

Krina T Zondervan

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

161
papers

17,568
citations

61
h-index

132
g-index

178
ext. papers

21,430
ext. citations

9
avg, IF

6.23
L-index

#	Paper	IF	Citations
161	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015 , 518, 197-206	50.4	2687
160	New genetic loci link adipose and insulin biology to body fat distribution. <i>Nature</i> , 2015 , 518, 187-196	50.4	920
159	Data quality control in genetic case-control association studies. <i>Nature Protocols</i> , 2010 , 5, 1564-73	18.8	741
158	Meta-analysis identifies 13 new loci associated with waist-hip ratio and reveals sexual dimorphism in the genetic basis of fat distribution. <i>Nature Genetics</i> , 2010 , 42, 949-60	36.3	724
157	Impact of endometriosis on quality of life and work productivity: a multicenter study across ten countries. <i>Fertility and Sterility</i> , 2011 , 96, 366-373.e8	4.8	699
156	Mapping cis- and trans-regulatory effects across multiple tissues in twins. <i>Nature Genetics</i> , 2012 , 44, 1084-93	36.3	572
155	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	6	501
154	The complex interplay among factors that influence allelic association. <i>Nature Reviews Genetics</i> , 2004 , 5, 89-100	30.1	429
153	Endometriosis. <i>Nature Reviews Disease Primers</i> , 2018 , 4, 9	51.1	397
152	Basic statistical analysis in genetic case-control studies. <i>Nature Protocols</i> , 2011 , 6, 121-33	18.8	344
151	The architecture of gene regulatory variation across multiple human tissues: the MuTHER study. <i>PLoS Genetics</i> , 2011 , 7, e1002003	6	336
150	Endometriosis. <i>New England Journal of Medicine</i> , 2020 , 382, 1244-1256	59.2	300
149	Genome-wide association identifies nine common variants associated with fasting proinsulin levels and provides new insights into the pathophysiology of type 2 diabetes. <i>Diabetes</i> , 2011 , 60, 2624-34	0.9	285
148	Global analysis of DNA methylation variation in adipose tissue from twins reveals links to disease-associated variants in distal regulatory elements. <i>American Journal of Human Genetics</i> , 2013 , 93, 876-90	11	269
147	Genome-wide associations for birth weight and correlations with adult disease. <i>Nature</i> , 2016 , 538, 248-252	52.4	266
146	Identification of an imprinted master trans regulator at the KLF14 locus related to multiple metabolic phenotypes. <i>Nature Genetics</i> , 2011 , 43, 561-4	36.3	253
145	Association Between Telomere Length and Risk of Cancer and Non-Neoplastic Diseases: A Mendelian Randomization Study. <i>JAMA Oncology</i> , 2017 , 3, 636-651	13.4	236

144	Genome-wide association study identifies a locus at 7p15.2 associated with endometriosis. <i>Nature Genetics</i> , 2011 , 43, 51-4	36.3	227
143	Priorities for endometriosis research: recommendations from an international consensus workshop. <i>Reproductive Sciences</i> , 2009 , 16, 335-46	3	227
142	What makes a good case-control study? Design issues for complex traits such as endometriosis. <i>Human Reproduction</i> , 2002 , 17, 1415-23	5.7	223
141	Prevalence and incidence of adults consulting for shoulder conditions in UK primary care; patterns of diagnosis and referral. <i>Rheumatology</i> , 2006 , 45, 215-21	3.9	222
140	Epidemiology of hip and knee pain and its impact on overall health status in older adults. <i>British Journal of Rheumatology</i> , 2004 , 43, 497-504		218
139	Prevalence and incidence of chronic pelvic pain in primary care: evidence from a national general practice database. <i>BJOG: an International Journal of Obstetrics and Gynaecology</i> , 1999 , 106, 1149-55	3.7	217
138	Genome-wide association meta-analysis identifies new endometriosis risk loci. <i>Nature Genetics</i> , 2012 , 44, 1355-9	36.3	214
137	Genome-wide analysis identifies 12 loci influencing human reproductive behavior. <i>Nature Genetics</i> , 2016 , 48, 1462-1472	36.3	198
136	Chronic pelvic pain in the community--symptoms, investigations, and diagnoses. <i>American Journal of Obstetrics and Gynecology</i> , 2001 , 184, 1149-55	6.4	190
135	Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. <i>Nature Genetics</i> , 2018 , 50, 26-41	36.3	186
134	The community prevalence of chronic pelvic pain in women and associated illness behaviour. <i>British Journal of General Practice</i> , 2001 , 51, 541-7	1.6	170
133	Linkage disequilibrium mapping via cladistic analysis of single-nucleotide polymorphism haplotypes. <i>American Journal of Human Genetics</i> , 2004 , 75, 35-43	11	168
132	The search for genes contributing to endometriosis risk. <i>Human Reproduction Update</i> , 2008 , 14, 447-57	15.8	156
131	Meta-analysis identifies five novel loci associated with endometriosis highlighting key genes involved in hormone metabolism. <i>Nature Communications</i> , 2017 , 8, 15539	17.4	151
130	Central changes associated with chronic pelvic pain and endometriosis. <i>Human Reproduction Update</i> , 2014 , 20, 737-47	15.8	147
129	Estimation and partitioning of polygenic variation captured by common SNPs for AlzheimerB disease, multiple sclerosis and endometriosis. <i>Human Molecular Genetics</i> , 2013 , 22, 832-41	5.6	147
128	Differences in characteristics among 1,000 women with endometriosis based on extent of disease. <i>Fertility and Sterility</i> , 2008 , 89, 538-45	4.8	140
127	Genetic variants underlying risk of endometriosis: insights from meta-analysis of eight genome-wide association and replication datasets. <i>Human Reproduction Update</i> , 2014 , 20, 702-16	15.8	131

126	Designing candidate gene and genome-wide case-control association studies. <i>Nature Protocols</i> , 2007 , 2, 2492-501	18.8	124
125	World Endometriosis Research Foundation Endometriosis Phenome and Biobanking Harmonization Project: II. Clinical and covariate phenotype data collection in endometriosis research. <i>Fertility and Sterility</i> , 2014 , 102, 1223-32	4.8	119
124	Human metabolic profiles are stably controlled by genetic and environmental variation. <i>Molecular Systems Biology</i> , 2011 , 7, 525	12.2	119
123	World Endometriosis Research Foundation Endometriosis Phenome and Biobanking Harmonisation Project: I. Surgical phenotype data collection in endometriosis research. <i>Fertility and Sterility</i> , 2014 , 102, 1213-22	4.8	116
122	A genome-wide metabolic QTL analysis in Europeans implicates two loci shaped by recent positive selection. <i>PLoS Genetics</i> , 2011 , 7, e1002270	6	109
121	Distinct developmental profile of lower-body adipose tissue defines resistance against obesity-associated metabolic complications. <i>Diabetes</i> , 2014 , 63, 3785-97	0.9	107
120	Peripheral changes in endometriosis-associated pain. <i>Human Reproduction Update</i> , 2014 , 20, 717-36	15.8	106
119	World Endometriosis Research Foundation Endometriosis Phenome and Biobanking Harmonisation Project: IV. Tissue collection, processing, and storage in endometriosis research. <i>Fertility and Sterility</i> , 2014 , 102, 1244-53	4.8	100
118	Chronic pelvic pain in New Zealand: prevalence, pain severity, diagnoses and use of the health services. <i>Australian and New Zealand Journal of Public Health</i> , 2004 , 28, 369-75	2.3	99
117	Is early age at menarche a risk factor for endometriosis? A systematic review and meta-analysis of case-control studies. <i>Fertility and Sterility</i> , 2012 , 98, 702-712.e6	4.8	92
116	World Endometriosis Research Foundation Endometriosis Phenome and Biobanking Harmonization Project: III. Fluid biospecimen collection, processing, and storage in endometriosis research. <i>Fertility and Sterility</i> , 2014 , 102, 1233-43	4.8	91
115	The genetic basis of endometriosis. <i>Current Opinion in Obstetrics and Gynecology</i> , 2001 , 13, 309-14	2.4	91
114	High-quality genomic DNA extraction from formalin-fixed and paraffin-embedded samples deparaffinized using mineral oil. <i>Analytical Biochemistry</i> , 2009 , 395, 265-7	3.1	83
113	Genome-Wide Association Study of Susceptibility to Idiopathic Pulmonary Fibrosis. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2020 , 201, 564-574	10.2	81
112	The prevalence of chronic pelvic pain in women in the United Kingdom: a systematic review. <i>BJOG: an International Journal of Obstetrics and Gynaecology</i> , 1998 , 105, 93-9	3.7	80
111	Do dietary and supplementary intakes of antioxidants differ with smoking status?. <i>International Journal of Epidemiology</i> , 1996 , 25, 70-9	7.8	80
110	The association between endometriosis and autoimmune diseases: a systematic review and meta-analysis. <i>Human Reproduction Update</i> , 2019 , 25, 486-503	15.8	76
109	The vascular endothelial growth factor (VEGF) +405G>C 5Puntranslated region polymorphism and increased risk of endometriosis in South Indian women: a case control study. <i>Human Reproduction</i> , 2005 , 20, 1844-9	5.7	76

108	Patterns of diagnosis and referral in women consulting for chronic pelvic pain in UK primary care. <i>BJOG: an International Journal of Obstetrics and Gynaecology</i> , 1999 , 106, 1156-61	3.7	76
107	Developing symptom-based predictive models of endometriosis as a clinical screening tool: results from a multicenter study. <i>Fertility and Sterility</i> , 2012 , 98, 692-701.e5	4.8	75
106	Familial aggregation of endometriosis in a large pedigree of rhesus macaques. <i>Human Reproduction</i> , 2004 , 19, 448-55	5.7	75
105	Research Priorities for Endometriosis. <i>Reproductive Sciences</i> , 2017 , 24, 202-226	3	72
104	Recent insights on the genetics and epigenetics of endometriosis. <i>Clinical Genetics</i> , 2017 , 91, 254-264	4	71
103	Variability of gene expression profiles in human blood and lymphoblastoid cell lines. <i>BMC Genomics</i> , 2010 , 11, 96	4.5	64
102	Lymphotoxin-alpha gene and risk of myocardial infarction in 6,928 cases and 2,712 controls in the ISIS case-control study. <i>PLoS Genetics</i> , 2006 , 2, e107	6	64
101	Genome-wide linkage and association analyses implicate FASN in predisposition to Uterine Leiomyomata. <i>American Journal of Human Genetics</i> , 2012 , 91, 621-8	11	62
100	The management of menopause in women with a history of endometriosis: a systematic review. <i>Human Reproduction Update</i> , 2017 , 23, 481-500	15.8	60
99	Epidemiology of chronic pelvic pain. <i>Best Practice and Research in Clinical Obstetrics and Gynaecology</i> , 2000 , 14, 403-14	4.6	60
98	Genome-wide genetic analyses highlight mitogen-activated protein kinase (MAPK) signaling in the pathogenesis of endometriosis. <i>Human Reproduction</i> , 2017 , 32, 780-793	5.7	59
97	Genome-wide enrichment analysis between endometriosis and obesity-related traits reveals novel susceptibility loci. <i>Human Molecular Genetics</i> , 2015 , 24, 1185-99	5.6	57
96	Impact of persistent hip or knee pain on overall health status in elderly people: a longitudinal population study. <i>Arthritis and Rheumatism</i> , 2005 , 53, 368-74		57
95	The miRNA Mirage: How Close Are We to Finding a Non-Invasive Diagnostic Biomarker in Endometriosis? A Systematic Review. <i>International Journal of Molecular Sciences</i> , 2018 , 19,	6.3	55
94	Coexpression network analysis in abdominal and gluteal adipose tissue reveals regulatory genetic loci for metabolic syndrome and related phenotypes. <i>PLoS Genetics</i> , 2012 , 8, e1002505	6	52
93	Insights into Assessing the Genetics of Endometriosis. <i>Current Obstetrics and Gynecology Reports</i> , 2012 , 1, 124-137	0.6	50
92	Beyond Endometriosis Genome-Wide Association Study: From Genomics to Phenomics to the Patient. <i>Seminars in Reproductive Medicine</i> , 2016 , 34, 242-54	1.4	50
91	Association between endometriosis and the interleukin 1A (IL1A) locus. <i>Human Reproduction</i> , 2015 , 30, 239-48	5.7	49

90	Chronic pelvic pain in women in New Zealand: comparative well-being, comorbidity, and impact on work and other activities. <i>Health Care for Women International</i> , 2006 , 27, 585-99	1.5	49
89	Oral contraceptives and cervical cancer--further findings from the Oxford Family Planning Association contraceptive study. <i>British Journal of Cancer</i> , 1996 , 73, 1291-7	8.7	49
88	Shared genetics underlying epidemiological association between endometriosis and ovarian cancer. <i>Human Molecular Genetics</i> , 2015 , 24, 5955-64	5.6	48
87	Association between endometriosis and N-acetyl transferase 2 polymorphisms in a UK population. <i>Molecular Human Reproduction</i> , 2001 , 7, 1079-83	4.4	48
86	Episiotomies and the occurrence of severe perineal lacerations. <i>BJOG: an International Journal of Obstetrics and Gynaecology</i> , 1994 , 101, 1064-7	3.7	48
85	Significant evidence of one or more susceptibility loci for endometriosis with near-Mendelian inheritance on chromosome 7p13-15. <i>Human Reproduction</i> , 2007 , 22, 717-28	5.7	42
84	Genetic overlap between endometriosis and endometrial cancer: evidence from cross-disease genetic correlation and GWAS meta-analyses. <i>Cancer Medicine</i> , 2018 , 7, 1978-1987	4.8	40
83	MicroRNA expression in abdominal and gluteal adipose tissue is associated with mRNA expression levels and partly genetically driven. <i>PLoS ONE</i> , 2011 , 6, e27338	3.7	39
82	Marker selection for genetic case-control association studies. <i>Nature Protocols</i> , 2009 , 4, 743-52	18.8	38
81	Vascular endothelial growth factor +936 C/T polymorphism is associated with an increased risk of endometriosis in a Japanese population. <i>Acta Obstetrica Et Gynecologica Scandinavica</i> , 2007 , 86, 1352-8	3.8	38
80	Challenges in endometriosis miRNA studies - From tissue heterogeneity to disease specific miRNAs. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2017 , 1863, 2282-2292	6.9	36
79	The influence of menstrual cycle and endometriosis on endometrial methylome. <i>Clinical Epigenetics</i> , 2016 , 8, 2	7.7	35
78	Binding of proteinase 3 and myeloperoxidase to endothelial cells: ANCA-mediated endothelial damage through ADCC?. <i>Clinical and Experimental Immunology</i> , 1994 , 97, 52-60	6.2	35
77	The use of genome-wide eQTL associations in lymphoblastoid cell lines to identify novel genetic pathways involved in complex traits. <i>PLoS ONE</i> , 2011 , 6, e22070	3.7	35
76	Genetics of endometriosis. <i>Women's Health</i> , 2015 , 11, 577-86	3	34
75	Genome-wide association and epidemiological analyses reveal common genetic origins between uterine leiomyomata and endometriosis. <i>Nature Communications</i> , 2019 , 10, 4857	17.4	34
74	The future for genetic studies in reproduction. <i>Molecular Human Reproduction</i> , 2014 , 20, 1-14	4.4	34
73	The ENDOCARE questionnaire (ECQ): a valid and reliable instrument to measure the patient-centeredness of endometriosis care in Europe. <i>Human Reproduction</i> , 2011 , 26, 2988-99	5.7	32

72	Variants in EMX2 and PTEN do not contribute to risk of endometriosis. <i>Molecular Human Reproduction</i> , 2007 , 13, 587-94	4.4	31
71	The role of gene polymorphisms in endometriosis. <i>Molecular Medicine Reports</i> , 2017 , 16, 5881-5886	2.9	30
70	High-density fine-mapping of a chromosome 10q26 linkage peak suggests association between endometriosis and variants close to CYP2C19. <i>Fertility and Sterility</i> , 2011 , 95, 2236-40	4.8	30
69	A simple and fast two-locus quality control test to detect false positives due to batch effects in genome-wide association studies. <i>Genetic Epidemiology</i> , 2010 , 34, 854-62	2.6	30
68	Multivariate genetic analysis of chronic pelvic pain and associated phenotypes. <i>Behavior Genetics</i> , 2005 , 35, 177-88	3.2	28
67	Assessment of the Lequesne index of severity for osteoarthritis of the hip in an elderly population. <i>Osteoarthritis and Cartilage</i> , 2005 , 13, 854-60	6.2	27
66	Endometriosis and cancer: a systematic review and meta-analysis. <i>Human Reproduction Update</i> , 2021 , 27, 393-420	15.8	27
65	DNA methylation changes in endometrium and correlation with gene expression during the transition from pre-receptive to receptive phase. <i>Scientific Reports</i> , 2017 , 7, 3916	4.9	23
64	Population survey comparing older adults with hip versus knee pain in primary care. <i>British Journal of General Practice</i> , 2005 , 55, 192-8	1.6	23
63	Genetic burden associated with varying degrees of disease severity in endometriosis. <i>Molecular Human Reproduction</i> , 2015 , 21, 594-602	4.4	22
62	Prospective study of elderly people comparing treatments following first primary care consultation for a symptomatic hip or knee. <i>Family Practice</i> , 2005 , 22, 118-25	1.9	22
61	The genetic epidemiology of spontaneous endometriosis in the rhesus monkey. <i>Annals of the New York Academy of Sciences</i> , 2002 , 955, 233-8; discussion 293-5, 396-406	6.5	22
60	Gabapentin for chronic pelvic pain in women (GaPP2): a multicentre, randomised, double-blind, placebo-controlled trial. <i>Lancet, The</i> , 2020 , 396, 909-917	4.0	22
59	Mass cytometry analysis reveals a distinct immune environment in peritoneal fluid in endometriosis: a characterisation study. <i>BMC Medicine</i> , 2020 , 18, 3	11.4	21
58	Variability of genome-wide DNA methylation and mRNA expression profiles in reproductive and endocrine disease related tissues. <i>Epigenetics</i> , 2017 , 12, 897-908	5.7	19
57	Uterine fibroids and cardiovascular risk. <i>Human Reproduction</i> , 2016 , 31, 2689-2703	5.7	18
56	Development of a Web site for the genetic epidemiology of endometriosis. <i>Fertility and Sterility</i> , 2002 , 78, 777-81	4.8	18
55	Pain and overall health status in older people with hip and knee replacement: a population perspective. <i>Journal of Public Health</i> , 2006 , 28, 267-73	3.5	17

54	Multi-Center Studies of the Global Impact of Endometriosis and the Predictive Value of Associated Symptoms. <i>Journal of Endometriosis</i> , 2009 , 1, 36-45		16
53	N-acetyl transferase 2 polymorphism and advanced stages of endometriosis in South Indian women. <i>Reproductive BioMedicine Online</i> , 2004 , 9, 533-40	4	15
52	The role of IL-16 gene polymorphisms in endometriosis. <i>International Journal of Molecular Medicine</i> , 2018 , 41, 1469-1476	4.4	13
51	Optimizing the power of genome-wide association studies by using publicly available reference samples to expand the control group. <i>Genetic Epidemiology</i> , 2010 , 34, 319-26	2.6	13
50	The exon 1-8C/G SNP in the PSMA6 gene contributes only a small amount to the burden of myocardial infarction in 6946 cases and 2720 controls from a United Kingdom population. <i>European Journal of Human Genetics</i> , 2008 , 16, 480-6	5.3	13
49	Endometriosis and associated symptoms among Nigerian women. <i>International Journal of Gynecology and Obstetrics</i> , 2015 , 130, 190-4	4	12
48	Common variants in the CYP2C19 gene are associated with susceptibility to endometriosis. <i>Fertility and Sterility</i> , 2014 , 102, 496-502.e5	4.8	12
47	The endothelial nitric oxide synthase Glu298Asp polymorphism is not a risk factor for endometriosis in south Indian women. <i>European Journal of Obstetrics, Gynecology and Reproductive Biology</i> , 2008 , 139, 53-8	2.4	12
46	Characterization of exosomes in peritoneal fluid of endometriosis patients. <i>Fertility and Sterility</i> , 2020 , 113, 364-373.e2	4.8	11
45	The genetic architecture of sporadic and multiple consecutive miscarriage. <i>Nature Communications</i> , 2020 , 11, 5980	17.4	11
44	Large-scale meta-analysis highlights the hypothalamic-pituitary-gonadal axis in the genetic regulation of menstrual cycle length. <i>Human Molecular Genetics</i> , 2018 , 27, 4323-4332	5.6	11
43	No evidence for genetic association with the let-7 microRNA-binding site or other common KRAS variants in risk of endometriosis. <i>Human Reproduction</i> , 2012 , 27, 3616-21	5.7	10
42	DNA methylation alterations-potential cause of endometriosis pathogenesis or a reflection of tissue heterogeneity?. <i>Biology of Reproduction</i> , 2018 , 99, 273-282	3.9	8
41	Coding regions of INHBA, SFRP4 and HOXA10 are not implicated in familial endometriosis linked to chromosome 7p13-15. <i>Molecular Human Reproduction</i> , 2011 , 17, 605-11	4.4	8
40	Genetic analysis of endometriosis and depression identifies shared loci and implicates causal links with gastric mucosa abnormality. <i>Human Genetics</i> , 2021 , 140, 529-552	6.3	8
39	An international terminology for endometriosis, 2021. <i>Human Reproduction Open</i> , 2021 , 2021, hoab029	6.1	8
38	MULTI-CENTRE STUDIES OF THE GLOBAL IMPACT OF ENDOMETRIOSIS AND THE PREDICTIVE VALUE OF ASSOCIATED SYMPTOMS. <i>Journal of Endometriosis</i> , 2009 , 1, 36-45		6
37	Endometriosis classification, staging and reporting systems: a review on the road to a universally accepted endometriosis classification. <i>Human Reproduction Open</i> , 2021 , 2021, hoab025	6.1	6

36	Role of FN1 and GREB1 gene polymorphisms in endometriosis. <i>Molecular Medicine Reports</i> , 2019 , 20, 111-116	2.9	5
35	Endometriosis classification, staging and reporting systems: a review on the road to a universally accepted endometriosis classification. <i>Facts, Views & Vision in ObGyn</i> , 2021 , 13,	1.4	5
34	Machine Learning based histology phenotyping to investigate the epidemiologic and genetic basis of adipocyte morphology and cardiometabolic traits. <i>PLoS Computational Biology</i> , 2020 , 16, e1008044	5	5
33	Genetic analyses of gynecological disease identify genetic relationships between uterine fibroids and endometrial cancer, and a novel endometrial cancer genetic risk region at the WNT4 1p36.12 locus. <i>Human Genetics</i> , 2021 , 140, 1353-1365	6.3	5
32	Amine oxidase 3 is a novel pro-inflammatory marker of oxidative stress in peritoneal endometriosis lesions. <i>Scientific Reports</i> , 2020 , 10, 1495	4.9	4
31	Genome-wide association analysis identifies 27 novel loci associated with uterine leiomyomata revealing common genetic origins with endometriosis		4
30	An International Terminology for Endometriosis, 2021. <i>Facts, Views & Vision in ObGyn</i> , 2021 , 13,	1.4	3
29	Harmonization of Clinical and Laboratory Data to Improve Biomarker Discovery in Endometriosis: WERF EPHect 2017 , 199-226		3
28	Epidemiological and Clinical Risk Factors for Endometriosis 2017 , 95-121		3
27	Genome-wide association analysis and replication in 810,625 individuals identifies novel therapeutic targets for varicose veins		3
26	Epidemiology of chronic pelvic pain. <i>International Congress Series</i> , 2005 , 1279, 77-84		2
25	Protocol for a longitudinal, prospective cohort study investigating the biology of uterine fibroids and endometriosis, and patients' quality of life: the FENOX study. <i>BMJ Open</i> , 2020 , 10, e032220	3	2
24	Genetic overlap analysis of endometriosis and asthma identifies shared loci implicating sex hormones and thyroid signalling pathways.. <i>Human Reproduction</i> , 2022 , 37, 366-383	5.7	2
23	An International Terminology for Endometriosis, 2021. <i>Journal of Minimally Invasive Gynecology</i> , 2021 , 28, 1849-1859	2.2	2
22	Endometriosis Classification, Staging and Reporting Systems: A Review on the Road to a Universally Accepted Endometriosis Classification. <i>Journal of Minimally Invasive Gynecology</i> , 2021 , 28, 1822-1848	2.2	2
21	The genetic architecture of sporadic and recurrent miscarriage		2
20	Neuropeptide S receptor 1 is a nonhormonal treatment target in endometriosis. <i>Science Translational Medicine</i> , 2021 , 13,	17.5	2
19	Understanding the Pathogenesis of Endometriosis: Gene Mapping Studies 54-64		2

18	The role of endometrial B cells in normal endometrium and benign female reproductive pathologies: a systematic review.. <i>Human Reproduction Open</i> , 2022 , 2022, hoab043	6.1	2
17	Cyprus Women's Health Research (COHERE) initiative: determining the relative burden of women's health conditions and related co-morbidities in an Eastern Mediterranean population. <i>BMC Women's Health</i> , 2019 , 19, 50	2.9	1
16	Genetic Association Study Design 2011 , 25-48		1
15	Obesity and risk of female reproductive conditions: A Mendelian randomisation study.. <i>PLoS Medicine</i> , 2022 , 19, e1003679	11.6	1
14	Protocol for the Cultural Translation and Adaptation of the World Endometriosis Research Foundation Endometriosis Phenome and Biobanking Harmonization Project Endometriosis Participant Questionnaire (EPHect). <i>Frontiers in Global Women's Health</i> , 2021 , 2, 644609	8.4	1
13	Genetics and Genomics of Endometriosis 2019 , 399-426		1
12	Impact of Endometriosis in Women of Arab Ancestry on: Health-Related Quality of Life, Work Productivity, and Diagnostic Delay. <i>Frontiers in Global Women's Health</i> , 2021 , 2, 708410	8.4	1
11	Molecular Genetics of Endometriosis		1
10	Using Polygenic Scores in Social Science Research: Unraveling Childlessness. <i>Frontiers in Sociology</i> , 2019 , 4, 74	1.7	0
9	Pathogenesis of Endometriosis: Genetics 2022 , 75-84		0
8	Endometriosis classification systems: an international survey to map current knowledge and uptake.. <i>Human Reproduction Open</i> , 2022 , 2022, hoac002	6.1	
7	The Cyprus Women's Health Research (COHERE) initiative: normative data from the SF-36v2 questionnaire for reproductive aged women from the Eastern Mediterranean.. <i>Quality of Life Research</i> , 2022 , 1	3.7	
6	Machine Learning based histology phenotyping to investigate the epidemiologic and genetic basis of adipocyte morphology and cardiometabolic traits 2020 , 16, e1008044		
5	Machine Learning based histology phenotyping to investigate the epidemiologic and genetic basis of adipocyte morphology and cardiometabolic traits 2020 , 16, e1008044		
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