient views and experiences. <i>PLoS ONE</i> , 2022, 17, e0262422.	1.1	3
8	The association of socio-economic and psychological factors with limitations in day-to-day activity over 7 years in newly diagnosed osteoarthritis patients. <i>Scientific Reports</i> , 2022, 12, 943.	1.6	4
9	Validity of continuous glucose monitoring for categorizing glycemic responses to diet: implications for use in personalized nutrition. <i>American Journal of Clinical Nutrition</i> , 2022, 115, 1569-1576.	2.2	15
10	Metabolome Genome-Wide Association Study Identifies 74 Novel Genomic Regions Influencing Plasma Metabolites Levels. <i>Metabolites</i> , 2022, 12, 61.	1.3	18
11	Different genes may be involved in distal and local sensitization: A genome-wide gene-based association study and meta-analysis. <i>European Journal of Pain</i> , 2022, 26, 740-753.	1.4	3
12	Incremental Value of a Panel of Serum Metabolites for Predicting Risk of Atherosclerotic Cardiovascular Disease. <i>Journal of the American Heart Association</i> , 2022, 11, e024590.	1.6	1
13	HLA-DR polymorphism in SARS-CoV-2 infection and susceptibility to symptomatic COVID-19. <i>Immunology</i> , 2022, 166, 68-77.	2.0	18
14	Symptom prevalence, duration, and risk of hospital admission in individuals infected with SARS-CoV-2 during periods of omicron and delta variant dominance: a prospective observational study from the ZOE COVID Study. <i>Lancet</i> , 2022, 399, 1618-1624.	6.3	547
15	COVID-19 vaccine waning and effectiveness and side-effects of boosters: a prospective community study from the ZOE COVID Study. <i>Lancet Infectious Diseases</i> , 2022, 22, 1002-1010.	4.6	192
16	Has a change in established care pathways during the first wave of the COVID-19 pandemic led to an excess death rate in the fragility fracture population? A longitudinal cohort study of 1846 patients. <i>BMJ Open</i> , 2022, 12, e058526.	0.8	2
17	Effects of temporarily suspending low-dose methotrexate treatment for 2 weeks after SARS-CoV-2 vaccine booster on vaccine response in immunosuppressed adults with inflammatory conditions: protocol for a multicentre randomised controlled trial and nested mechanistic substudy (Vaccine) <i>Tj ETQq1 1 0.784314 rgBT<sup>3</sup>Overloc</i>	0.8	3
18	Postprandial and Fasting Metabolic Signatures: Insights From the ZOE PREDICT 1 Study. <i>Current Developments in Nutrition</i> , 2022, 6, 448.	0.1	0

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19	Comorbidities and use of analgesics in people with knee pain: a study in the Nottingham Knee Pain and Health in the Community (KPIC) cohort. <i>Rheumatology Advances in Practice</i> , 2022, 6, .	0.3	2
20	Effect of a 2-week interruption in methotrexate treatment versus continued treatment on COVID-19 booster vaccine immunity in adults with inflammatory conditions (VROOM study): a randomised, open label, superiority trial. <i>Lancet Respiratory Medicine</i> , 2022, 10, 840-850.	5.2	52
21	IgG N-glycome changes during the course of severe COVID-19: An observational study. <i>EBioMedicine</i> , 2022, 81, 104101.	2.7	18
22	Perspective: Leveraging the Gut Microbiota to Predict Personalized Responses to Dietary, Prebiotic, and Probiotic Interventions. <i>Advances in Nutrition</i> , 2022, 13, 1450-1461.	2.9	21
23	The role of short-chain fatty acids in the interplay between gut microbiota and diet in cardio-metabolic health. <i>Gut Microbes</i> , 2021, 13, 1-24.	4.3	259
24	Gut Microbial Profile Is Associated With Residential Settings and Not Nutritional Status in Adults in Karnataka, India. <i>Frontiers in Nutrition</i> , 2021, 8, 595756.	1.6	1
25	High intake of vegetables is linked to lower white blood cell profile and the effect is mediated by the gut microbiome. <i>BMC Medicine</i> , 2021, 19, 37.	2.3	30
26	Effectiveness of Internet-Based Exercises Aimed at Treating Knee Osteoarthritis. <i>JAMA Network Open</i> , 2021, 4, e210012.	2.8	59
27	Genome-wide association study in almost 195,000 individuals identifies 50 previously unidentified genetic loci for eye color. <i>Science Advances</i> , 2021, 7, .	4.7	36
28	Metabolic signatures of osteoarthritis in urine using liquid chromatography–high resolution tandem mass spectrometry. <i>Metabolomics</i> , 2021, 17, 29.	1.4	14
29	β-blocker prescription is associated with lower cumulative risk of knee osteoarthritis and knee pain consultations in primary care: a propensity score–matched cohort study. <i>Rheumatology</i> , 2021, 60, 5686-5696.	0.9	10
30	Blue poo: impact of gut transit time on the gut microbiome using a novel marker. <i>Gut</i> , 2021, 70, 1665-1674.	6.1	84
31	Longitudinal assessment of symptoms and risk of SARS-CoV-2 infection in healthcare workers across 5 hospitals to understand ethnic differences in infection risk.. <i>EClinicalMedicine</i> , 2021, 34, 100835.	3.2	20
32	Prior SARS-CoV-2 infection rescues B and T cell responses to variants after first vaccine dose. <i>Science</i> , 2021, 372, 1418-1423.	6.0	286
33	Postprandial glycaemic dips predict appetite and energy intake in healthy individuals. <i>Nature Metabolism</i> , 2021, 3, 523-529.	5.1	47
34	Modest effects of dietary supplements during the COVID-19 pandemic: insights from 445 850 users of the COVID-19 Symptom Study app. <i>BMJ Nutrition, Prevention and Health</i> , 2021, 4, 149-157.	1.9	91
35	Gut microbiome diversity and composition is associated with hypertension in women. <i>Journal of Hypertension</i> , 2021, 39, 1810-1816.	0.3	22
36	Meal-induced inflammation: postprandial insights from the Personalised REsponses to Dietary Composition Trial (PREDICT) study in 1000 participants. <i>American Journal of Clinical Nutrition</i> , 2021, 114, 1028-1038.	2.2	43

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37	Fidelity assessment of nurse-led non-pharmacological package of care for knee pain in the package development phase of a feasibility randomised controlled trial based in secondary care: a mixed methods study. <i>BMJ Open</i> , 2021, 11, e045242.	0.8	4
38	Vaccine side-effects and SARS-CoV-2 infection after vaccination in users of the COVID Symptom Study app in the UK: a prospective observational study. <i>Lancet Infectious Diseases</i> , The, 2021, 21, 939-949.	4.6	744
39	N-glycosylation of immunoglobulin G predicts incident hypertension. <i>Journal of Hypertension</i> , 2021, 39, 2527-2533.	0.3	13
40	Dietary Interventions Reduce Traditional and Novel Cardiovascular Risk Markers by Altering the Gut Microbiome and Their Metabolites. <i>Frontiers in Cardiovascular Medicine</i> , 2021, 8, 691564.	1.1	25
41	Circulating Levels of the Short-Chain Fatty Acid Acetate Mediate the Effect of the Gut Microbiome on Visceral Fat. <i>Frontiers in Microbiology</i> , 2021, 12, 711359.	1.5	86
42	Two doses of the SARS-CoV-2 BNT162b2 vaccine enhance antibody responses to variants in individuals with prior SARS-CoV-2 infection. <i>Science Translational Medicine</i> , 2021, 13, eabj0847.	5.8	40
43	Deciphering osteoarthritis genetics across 826,690 individuals from 9 populations. <i>Cell</i> , 2021, 184, 4784-4818.e17.	13.5	188
44	Microbiome connections with host metabolism and habitual diet from 1,098 deeply phenotyped individuals. <i>Nature Medicine</i> , 2021, 27, 321-332.	15.2	477
45	The prebiotic effects of omega-3 fatty acid supplementation: A six-week randomised intervention trial. <i>Gut Microbes</i> , 2021, 13, 1-11.	4.3	63
46	Effects of an isoenergetic low Glycaemic Index (GI) diet on liver fat accumulation and gut microbiota composition in patients with non-alcoholic fatty liver disease (NAFLD): a study protocol of an efficacy mechanism evaluation. <i>BMJ Open</i> , 2021, 11, e045802.	0.8	2
47	The anti-inflammatory effect of bacterial short chain fatty acids is partially mediated by endocannabinoids. <i>Gut Microbes</i> , 2021, 13, 1997559.	4.3	34
48	A High Protein Diet Is More Effective in Improving Insulin Resistance and Glycemic Variability Compared to a Mediterranean Diet—A Cross-Over Controlled Inpatient Dietary Study. <i>Nutrients</i> , 2021, 13, 4380.	1.7	25
49	Metabolic syndrome and osteoarthritis pain: common molecular mechanisms and potential therapeutic implications. <i>Osteoarthritis and Cartilage</i> , 2020, 28, 7-9.	0.6	18
50	Lower gut microbiome diversity and higher abundance of proinflammatory genus <i>Collinsella</i> are associated with biopsy-proven nonalcoholic steatohepatitis. <i>Gut Microbes</i> , 2020, 11, 569-580.	4.3	125
51	Baseline self-report "central mechanisms" trait predicts persistent knee pain in the Knee Pain in the Community (KPIC) cohort. <i>Osteoarthritis and Cartilage</i> , 2020, 28, 173-181.	0.6	15
52	IL-15 and IL15RA in Osteoarthritis: Association With Symptoms and Protease Production, but Not Structural Severity. <i>Frontiers in Immunology</i> , 2020, 11, 1385.	2.2	19
53	East Midlands knee pain multiple randomised controlled trial cohort study: cohort establishment and feasibility study protocol. <i>BMJ Open</i> , 2020, 10, e037760.	0.8	5
54	Role of Drugs Used for Chronic Disease Management on Susceptibility and Severity of COVID-19: A Large Case-Control Study. <i>Clinical Pharmacology and Therapeutics</i> , 2020, 108, 1185-1194.	2.3	49

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55	Are facemasks a priority for all staff in theatre to prevent surgical site infections during shortages of supply? A systematic review and meta-analysis. <i>Journal of the Royal College of Surgeons of Edinburgh</i> , 2020, 19, e132-e139.	0.8	7
56	Real-time tracking of self-reported symptoms to predict potential COVID-19. <i>Nature Medicine</i> , 2020, 26, 1037-1040.	15.2	1,173
57	Brain perfusion patterns are altered in chronic knee pain: a spatial covariance analysis of arterial spin labelling MRI. <i>Pain</i> , 2020, 161, 1255-1263.	2.0	17
58	Human postprandial responses to food and potential for precision nutrition. <i>Nature Medicine</i> , 2020, 26, 964-973.	15.2	418
59	Serum metabolites reflecting gut microbiome alpha diversity predict type 2 diabetes. <i>Gut Microbes</i> , 2020, 11, 1632-1642.	4.3	65
60	Investigating musculoskeletal health and wellbeing; a cohort study protocol. <i>BMC Musculoskeletal Disorders</i> , 2020, 21, 182.	0.8	10
61	Consumption of Stilbenes and Flavonoids is Linked to Reduced Risk of Obesity Independently of Fiber Intake. <i>Nutrients</i> , 2020, 12, 1871.	1.7	19
62	Deficiency of Prebiotic Fiber and Insufficient Signaling Through Gut Metabolite-Sensing Receptors Leads to Cardiovascular Disease. <i>Circulation</i> , 2020, 141, 1393-1403.	1.6	176
63	Effects of Environmental Factors on Severity and Mortality of COVID-19. <i>Frontiers in Medicine</i> , 2020, 7, 607786.	1.2	40
64	Do $\beta$ -adrenoreceptor blocking drugs associate with reduced risk of symptomatic osteoarthritis and total joint replacement in the general population? A primary care-based, prospective cohort study using the Clinical Practice Research Datalink. <i>BMJ Open</i> , 2019, 9, e032050.	0.8	3
65	Gut microbiota and osteoarthritis management: An expert consensus of the European society for clinical and economic aspects of osteoporosis, osteoarthritis and musculoskeletal diseases (ESCEO). <i>Ageing Research Reviews</i> , 2019, 55, 100946.	5.0	103
66	Self-report central mechanisms trait predicts knee pain persistence in the Knee Pain In the Community (KPIC) cohort. <i>Rheumatology</i> , 2019, 58, .	0.9	0
67	Circulating levels of the anti-oxidant indolepropionic acid are associated with higher gut microbiome diversity. <i>Gut Microbes</i> , 2019, 10, 688-695.	4.3	67
68	The Metabolomic Signatures of Weight Change. <i>Metabolites</i> , 2019, 9, 67.	1.3	11
69	Microbiome genetics links short-chain fatty acids to metabolic diseases. <i>Nature Metabolism</i> , 2019, 1, 420-421.	5.1	2
70	The impact of anxiety on chronic musculoskeletal pain and the role of astrocyte activation. <i>Pain</i> , 2019, 160, 658-669.	2.0	36
71	Thresholds of ultrasound synovial abnormalities for knee osteoarthritis – a cross sectional study in the general population. <i>Osteoarthritis and Cartilage</i> , 2019, 27, 435-443.	0.6	12
72	Evaluating the efficacy of Internet-Based Exercise programme Aimed at Treating knee Osteoarthritis (iBEAT-OA) in the community: a study protocol for a randomised controlled trial. <i>BMJ Open</i> , 2019, 9, e030564.	0.8	9

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73	Genome-wide association meta-analysis of individuals of European ancestry identifies new loci explaining a substantial fraction of hair color variation and heritability. <i>Nature Genetics</i> , 2018, 50, 652-656.	9.4	86
74	Mitochondrial DNA variation and the pathogenesis of osteoarthritis phenotypes. <i>Nature Reviews Rheumatology</i> , 2018, 14, 327-340.	3.5	112
75	Traits associated with central pain augmentation in the Knee Pain In the Community (KPIC) cohort. <i>Pain</i> , 2018, 159, 1035-1044.	2.0	31
76	Genome-wide association study in 79,366 European-ancestry individuals informs the genetic architecture of 25-hydroxyvitamin D levels. <i>Nature Communications</i> , 2018, 9, 260.	5.8	295
77	Glycosylation Profile of Immunoglobulin G Is Cross-Sectionally Associated With Cardiovascular Disease Risk Score and Subclinical Atherosclerosis in Two Independent Cohorts. <i>Circulation Research</i> , 2018, 122, 1555-1564.	2.0	87
78	First validation of the gout activity score against gout impact scale in a primary care based gout cohort. <i>Joint Bone Spine</i> , 2018, 85, 323-325.	0.8	12
79	090â€fDNA methylation and its relationship with musculoskeletal health in older adults from the Hertfordshire Cohort Study: findings from an epigenome-wide association study. <i>Rheumatology</i> , 2018, 57, .	0.9	1
80	Familial aggregation and heritability of type 1 diabetes mellitus and coaggregation of chronic diseases in affected families. <i>Clinical Epidemiology</i> , 2018, Volume 10, 1447-1455.	1.5	16
81	The fecal metabolome as a functional readout of the gut microbiome. <i>Nature Genetics</i> , 2018, 50, 790-795.	9.4	482
82	Effect of dietary omega-3 fatty acid supplementation on frailty-related phenotypes in older adults: a systematic review and meta-analysis protocol. <i>BMJ Open</i> , 2018, 8, e021344.	0.8	6
83	Osteoarthritis: Genetic Studies of Monogenic and Complex Forms. , 2018, , 421-438.		0
84	Bidirectional association between disturbed sleep and neuropathic pain symptoms: a prospective cohort study in post-total joint replacement participants. <i>Journal of Pain Research</i> , 2018, Volume 11, 1087-1093.	0.8	20
85	Genetic and microbiome influence on lipid metabolism and dyslipidemia. <i>Physiological Genomics</i> , 2018, 50, 117-126.	1.0	84
86	Omega-6 oxylipins generated by soluble epoxide hydrolase are associated with knee osteoarthritis. <i>Journal of Lipid Research</i> , 2018, 59, 1763-1770.	2.0	41
87	Gut microbial diversity is associated with lower arterial stiffness in women. <i>European Heart Journal</i> , 2018, 39, 2390-2397.	1.0	181
88	Contribution of central and peripheral risk factors to prevalence, incidence and progression of knee pain: a community-based cohort study. <i>Osteoarthritis and Cartilage</i> , 2018, 26, 1461-1473.	0.6	17
89	Role of the gut microbiota in nutrition and health. <i>BMJ: British Medical Journal</i> , 2018, 361, k2179.	2.4	1,228
90	Big data boost for osteoarthritis genetics. <i>Nature Reviews Rheumatology</i> , 2018, 14, 387-388.	3.5	8

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91	Metabolomic signatures of low birthweight: Pathways to insulin resistance and oxidative stress. PLoS ONE, 2018, 13, e0194316.	1.1	21
92	Inflammatory markers and mediators in heart disease. Aging, 2018, 10, 3061-3062.	1.4	1
93	Genome-wide association scan of neuropathic pain symptoms post total joint replacement highlights a variant in the protein-kinase C gene. European Journal of Human Genetics, 2017, 25, 446-451.	1.4	39
94	Mixing omics: combining genetics and metabolomics to study rheumatic diseases. Nature Reviews Rheumatology, 2017, 13, 174-181.	3.5	63
95	Pain in knee osteoarthritis is associated with variation in the neurokinin 1/substance P receptor ( <i>TACR1</i> ) gene. European Journal of Pain, 2017, 21, 1277-1284.	1.4	21
96	Familial aggregation of rheumatoid arthritis and co-aggregation of autoimmune diseases in affected families: a nationwide population-based study. Rheumatology, 2017, 56, 928-933.	0.9	46
97	Gut microbiome diversity and high-fibre intake are related to lower long-term weight gain. International Journal of Obesity, 2017, 41, 1099-1105.	1.6	268
98	Untangling the relationship between diet and visceral fat mass through blood metabolomics and gut microbiome profiling. International Journal of Obesity, 2017, 41, 1106-1113.	1.6	68
99	Mitochondrial DNA haplogroups and ageing mechanisms in osteoarthritis. Annals of the Rheumatic Diseases, 2017, 76, 939-941.	0.5	13
100	Genome-wide association and functional studies identify a role for matrix Gla protein in osteoarthritis of the hand. Annals of the Rheumatic Diseases, 2017, 76, 2046-2053.	0.5	64
101	Association of the resolvin precursor 17-HDHA, but not D- or E- series resolvins, with heat pain sensitivity and osteoarthritis pain in humans. Scientific Reports, 2017, 7, 10748.	1.6	47
102	Omega-3 fatty acids correlate with gut microbiome diversity and production of N-carbamylglutamate in middle aged and elderly women. Scientific Reports, 2017, 7, 11079.	1.6	174
103	Metabolomic Profiling of Long-Term Weight Change: Role of Oxidative Stress and Urate Levels in Weight Gain. Obesity, 2017, 25, 1618-1624.	1.5	23
104	Genetic association studies in osteoarthritis: is it fairytale?. Current Opinion in Rheumatology, 2017, 29, 103-109.	2.0	32
105	Association of Beta-Blocker Use With Less Prevalent Joint Pain and Lower Opioid Requirement in People With Osteoarthritis. Arthritis Care and Research, 2017, 69, 1076-1081.	1.5	40
106	Familial Aggregation and Heritability of Schizophrenia and Co-aggregation of Psychiatric Illnesses in Affected Families. Schizophrenia Bulletin, 2017, 43, 1070-1078.	2.3	51
107	Triggers of acute attacks of gout, does age of gout onset matter? A primary care based cross-sectional study. PLoS ONE, 2017, 12, e0186096.	1.1	19
108	Association between ultrasound-detected synovitis and knee pain: a population-based case-control study with both cross-sectional and follow-up data. Arthritis Research and Therapy, 2017, 19, 281.	1.6	32

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109	Neuropathic pain-like symptoms and pre-surgery radiographic severity contribute to patient satisfaction 4.8 years post-total joint replacement. <i>World Journal of Orthopedics</i> , 2017, 8, 761-769.	0.8	6
110	Molecular pathways associated with blood pressure and hexadecanedioate levels. <i>PLoS ONE</i> , 2017, 12, e0175479.	1.1	8
111	The Genetics of Osteoarthritis: A Review. <i>Journal of Functional Morphology and Kinesiology</i> , 2016, 1, 140-153.	1.1	42
112	Novel Genetic Variants for Cartilage Thickness and Hip Osteoarthritis. <i>PLoS Genetics</i> , 2016, 12, e1006260.	1.5	76
113	The Pharmacogenetic Footprint of ACE Inhibition: A Population-Based Metabolomics Study. <i>PLoS ONE</i> , 2016, 11, e0153163.	1.1	13
114	Analysis and Visualization Tool for Targeted Amplicon Bisulfite Sequencing on Ion Torrent Sequencers. <i>PLoS ONE</i> , 2016, 11, e0160227.	1.1	24
115	Association of Serum Uric Acid and Disease Duration With Frequent Gout Attacks: A Caseâ€“Control Study. <i>Arthritis Care and Research</i> , 2016, 68, 1573-1577.	1.5	33
116	Replication of Associations of Genetic Loci Outside the HLA Region With Susceptibility to Antiâ€“Cyclic Citrullinated Peptideâ€“Negative Rheumatoid Arthritis. <i>Arthritis and Rheumatology</i> , 2016, 68, 1603-1613.	2.9	33
117	Metabolomic profiling to dissect the role of visceral fat in cardiometabolic health. <i>Obesity</i> , 2016, 24, 1380-1388.	1.5	41
118	Intercritical circulating levels of neo-epitopes reflecting matrixmetalloprotease-driven degradation as markers of gout and frequent gout attacks. <i>Rheumatology</i> , 2016, 55, 1642-1646.	0.9	3
119	Low omega-3 fatty acid levels associate with frequent gout attacks: a case control study. <i>Annals of the Rheumatic Diseases</i> , 2016, 75, 784-785.	0.5	18
120	KIR haplotypes are associated with late-onset type 1 diabetes in Europeanâ€“American families. <i>Genes and Immunity</i> , 2016, 17, 8-12.	2.2	17
121	A Metabolome-Wide Association Study of Kidney Function and Disease in the General Population. <i>Journal of the American Society of Nephrology: JASN</i> , 2016, 27, 1175-1188.	3.0	159
122	Investigating the Causal Relationship of C-Reactive Protein with 32 Complex Somatic and Psychiatric Outcomes: A Large-Scale Cross-Consortium Mendelian Randomization Study. <i>PLoS Medicine</i> , 2016, 13, e1001976.	3.9	150
123	Osteoarthritis Genetics. , 2016, , 1041-1047.		0
124	DNA Methylation Changes in the <i>IGF1R</i> Gene in Birth Weight Discordant Adult Monozygotic Twins. <i>Twin Research and Human Genetics</i> , 2015, 18, 635-646.	0.3	23
125	Circulating Proteomic Signatures of Chronological Age. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2015, 70, 809-816.	1.7	106
126	Circulating Levels of Antioxidant Vitamins Correlate with Better Lung Function and Reduced Exposure to Ambient Pollution. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2015, 191, 1203-1207.	2.5	39



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127	Cardiovascular disease and osteoarthritis: common pathways and patient outcomes. <i>European Journal of Clinical Investigation</i> , 2015, 45, 405-414.	1.7	90
128	Familial Aggregation of Systemic Lupus Erythematosus and Coaggregation of Autoimmune Diseases in Affected Families. <i>JAMA Internal Medicine</i> , 2015, 175, 1518.	2.6	221
129	Familial Risk of Sjögren's Syndrome and Coaggregation of Autoimmune Diseases in Affected Families: A Nationwide Population Study. <i>Arthritis and Rheumatology</i> , 2015, 67, 1904-1912.	2.9	79
130	Metabolomic study of carotid-femoral pulse-wave velocity in women. <i>Journal of Hypertension</i> , 2015, 33, 791-796.	0.3	57
131	Familial aggregation of gout and relative genetic and environmental contributions: a nationwide population study in Taiwan. <i>Annals of the Rheumatic Diseases</i> , 2015, 74, 369-374.	0.5	67
132	Metabolomic Identification of a Novel Pathway of Blood Pressure Regulation Involving Hexadecanedioate. <i>Hypertension</i> , 2015, 66, 422-429.	1.3	90
133	The UK10K project identifies rare variants in health and disease. <i>Nature</i> , 2015, 526, 82-90.	13.7	1,014
134	Genome-wide association and functional studies identify a role for <i>IGFBP3</i> in hip osteoarthritis. <i>Annals of the Rheumatic Diseases</i> , 2015, 74, 1861-1867.	0.5	47
135	Use of prescription analgesic medication and pain catastrophizing after total joint replacement surgery. <i>Seminars in Arthritis and Rheumatism</i> , 2015, 45, 150-155.	1.6	24
136	Genetics of osteoarthritis. , 2015, , 1477-1482.		0
137	Osteoarthritis Genetics. , 2015, , 1-8.		0
138	Meta-analysis identifies loci affecting levels of the potential osteoarthritis biomarkers sCOMP and uCTX-II with genome wide significance. <i>Journal of Medical Genetics</i> , 2014, 51, 596-604.	1.5	18
139	Assessment of Osteoarthritis Candidate Genes in a Meta-Analysis of Nine Genome-Wide Association Studies. <i>Arthritis and Rheumatology</i> , 2014, 66, 940-949.	2.9	108
140	The effect of <i>FTO</i> variation on increased osteoarthritis risk is mediated through body mass index: a mendelian randomisation study. <i>Annals of the Rheumatic Diseases</i> , 2014, 73, 2082-2086.	0.5	66
141	A meta-analysis of genome-wide association studies identifies novel variants associated with osteoarthritis of the hip. <i>Annals of the Rheumatic Diseases</i> , 2014, 73, 2130-2136.	0.5	108
142	Glycans Are a Novel Biomarker of Chronological and Biological Ages. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2014, 69, 779-789.	1.7	297
143	Variants Close to <i>NTRK2</i> Gene Are Associated With Birth Weight in Female Twins. <i>Twin Research and Human Genetics</i> , 2014, 17, 254-261.	0.3	16
144	An atlas of genetic influences on human blood metabolites. <i>Nature Genetics</i> , 2014, 46, 543-550.	9.4	1,084

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145	Prediction model for knee osteoarthritis incidence, including clinical, genetic and biochemical risk factors. <i>Annals of the Rheumatic Diseases</i> , 2014, 73, 2116-2121.	0.5	111
146	Association of adiponectin and leptin with relative telomere length in seven independent cohorts including 11,448 participants. <i>European Journal of Epidemiology</i> , 2014, 29, 629-638.	2.5	23
147	Contribution of the COMT Val158Met variant to symptomatic knee osteoarthritis. <i>Annals of the Rheumatic Diseases</i> , 2014, 73, 315-317.	0.5	18
148	Use of statins is associated with a lower prevalence of generalised osteoarthritis. <i>Annals of the Rheumatic Diseases</i> , 2014, 73, 943-945.	0.5	30
149	Severe osteoarthritis of the hand associates with common variants within the ALDH1A2 gene and with rare variants at 1p31. <i>Nature Genetics</i> , 2014, 46, 498-502.	9.4	136
150	Design and Analysis of Metabolomics Studies in Epidemiologic Research: A Primer on -Omics Technologies. <i>American Journal of Epidemiology</i> , 2014, 180, 129-139.	1.6	152
151	History of knee surgery is associated with higher prevalence of neuropathic pain-like symptoms in patients with severe osteoarthritis of the knee. <i>Seminars in Arthritis and Rheumatism</i> , 2014, 43, 588-592.	1.6	81
152	Large scale meta-analysis of urinary C-terminal telopeptide, serum cartilage oligomeric protein and matrix metalloprotease degraded type II collagen and their role in prevalence, incidence and progression of osteoarthritis. <i>Osteoarthritis and Cartilage</i> , 2014, 22, 683-689.	0.6	72
153	Association of interleukin-6 gene polymorphisms with hand osteoarthritis and hand osteoporosis. <i>Cytokine</i> , 2014, 69, 94-101.	1.4	20
154	SMAD3 Is Associated with the Total Burden of Radiographic Osteoarthritis: The Chingford Study. <i>PLoS ONE</i> , 2014, 9, e97786.	1.1	17
155	Omics technologies and the study of human ageing. <i>Nature Reviews Genetics</i> , 2013, 14, 601-607.	7.7	108
156	Next Generation Sequencing Reveals the Association of DRB3*02:02 With Type 1 Diabetes. <i>Diabetes</i> , 2013, 62, 2618-2622.	0.3	42
157	Targeted metabolomics profiles are strongly correlated with nutritional patterns in women. <i>Metabolomics</i> , 2013, 9, 506-514.	1.4	110
158	Inference of the Genetic Architecture Underlying BMI and Height with the Use of 20,240 Sibling Pairs. <i>American Journal of Human Genetics</i> , 2013, 93, 865-875.	2.6	104
159	Receiver Operating Characteristic Analysis of HLA, CTLA4, and Insulin Genotypes for Type 1 Diabetes. <i>Diabetes Care</i> , 2013, 36, 2504-2507.	4.3	5
160	Metabolomic markers reveal novel pathways of ageing and early development in human populations. <i>International Journal of Epidemiology</i> , 2013, 42, 1111-1119.	0.9	241
161	Identification of seven loci affecting mean telomere length and their association with disease. <i>Nature Genetics</i> , 2013, 45, 422-427.	9.4	808
162	Prediction of Type 1 Diabetes. <i>Diabetes</i> , 2013, 62, 1020-1021.	0.3	9

#	ARTICLE	IF	CITATIONS
163	The genetic contribution to severe post-traumatic osteoarthritis. <i>Annals of the Rheumatic Diseases</i> , 2013, 72, 1687-1690.	0.5	22
164	The <i>DOT1L</i> rs12982744 polymorphism is associated with osteoarthritis of the hip with genome-wide statistical significance in males. <i>Annals of the Rheumatic Diseases</i> , 2013, 72, 1264-1265.	0.5	51
165	HLA Class II Genotyping of African American Type 1 Diabetic Patients Reveals Associations Unique to African Haplotypes. <i>Diabetes</i> , 2013, 62, 3292-3299.	0.3	52
166	Genome-wide association study meta-analysis of chronic widespread pain: evidence for involvement of the 5p15.2 region. <i>Annals of the Rheumatic Diseases</i> , 2013, 72, 427-436.	0.5	112
167	Evaluation of the genetic overlap between osteoarthritis with body mass index and height using genome-wide association scan data. <i>Annals of the Rheumatic Diseases</i> , 2013, 72, 935-941.	0.5	52
168	Cohort Profile: TwinsUK and Healthy Ageing Twin Study. <i>International Journal of Epidemiology</i> , 2013, 42, 76-85.	0.9	224
169	Osteoarthritis – Genetic Studies of Monogenic and Complex Forms. , 2013, , 275-293.		3
170	No evidence of an association between mitochondrial DNA variants and osteoarthritis in 7393 cases and 5122 controls. <i>Annals of the Rheumatic Diseases</i> , 2013, 72, 136-139.	0.5	39
171	The Contribution of Diet and Genotype to Iron Status in Women: A Classical Twin Study. <i>PLoS ONE</i> , 2013, 8, e83047.	1.1	7
172	Glycosylation of Immunoglobulin G: Role of Genetic and Epigenetic Influences. <i>PLoS ONE</i> , 2013, 8, e82558.	1.1	105
173	Genetic contribution to radiographic severity in osteoarthritis of the knee. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 1537-1540.	0.5	17
174	Variants in the genes encoding TNF- $\alpha$ , IL-10, and GSTP1 influence the effect of $\alpha$ -tocopherol on inflammatory cell responses in healthy men. <i>American Journal of Clinical Nutrition</i> , 2012, 95, 1461-1467.	2.2	10
175	Genome-wide association and functional studies identify the <i>DOT1L</i> gene to be involved in cartilage thickness and hip osteoarthritis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 8218-8223.	3.3	154
176	Genome-wide meta-analysis points to CTC1 and ZNF676 as genes regulating telomere homeostasis in humans. <i>Human Molecular Genetics</i> , 2012, 21, 5385-5394.	1.4	210
177	A role for PACE4 in osteoarthritis pain: evidence from human genetic association and null mutant phenotype. <i>Annals of the Rheumatic Diseases</i> , 2012, 71, 1042-1048.	0.5	49
178	Use of class I and class II HLA loci for predicting age at onset of type 1 diabetes in multiple populations. <i>Diabetologia</i> , 2012, 55, 2394-2401.	2.9	36
179	Identification of new susceptibility loci for osteoarthritis (arcOGEN): a genome-wide association study. <i>Lancet</i> , The, 2012, 380, 815-823.	6.3	373
180	Genetic factors in OA pathogenesis. <i>Bone</i> , 2012, 51, 258-264.	1.4	71

#	ARTICLE	IF	CITATIONS
181	Epigenome-Wide Scans Identify Differentially Methylated Regions for Age and Age-Related Phenotypes in a Healthy Ageing Population. <i>PLoS Genetics</i> , 2012, 8, e1002629.	1.5	620
182	Inverse Relationship Between Preoperative Radiographic Severity and Postoperative Pain in Patients with Osteoarthritis who Have Undergone Total Joint Arthroplasty. <i>Seminars in Arthritis and Rheumatism</i> , 2012, 41, 568-575.	1.6	87
183	Reply to "Human genetic studies on osteoarthritis from clinicians' viewpoints". <i>Osteoarthritis and Cartilage</i> , 2012, 20, 250-251.	0.6	1
184	Role of the Na <sup>v</sup> 1.7 R1150W amino acid change in susceptibility to symptomatic knee osteoarthritis and multiple regional pain. <i>Arthritis Care and Research</i> , 2011, 63, 440-444.	1.5	29
185	Genetic epidemiology of hip and knee osteoarthritis. <i>Nature Reviews Rheumatology</i> , 2011, 7, 23-32.	3.5	203
186	Insights into the genetic architecture of osteoarthritis from stage 1 of the arcOGEN study. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 864-867.	0.5	119
187	Meta-analysis of genome-wide association studies confirms a susceptibility locus for knee osteoarthritis on chromosome 7q22. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 349-355.	0.5	126
188	Race-specific type 1 diabetes risk of HLA-DR7 haplotypes. <i>Tissue Antigens</i> , 2011, 78, 348-351.	1.0	18
189	The effect of genome-wide association scan quality control on imputation outcome for common variants. <i>European Journal of Human Genetics</i> , 2011, 19, 610-614.	1.4	27
190	Involvement of the CLEC3B gene in osteoarthritis. <i>Osteoarthritis and Cartilage</i> , 2011, 19, 249.	0.6	2
191	Recommendations for standardization and phenotype definitions in genetic studies of osteoarthritis: the TREAT-OA consortium. <i>Osteoarthritis and Cartilage</i> , 2011, 19, 254-264.	0.6	82
192	Large-scale meta-analysis of interleukin-1 beta and interleukin-1 receptor antagonist polymorphisms on risk of radiographic hip and knee osteoarthritis and severity of knee osteoarthritis. <i>Osteoarthritis and Cartilage</i> , 2011, 19, 265-271.	0.6	72
193	A Variant in MCF2L Is Associated with Osteoarthritis. <i>American Journal of Human Genetics</i> , 2011, 89, 446-450.	2.6	115
194	Genetics of the HLA Region in the Prediction of Type 1 Diabetes. <i>Current Diabetes Reports</i> , 2011, 11, 533-542.	1.7	302
195	GDF5 single-nucleotide polymorphism rs143383 is associated with lumbar disc degeneration in Northern European women. <i>Arthritis and Rheumatism</i> , 2011, 63, 708-712.	6.7	100
196	The Ile585Val TRPV1 variant is involved in risk of painful knee osteoarthritis. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 1556-1561.	0.5	111
197	The GDF5 rs143383 polymorphism is associated with osteoarthritis of the knee with genome-wide statistical significance. <i>Annals of the Rheumatic Diseases</i> , 2011, 70, 873-875.	0.5	137
198	A Twin Study of Mitochondrial DNA Polymorphisms Shows that Heteroplasmy at Multiple Sites Is Associated with mtDNA Variant 16093 but Not with Zygosity. <i>PLoS ONE</i> , 2011, 6, e22332.	1.1	21

#	ARTICLE	IF	CITATIONS
199	Large Scale Replication Study of the Association between HLA Class II/BTNL2 Variants and Osteoarthritis of the Knee in European-Descent Populations. <i>PLoS ONE</i> , 2011, 6, e23371.	1.1	32
200	The genetic epidemiology of osteoarthritis. <i>Current Opinion in Rheumatology</i> , 2010, 22, 139-143.	2.0	53
201	Genetic Markers of Osteoarthritis. <i>Current Rheumatology Reviews</i> , 2010, 6, 257-267.	0.4	4
202	Association of JAG1 with Bone Mineral Density and Osteoporotic Fractures: A Genome-wide Association Study and Follow-up Replication Studies. <i>American Journal of Human Genetics</i> , 2010, 86, 229-239.	2.6	188
203	A meta-analysis of interleukin-6 promoter polymorphisms on risk of hip and knee osteoarthritis. <i>Osteoarthritis and Cartilage</i> , 2010, 18, 699-704.	0.6	36
204	The clinical relevance of genetic susceptibility to osteoarthritis. <i>Best Practice and Research in Clinical Rheumatology</i> , 2010, 24, 3-14.	1.4	52
205	A genome-wide association study identifies an osteoarthritis susceptibility locus on chromosome 7q22. <i>Arthritis and Rheumatism</i> , 2010, 62, 499-510.	6.7	178
206	Genetic variation in the <i>SMAD3</i> gene is associated with hip and knee osteoarthritis. <i>Arthritis and Rheumatism</i> , 2010, 62, 2347-2352.	6.7	145
207	Involvement of different risk factors in clinically severe large joint osteoarthritis according to the presence of hand interphalangeal nodes. <i>Arthritis and Rheumatism</i> , 2010, 62, 2688-2695.	6.7	46
208	Common genetic variation in the Estrogen Receptor Beta (ESR2) gene and osteoarthritis: results of a meta-analysis. <i>BMC Medical Genetics</i> , 2010, 11, 164.	2.1	8
209	Genetic variation within the HLA class III influences T1D susceptibility conferred by high-risk HLA haplotypes. <i>Genes and Immunity</i> , 2010, 11, 209-218.	2.2	9
210	Sequence variants at <i>CHRNA6</i> and <i>CYP2A6</i> affect smoking behavior. <i>Nature Genetics</i> , 2010, 42, 448-453.	9.4	649
211	A Common Variant in the Telomerase RNA Component Is Associated with Short Telomere Length. <i>PLoS ONE</i> , 2010, 5, e13048.	1.1	39
212	HLA Class I and Genetic Susceptibility to Type 1 Diabetes. <i>Diabetes</i> , 2010, 59, 2972-2979.	0.3	202
213	HLA DPA1, DPB1 Alleles and Haplotypes Contribute to the Risk Associated With Type 1 Diabetes. <i>Diabetes</i> , 2010, 59, 2055-2062.	0.3	64
214	Leukocyte telomere length is associated with cognitive performance in healthy women. <i>Neurobiology of Aging</i> , 2010, 31, 986-992.	1.5	104
215	Common variants near <i>TERC</i> are associated with mean telomere length. <i>Nature Genetics</i> , 2010, 42, 197-199.	9.4	296
216	Human aging-associated DNA hypermethylation occurs preferentially at bivalent chromatin domains. <i>Genome Research</i> , 2010, 20, 434-439.	2.4	646

#	ARTICLE	IF	CITATIONS
217	Molecular pathogenesis and genetics of osteoarthritis: implications for personalized medicine. <i>Personalized Medicine</i> , 2010, 7, 49-63.	0.8	5
218	Homozygosity for the Ala allele of the PPAR $\gamma$ 3 Pro12Ala polymorphism is associated with reduced risk of coronary artery disease. <i>Disease Markers</i> , 2010, 29, 259-64.	0.6	10
219	A genome-wide association study suggests that a locus within the ataxin 2 binding protein 1 gene is associated with hand osteoarthritis: the Treat-OA consortium. <i>Journal of Medical Genetics</i> , 2009, 46, 614-616.	1.5	58
220	Association of the DVWA and GDF5 polymorphisms with osteoarthritis in UK populations. <i>Annals of the Rheumatic Diseases</i> , 2009, 68, 1916-1920.	0.5	70
221	The genetic predisposition to osteoarthritis. <i>IBMS BoneKEy</i> , 2009, 6, 181-189.	0.1	3
222	A genome-wide association study identifies a novel locus on chromosome 18q12.2 influencing white cell telomere length. <i>Journal of Medical Genetics</i> , 2009, 46, 451-454.	1.5	76
223	Association of a nsSNP in ADAMTS14 to some osteoarthritis phenotypes. <i>Osteoarthritis and Cartilage</i> , 2009, 17, 321-327.	0.6	62
224	Large-scale analysis of association between <i>GDF5</i> and <i>FRZB</i> variants and osteoarthritis of the hip, knee, and hand. <i>Arthritis and Rheumatism</i> , 2009, 60, 1710-1721.	6.7	181
225	Variation at the ANP32A gene is associated with risk of hip osteoarthritis in women. <i>Arthritis and Rheumatism</i> , 2009, 60, 2046-2054.	6.7	27
226	The Contribution of Genes to Osteoarthritis. <i>Medical Clinics of North America</i> , 2009, 93, 45-66.	1.1	99
227	Natural history and risk factors for bone loss in postmenopausal Caucasian women: a 15-year follow-up population-based study. <i>Osteoporosis International</i> , 2008, 19, 1211-1217.	1.3	39
228	Genome-wide Association Scan Identifies a Prostaglandin-Endoperoxide Synthase 2 Variant Involved in Risk of Knee Osteoarthritis. <i>American Journal of Human Genetics</i> , 2008, 82, 1231-1240.	2.6	109
229	Radiographic osteoarthritis at three joint sites and FRZB, LRP5, and LRP6 polymorphisms in two population-based cohorts. <i>Osteoarthritis and Cartilage</i> , 2008, 16, 1141-1149.	0.6	47
230	Type 1 diabetes risk for human leukocyte antigen (HLA)-DR3 haplotypes depends on genotypic context: Association of DPB1 and HLA class I loci among DR3- and DR4-matched Italian patients and controls. <i>Human Immunology</i> , 2008, 69, 291-300.	1.2	21
231	The additive effect of individual genes in predicting risk of knee osteoarthritis. <i>Annals of the Rheumatic Diseases</i> , 2008, 67, 124-127.	0.5	53
232	Homocysteine levels and leukocyte telomere length. <i>Atherosclerosis</i> , 2008, 200, 271-277.	0.4	78
233	Bone mineral density, osteoporosis, and osteoporotic fractures: a genome-wide association study. <i>Lancet</i> , The, 2008, 371, 1505-1512.	6.3	612
234	The Contribution of Genes to Osteoarthritis. <i>Rheumatic Disease Clinics of North America</i> , 2008, 34, 581-603.	0.8	75

#	ARTICLE	IF	CITATIONS
235	Influence of apoA-V gene variants on postprandial triglyceride metabolism: impact of gender. Journal of Lipid Research, 2008, 49, 945-953.	2.0	31
236	HLA DR-DQ Haplotypes and Genotypes and Type 1 Diabetes Risk. Diabetes, 2008, 57, 1084-1092.	0.3	631
237	Searching for Additional Disease Loci in a Genomic Region. Advances in Genetics, 2008, 60, 253-292.	0.8	15
238	Serum Adiponectin and Bone Mineral Density in Women. Journal of Clinical Endocrinology and Metabolism, 2007, 92, 1517-1523.	1.8	191
239	Nevus Size and Number Are Associated with Telomere Length and Represent Potential Markers of a Decreased Senescence <i>in vivo</i> . Cancer Epidemiology Biomarkers and Prevention, 2007, 16, 1499-1502.	1.1	115
240	The Interaction of Genes and Smoking on Forced Expiratory Volume. Chest, 2007, 132, 1772-1777.	0.4	22
241	Joint effect of dopaminergic genes on likelihood of smoking following treatment with bupropion SR.. Health Psychology, 2007, 26, 361-368.	1.3	44
242	Higher serum vitamin D concentrations are associated with longer leukocyte telomere length in women. American Journal of Clinical Nutrition, 2007, 86, 1420-1425.	2.2	208
243	Gene-gene interactions between CYP2B6 and CYP2A6 in nicotine metabolism. Pharmacogenetics and Genomics, 2007, 17, 1007-1015.	0.7	36
244	Improved weight management using genetic information to personalize a calorie controlled diet. Nutrition Journal, 2007, 6, 29.	1.5	122
245	Sex and ethnic differences in the association of ASPN, CALM1, COL2A1, COMP, and FRZB with genetic susceptibility to osteoarthritis of the knee. Arthritis and Rheumatism, 2007, 56, 137-146.	6.7	178
246	Relative predispositional effects of HLA class II DRB1-DQB1 haplotypes and genotypes on type 1 diabetes: a meta-analysis. Tissue Antigens, 2007, 70, 110-127.	1.0	153
247	Telomere length in leukocytes correlates with bone mineral density and is shorter in women with osteoporosis. Osteoporosis International, 2007, 18, 1203-1210.	1.3	143
248	Mapping Genetic Loci That Determine Leukocyte Telomere Length in a Large Sample of Unselected Female Sibling Pairs. American Journal of Human Genetics, 2006, 78, 480-486.	2.6	242
249	Linkage Disequilibrium With Predisposing DR3 Haplotypes Accounts for Apparent Effects of Tumor Necrosis Factor and Lymphotoxin- $\alpha$ Polymorphisms on Type 1 Diabetes Susceptibility. Human Immunology, 2006, 67, 999-1004.	1.2	33
250	The effects of social status on biological aging as measured by white-blood-cell telomere length. Aging Cell, 2006, 5, 361-365.	3.0	288
251	Reproducible genetic associations between candidate genes and clinical knee osteoarthritis in men and women. Arthritis and Rheumatism, 2006, 54, 533-539.	6.7	157
252	Association between a variation in LRCH1 and knee osteoarthritis: A genome-wide single-nucleotide polymorphism association study using DNA pooling. Arthritis and Rheumatism, 2006, 54, 524-532.	6.7	60

#	ARTICLE	IF	CITATIONS
253	When is a replication not a replication? Or how to spot a good genetic association study. <i>Arthritis and Rheumatism</i> , 2006, 54, 1051-1054.	6.7	13
254	Menopause Modifies the Association of Leukocyte Telomere Length with Insulin Resistance and Inflammation. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2006, 91, 635-640.	1.8	158
255	Reduction of leucocyte telomere length in radiographic hand osteoarthritis: a population-based study. <i>Annals of the Rheumatic Diseases</i> , 2006, 65, 1444-1448.	0.5	62
256	Human telomere biology: pitfalls of moving from the laboratory to epidemiology. <i>International Journal of Epidemiology</i> , 2006, 35, 1424-1429.	0.9	161
257	Radiographic Progression of Lumbar Spine Disc Degeneration Is Influenced by Variation at Inflammatory Genes. <i>Spine</i> , 2005, 30, 2445-2451.	1.0	80
258	Extended DR3-D6S273-HLA-B haplotypes are associated with increased susceptibility to type 1 diabetes in US Caucasians. <i>Tissue Antigens</i> , 2005, 65, 115-119.	1.0	27
259	Dopamine receptor DRD2 genotype and smoking cessation outcome following treatment with bupropion SR. <i>Pharmacogenomics Journal</i> , 2005, 5, 21-29.	0.9	114
260	D6S265*15 marks a DRB1*15, DQB1*0602 haplotype associated with attenuated protection from type 1 diabetes mellitus. <i>Diabetologia</i> , 2005, 48, 2540-2543.	2.9	21
261	Association of Non-HLA Genes With Type 1 Diabetes Autoimmunity. <i>Diabetes</i> , 2005, 54, 2482-2486.	0.3	55
262	A $\beta$ -lactamase with reduced immunogenicity for the targeted delivery of chemotherapeutics using antibody-directed enzyme prodrug therapy. <i>Molecular Cancer Therapeutics</i> , 2005, 4, 1791-1800.	1.9	73
263	Obesity, cigarette smoking, and telomere length in women. <i>Lancet</i> , The, 2005, 366, 662-664.	6.3	1,215
264	Association of the single nucleotide polymorphism C1858T of the PTPN22 gene with type 1 diabetes. <i>Human Immunology</i> , 2005, 66, 60-64.	1.2	107
265	Human leukocyte antigen class I B and C loci contribute to Type 1 Diabetes (T1D) susceptibility and age at T1D onset. <i>Human Immunology</i> , 2005, 66, 301-313.	1.2	99
266	DPB1 Alleles Are Associated With Type 1 Diabetes Susceptibility in Multiple Ethnic Groups. <i>Diabetes</i> , 2004, 53, 2158-2163.	0.3	39
267	The HLA-DR2 haplotype is associated with an increased proliferative response to the immunodominant CD4+ T-cell epitope in human interferon- $\beta$ . <i>Genes and Immunity</i> , 2004, 5, 1-7.	2.2	35
268	Association study of candidate genes for the prevalence and progression of knee osteoarthritis. <i>Arthritis and Rheumatism</i> , 2004, 50, 2497-2507.	6.7	163
269	Human population-based identification of CD4+ T-cell peptide epitope determinants. <i>Journal of Immunological Methods</i> , 2003, 281, 95-108.	0.6	16
270	Association and Interaction of the IL4R, IL4, and IL13 Loci with Type 1 Diabetes among Filipinos. <i>American Journal of Human Genetics</i> , 2003, 72, 1505-1514.	2.6	70



#	ARTICLE	IF	CITATIONS
271	A Polymorphism in the TCF7 Gene, C883A, Is Associated With Type 1 Diabetes. <i>Diabetes</i> , 2003, 52, 1579-1582.	0.3	39
272	Val64Ile Polymorphism in the C-C Chemokine Receptor 2 Is Associated With Reduced Coronary Artery Calcification. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2002, 22, 1924-1928.	1.1	70
273	The HLA class I A locus affects susceptibility to type 1 diabetes. <i>Human Immunology</i> , 2002, 63, 657-664.	1.2	106
274	Miniaturized sealed-tube allele-specific PCR. <i>Human Mutation</i> , 2002, 19, 543-553.	1.1	44
275	Association of IL4R Haplotypes With Type 1 Diabetes. <i>Diabetes</i> , 2002, 51, 3336-3341.	0.3	63
276	Association of Traditional Risk Factors with Coronary Calcification in Persons with a Family History of Premature Coronary Heart Disease: The Study of the Inherited Risk of Coronary Atherosclerosis. <i>Journal of Investigative Medicine</i> , 2001, 49, 353-361.	0.7	42
277	Modeling of HLA class II susceptibility to Type I diabetes reveals an effect associated with DPB1. <i>Genetic Epidemiology</i> , 2001, 21, 212-223.	0.6	30
278	Making drug discovery a SN(i)P. <i>Drug Discovery Today</i> , 2000, 5, 388-396.	3.2	11
279	The HLA class II locus DPB1 can influence susceptibility to type 1 diabetes. <i>Diabetes</i> , 2000, 49, 121-125.	0.3	97
280	Association between type 1 diabetes age of onset and HLA among sibling pairs. <i>Diabetes</i> , 1999, 48, 1658-1661.	0.3	54
281	Codon usage bias and base composition in MHC genes in humans and common chimpanzees. <i>Immunogenetics</i> , 1999, 49, 272-279.	1.2	13
282	Evidence for linkage and association to alcohol dependence on chromosome 19. <i>Genetic Epidemiology</i> , 1999, 17, S367-S372.	0.6	12
283	Mutational processes of simple-sequence repeat loci in human populations.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1994, 91, 3166-3170.	3.3	1,381
284	Molecular evolution of class A $\beta$ -lactamases: phylogeny and patterns of sequence conservation. <i>Molecular Microbiology</i> , 1990, 4, 1957-1965.	1.2	15
285	Primary structure of <i>Trypanosoma cruzi</i> small-subunit ribosomal RNA coding region: comparison with other trypanosomatids. <i>Molecular and Biochemical Parasitology</i> , 1990, 41, 207-212.	0.5	40
286	Real-time tracking of self-reported symptoms to predict potential COVID-19. , 0, .		1
287	Longitudinal Assessment of Symptoms and Risk of SARS-CoV-2 Infection in Healthcare Workers Across 5 Hospitals to Understand Ethnic Differences in Infection Risk. <i>SSRN Electronic Journal</i> , 0, , .	0.4	0