

Krina T Zondervan

List of Publications by Year in descending order

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Version: 2024-02-01

161
papers

24,477
citations

13854

67
h-index

8384

147
g-index

178
all docs

178
docs citations

178
times ranked

31754
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015, 518, 197-206.	13.7	3,823
2	New genetic loci link adipose and insulin biology to body fat distribution. <i>Nature</i> , 2015, 518, 187-196.	13.7	1,328
3	Data quality control in genetic case-control association studies. <i>Nature Protocols</i> , 2010, 5, 1564-1573.	5.5	1,030
4	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.	0.5	1,020
5	Endometriosis. <i>New England Journal of Medicine</i> , 2020, 382, 1244-1256.	13.9	924
6	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, 949-960.	9.4	836
7	Endometriosis. <i>Nature Reviews Disease Primers</i> , 2018, 4, 9.	18.1	726
8	Mapping cis- and trans-regulatory effects across multiple tissues in twins. <i>Nature Genetics</i> , 2012, 44, 1084-1089.	9.4	701
9	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
10	The complex interplay among factors that influence allelic association. <i>Nature Reviews Genetics</i> , 2004, 5, 89-100.	7.7	480
11	Basic statistical analysis in genetic case-control studies. <i>Nature Protocols</i> , 2011, 6, 121-133.	5.5	426
12	Genome-wide associations for birth weight and correlations with adult disease. <i>Nature</i> , 2016, 538, 248-252.	13.7	406
13	The Architecture of Gene Regulatory Variation across Multiple Human Tissues: The MuTHER Study. <i>PLoS Genetics</i> , 2011, 7, e1002003.	1.5	392
14	Association Between Telomere Length and Risk of Cancer and Non-Neoplastic Diseases. <i>JAMA Oncology</i> , 2017, 3, 636.	3.4	376
15	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, 2624-2634.	0.3	335
16	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, 876-890.	2.6	330
17	Identification of an imprinted master trans regulator at the KLF14 locus related to multiple metabolic phenotypes. <i>Nature Genetics</i> , 2011, 43, 561-564.	9.4	289
18	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.	9.4	286

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19	Priorities for Endometriosis Research: Recommendations From an International Consensus Workshop. <i>Reproductive Sciences</i> , 2009, 16, 335-346.	1.1	284
20	Genome-wide analysis identifies 12 loci influencing human reproductive behavior. <i>Nature Genetics</i> , 2016, 48, 1462-1472.	9.4	284
21	Prevalence and incidence of adults consulting for shoulder conditions in UK primary care; patterns of diagnosis and referral. <i>Rheumatology</i> , 2006, 45, 215-221.	0.9	266
22	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, 1149-1155.	1.1	264
23	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.	2.5	264
24	Genome-wide association study identifies a locus at 7p15.2 associated with endometriosis. <i>Nature Genetics</i> , 2011, 43, 51-54.	9.4	261
25	Genome-wide association meta-analysis identifies new endometriosis risk loci. <i>Nature Genetics</i> , 2012, 44, 1355-1359.	9.4	257
26	What makes a good case-control study?. <i>Human Reproduction</i> , 2002, 17, 1415-1423.	0.4	253
27	Chronic pelvic pain in the community—Symptoms, investigations, and diagnoses. <i>American Journal of Obstetrics and Gynecology</i> , 2001, 184, 1149-1155.	0.7	239
28	Meta-analysis identifies five novel loci associated with endometriosis highlighting key genes involved in hormone metabolism. <i>Nature Communications</i> , 2017, 8, 15539.	5.8	230
29	Genome-Wide Association Study of Susceptibility to Idiopathic Pulmonary Fibrosis. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2020, 201, 564-574.	2.5	208
30	Central changes associated with chronic pelvic pain and endometriosis. <i>Human Reproduction Update</i> , 2014, 20, 737-747.	5.2	192
31	The community prevalence of chronic pelvic pain in women and associated illness behaviour. <i>British Journal of General Practice</i> , 2001, 51, 541-7.	0.7	191
32	Differences in characteristics among 1,000 women with endometriosis based on extent of disease. <i>Fertility and Sterility</i> , 2008, 89, 538-545.	0.5	189
33	Estimation and partitioning of polygenic variation captured by common SNPs for Alzheimer's disease, multiple sclerosis and endometriosis. <i>Human Molecular Genetics</i> , 2013, 22, 832-841.	1.4	186
34	The search for genes contributing to endometriosis risk. <i>Human Reproduction Update</i> , 2008, 14, 447-457.	5.2	181
35	The association between endometriosis and autoimmune diseases: a systematic review and meta-analysis. <i>Human Reproduction Update</i> , 2019, 25, 486-503.	5.2	179
36	Linkage Disequilibrium Mapping via Cladistic Analysis of Single-Nucleotide Polymorphism Haplotypes. <i>American Journal of Human Genetics</i> , 2004, 75, 35-43.	2.6	173

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37	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, 1223-1232.	0.5	171
38	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, 702-716.	5.2	171
39	Human metabolic profiles are stably controlled by genetic and environmental variation. <i>Molecular Systems Biology</i> , 2011, 7, 525.	3.2	158
40	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, 1213-1222.	0.5	154
41	Designing candidate gene and genome-wide case-control association studies. <i>Nature Protocols</i> , 2007, 2, 2492-2501.	5.5	151
42	Distinct Developmental Profile of Lower-Body Adipose Tissue Defines Resistance Against Obesity-Associated Metabolic Complications. <i>Diabetes</i> , 2014, 63, 3785-3797.	0.3	148
43	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, 1233-1243.	0.5	147
44	Peripheral changes in endometriosis-associated pain. <i>Human Reproduction Update</i> , 2014, 20, 717-736.	5.2	135
45	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, 1244-1253.	0.5	134
46	A Genome-Wide Metabolic QTL Analysis in Europeans Implicates Two Loci Shaped by Recent Positive Selection. <i>PLoS Genetics</i> , 2011, 7, e1002270.	1.5	132
47	Research Priorities for Endometriosis: Recommendations From a Global Consortium of Investigators in Endometriosis. <i>Reproductive Sciences</i> , 2017, 24, 202-226.	1.1	124
48	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, 369-375.	0.8	123
49	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.	0.5	123
50	The genetic basis of endometriosis. <i>Current Opinion in Obstetrics and Gynecology</i> , 2001, 13, 309-314.	0.9	114
51	Endometriosis and cancer: a systematic review and meta-analysis. <i>Human Reproduction Update</i> , 2021, 27, 393-420.	5.2	112
52	High-quality genomic DNA extraction from formalin-fixed and paraffin-embedded samples deparaffinized using mineral oil. <i>Analytical Biochemistry</i> , 2009, 395, 265-267.	1.1	109
53	Recent insights on the genetics and epigenetics of endometriosis. <i>Clinical Genetics</i> , 2017, 91, 254-264.	1.0	106
54	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.	0.5	104

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55	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, 1156-1161.	1.1	97
56	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, 93-99.	1.1	90
57	Genome-wide association and epidemiological analyses reveal common genetic origins between uterine leiomyomata and endometriosis. <i>Nature Communications</i> , 2019, 10, 4857.	5.8	90
58	Familial aggregation of endometriosis in a large pedigree of rhesus macaques. <i>Human Reproduction</i> , 2004, 19, 448-455.	0.4	88
59	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, 599.	1.8	86
60	The management of menopause in women with a history of endometriosis: a systematic review. <i>Human Reproduction Update</i> , 2017, 23, 481-500.	5.2	84
61	Do Dietary and Supplementary Intakes of Antioxidants Differ with Smoking Status?. <i>International Journal of Epidemiology</i> , 1996, 25, 70-79.	0.9	83
62	Epidemiology of chronic pelvic pain. <i>Best Practice and Research in Clinical Obstetrics and Gynaecology</i> , 2000, 14, 403-414.	1.4	83
63	Genome-wide Linkage and Association Analyses Implicate FASN in Predisposition to Uterine Leiomyomata. <i>American Journal of Human Genetics</i> , 2012, 91, 621-628.	2.6	83
64	The vascular endothelial growth factor (VEGF) +405G>C 5'â€²-untranslated region polymorphism and increased risk of endometriosis in South Indian women: a case control study. <i>Human Reproduction</i> , 2005, 20, 1844-1849.	0.4	82
65	Genome-wide genetic analyses highlight mitogen-activated protein kinase (MAPK) signaling in the pathogenesis of endometriosis. <i>Human Reproduction</i> , 2017, 32, 780-793.	0.4	81
66	Lymphotoxin-Î± 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.	1.5	77
67	Variability of gene expression profiles in human blood and lymphoblastoid cell lines. <i>BMC Genomics</i> , 2010, 11, 96.	1.2	75
68	Genome-wide enrichment analysis between endometriosis and obesity-related traits reveals novel susceptibility loci. <i>Human Molecular Genetics</i> , 2015, 24, 1185-1199.	1.4	71
69	Shared genetics underlying epidemiological association between endometriosis and ovarian cancer. <i>Human Molecular Genetics</i> , 2015, 24, 5955-5964.	1.4	68
70	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, 368-374.	6.7	64
71	Episiotomies and the occurrence of severe perineal lacerations. <i>BJOG: an International Journal of Obstetrics and Gynaecology</i> , 1994, 101, 1064-1067.	1.1	63
72	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, 585-599.	0.6	63

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73	Beyond Endometriosis Genome-Wide Association Study: From Genomics to Phenomics to the Patient. <i>Seminars in Reproductive Medicine</i> , 2016, 34, 242-254.	0.5	62
74	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.	1.3	62
75	Insights into Assessing the Genetics of Endometriosis. <i>Current Obstetrics and Gynecology Reports</i> , 2012, 1, 124-137.	0.3	58
76	Association between endometriosis and the interleukin 1A (IL1A) locus. <i>Human Reproduction</i> , 2015, 30, 239-248.	0.4	58
77	Oral contraceptives and cervical cancer - further findings from the Oxford Family Planning Association contraceptive study. <i>British Journal of Cancer</i> , 1996, 73, 1291-1297.	2.9	57
78	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.	1.5	57
79	The influence of menstrual cycle and endometriosis on endometrial methylome. <i>Clinical Epigenetics</i> , 2016, 8, 2.	1.8	57
80	Association between endometriosis and N-acetyl transferase 2 polymorphisms in a UK population. <i>Molecular Human Reproduction</i> , 2001, 7, 1079-1083.	1.3	54
81	Significant evidence of one or more susceptibility loci for endometriosis with near-Mendelian inheritance on chromosome 7p13-15. <i>Human Reproduction</i> , 2007, 22, 717-728.	0.4	54
82	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.	1.8	52
83	The genetic architecture of sporadic and multiple consecutive miscarriage. <i>Nature Communications</i> , 2020, 11, 5980.	5.8	52
84	Obesity and risk of female reproductive conditions: A Mendelian randomisation study. <i>PLoS Medicine</i> , 2022, 19, e1003679.	3.9	50
85	Mass cytometry analysis reveals a distinct immune environment in peritoneal fluid in endometriosis: a characterisation study. <i>BMC Medicine</i> , 2020, 18, 3.	2.3	49
86	An International Terminology for Endometriosis, 2021. <i>Journal of Minimally Invasive Gynecology</i> , 2021, 28, 1849-1859.	0.3	49
87	The role of gene polymorphisms in endometriosis. <i>Molecular Medicine Reports</i> , 2017, 16, 5881-5886.	1.1	48
88	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.	1.1	46
89	Genetics of Endometriosis. <i>Women's Health</i> , 2015, 11, 577-586.	0.7	44
90	Marker selection for genetic case-control association studies. <i>Nature Protocols</i> , 2009, 4, 743-752.	5.5	43

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91	Binding of proteinase 3 and myeloperoxidase to endothelial cells: ANCA-mediated endothelial damage through ADCC?. <i>Clinical and Experimental Immunology</i> , 2008, 97, 52-60.	1.1	42
92	Gabapentin for chronic pelvic pain in women (GaPP2): a multicentre, randomised, double-blind, placebo-controlled trial. <i>Lancet, The</i> , 2020, 396, 909-917.	6.3	42
93	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, 2988-2999.	0.4	41
94	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, 1352-1358.	1.3	40
95	The future for genetic studies in reproduction. <i>Molecular Human Reproduction</i> , 2014, 20, 1-14.	1.3	38
96	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.	1.6	37
97	Assessment of the Lequesne index of severity for osteoarthritis of the hip in an elderly population. <i>Osteoarthritis and Cartilage</i> , 2005, 13, 854-860.	0.6	36
98	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, 2236-2240.	0.5	36
99	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.	1.8	36
100	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.	1.1	36
101	Characterization of exosomes in peritoneal fluid of endometriosis patients. <i>Fertility and Sterility</i> , 2020, 113, 364-373.e2.	0.5	35
102	Variants in EMX2 and PTEN do not contribute to risk of endometriosis. <i>Molecular Human Reproduction</i> , 2007, 13, 587-594.	1.3	34
103	An international terminology for endometriosis, 2021,. <i>Human Reproduction Open</i> , 2021, 2021, hoab029.	2.3	34
104	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, 854-862.	0.6	33
105	Uterine fibroids and cardiovascular risk. <i>Human Reproduction</i> , 2016, 31, 2689-2703.	0.4	33
106	Variability of genome-wide DNA methylation and mRNA expression profiles in reproductive and endocrine disease related tissues. <i>Epigenetics</i> , 2017, 12, 897-908.	1.3	33
107	Multivariate Genetic Analysis of Chronic Pelvic Pain and Associated Phenotypes. <i>Behavior Genetics</i> , 2005, 35, 177-188.	1.4	32
108	Prospective study of elderly people comparing treatments following first primary care consultation for a symptomatic hip or knee. <i>Family Practice</i> , 2004, 22, 118-125.	0.8	30

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109	Genetic burden associated with varying degrees of disease severity in endometriosis. <i>Molecular Human Reproduction</i> , 2015, 21, 594-602.	1.3	30
110	The Genetic Epidemiology of Spontaneous Endometriosis in the Rhesus Monkey. <i>Annals of the New York Academy of Sciences</i> , 2002, 955, 233-238.	1.8	23
111	Neuropeptide S receptor 1 is a nonhormonal treatment target in endometriosis. <i>Science Translational Medicine</i> , 2021, 13, .	5.8	23
112	Population survey comparing older adults with hip versus knee pain in primary care. <i>British Journal of General Practice</i> , 2005, 55, 192-8.	0.7	23
113	Pain and overall health status in older people with hip and knee replacement: a population perspective. <i>Journal of Public Health</i> , 2006, 28, 267-273.	1.0	22
114	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.	1.0	22
115	Endometriosis and associated symptoms among Nigerian women. <i>International Journal of Gynecology and Obstetrics</i> , 2015, 130, 190-194.	1.0	22
116	Development of a web site for the genetic epidemiology of endometriosis. <i>Fertility and Sterility</i> , 2002, 78, 777-781.	0.5	21
117	OUP accepted manuscript. <i>Human Reproduction Open</i> , 2021, 2021, hoab025.	2.3	21
118	Large-scale meta-analysis highlights the hypothalamicâ€“pituitaryâ€“gonadal axis in the genetic regulation of menstrual cycle length. <i>Human Molecular Genetics</i> , 2018, 27, 4323-4332.	1.4	20
119	An International Terminology for Endometriosis, 2021. <i>Facts, Views & Vision in ObGyn</i> , 2021, 13, 295-304.	0.5	20
120	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.	0.3	19
121	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.	0.4	19
122	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.	1.8	18
123	N-acetyl transferase 2 polymorphism and advanced stages of endometriosis in South Indian women. <i>Reproductive BioMedicine Online</i> , 2004, 9, 533-540.	1.1	17
124	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, 319-326.	0.6	17
125	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.	1.5	16
126	Common variants in the CYP2C19 gene are associated with susceptibility to endometriosis. <i>Fertility and Sterility</i> , 2014, 102, 496-502.e5.	0.5	15

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127	The role of IL16 gene polymorphisms in endometriosis. <i>International Journal of Molecular Medicine</i> , 2018, 41, 1469-1476.	1.8	15
128	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, 480-486.	1.4	14
129	Role of FN1 and GREB1 gene polymorphisms in endometriosis. <i>Molecular Medicine Reports</i> , 2019, 20, 111-116.	1.1	14
130	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, 3616-3621.	0.4	13
131	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.	1.1	13
132	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, 53-58.	0.5	12
133	DNA methylation alterations potential cause of endometriosis pathogenesis or a reflection of tissue heterogeneity? <i>Biology of Reproduction</i> , 2018, 99, 273-282.	1.2	11
134	Amine oxidase 3 is a novel pro-inflammatory marker of oxidative stress in peritoneal endometriosis lesions. <i>Scientific Reports</i> , 2020, 10, 1495.	1.6	11
135	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, 305-330.	0.5	10
136	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.	2.3	10
137	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, 605-611.	1.3	9
138	Genome-wide association analysis and replication in 810,625 individuals with varicose veins. <i>Nature Communications</i> , 2022, 13, .	5.8	8
139	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.	1.0	7
140	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.	0.8	6
141	Endometriosis Classification Systems: An International Survey to Map Current Knowledge and Uptake. <i>Journal of Minimally Invasive Gynecology</i> , 2022, 29, 716-725.e1.	0.3	6
142	Using Polygenic Scores in Social Science Research: Unraveling Childlessness. <i>Frontiers in Sociology</i> , 2019, 4, 74.	1.0	4
143	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.	0.8	4
144	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.	1.1	4

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145	Epidemiological and Clinical Risk Factors for Endometriosis. , 2017, , 95-121.		4
146	Harmonization of Clinical and Laboratory Data to Improve Biomarker Discovery in Endometriosis: WERF EPHect. , 2017, , 199-226.		3
147	The Cyprus Womenâ€™s Health Research (COHERE) initiative: normative data from the SF-36v2 questionnaire for reproductive aged women from the Eastern Mediterranean. Quality of Life Research, 2022, 31, 2011-2022.	1.5	3
148	Epidemiology of chronic pelvic pain. International Congress Series, 2005, 1279, 77-84.	0.2	2
149	Genetic Association Study Design. , 2011, , 25-48.		2
150	Genetics and Genomics of Endometriosis. , 2019, , 399-426.		2
151	Session 05: Endometriosis: Impact, Diagnosis and Surgery. Human Reproduction, 2010, 25, i9-i11.	0.4	1
152	Genome-Wide Association Study Identifies a Locus at 7p15.2 Associated With Endometriosis. Obstetrical and Gynecological Survey, 2011, 66, 214-216.	0.2	0
153	1625â€™Genome-Wide Association Analysis and Replication In 810,625 Individuals Identifies Novel Therapeutic Targets for Varicose Veins. British Journal of Surgery, 2021, 108, .	0.1	0
154	OUP accepted manuscript. Human Reproduction Open, 2022, 2022, hoac002.	2.3	0
155	Endometriosis classification systems: an international survey to map current knowledge and uptake. Facts, Views & Vision in ObGyn, 2022, 14, 5-15.	0.5	0
156	Title is missing!. , 2020, 16, e1008044.		0
157	Title is missing!. , 2020, 16, e1008044.		0
158	Title is missing!. , 2020, 16, e1008044.		0
159	Title is missing!. , 2020, 16, e1008044.		0
160	Title is missing!. , 2020, 16, e1008044.		0
161	Title is missing!. , 2020, 16, e1008044.		0