Kosuke Yoshihara

List of Publications by Year in descending order

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Version: 2024-02-01

95 papers 13,651 citations

33 h-index 84 g-index

101 all docs

101 docs citations

101 times ranked

21181 citing authors

#	Article	IF	CITATIONS
1	Racial and ethnic disparity in characteristics and outcomes of women with placenta accreta spectrum: a comparative study. Reproductive Sciences, 2022, 29, 1988-2000.	2.5	5
2	Genetic overlap analysis of endometriosis and asthma identifies shared loci implicating sex hormones and thyroid signalling pathways. Human Reproduction, 2022, 37, 366-383.	0.9	19
3	APOBEC mediated mutagenesis drives genomic heterogeneity in endometriosis. Journal of Human Genetics, 2022, 67, 323-329.	2.3	3
4	Spatiotemporal dynamics of clonal selection and diversification in normal endometrial epithelium. Nature Communications, 2022, 13, 943.	12.8	24
5	Population incidence and characteristics of secondary breast cancer after uterine cancer: a competing risk analysis. Archives of Gynecology and Obstetrics, 2022, , 1.	1.7	O
6	Fetal biometric and Doppler measurements following abdominal radical trachelectomy in the second trimester of the pregnancy. BMC Pregnancy and Childbirth, 2022, 22, 343.	2.4	0
7	Phase II study of niraparib in recurrent or persistent rare fraction of gynecologic malignancies with homologous recombination deficiency (JGOG2052). Journal of Gynecologic Oncology, 2022, 33, .	2.2	4
8	Therapeutic Strategies Focused on Cancer-Associated Hypercoagulation for Ovarian Clear Cell Carcinoma. Cancers, 2022, 14, 2125.	3.7	7
9	Sentinel node navigation surgery in cervical cancer: a systematic review and metaanalysis. International Journal of Clinical Oncology, 2022, 27, 1247-1255.	2.2	6
10	Trends in Pregnancy-Associated Cervical Cancer in Japan between 2012 and 2017: A Multicenter Survey. Cancers, 2022, 14, 3072.	3.7	1
11	Genetic analysis of endometriosis and depression identifies shared loci and implicates causal links with gastric mucosa abnormality. Human Genetics, 2021, 140, 529-552.	3.8	36
12	ASO Author Reflections: Clinical Significance of Mesenteric Lymph Node Involvement in Patients with Ovarian Cancer. Annals of Surgical Oncology, 2021, 28, 7614-7615.	1.5	0
13	How Does Endometriosis Lead to Ovarian Cancer? The Molecular Mechanism of Endometriosis-Associated Ovarian Cancer Development. Cancers, 2021, 13, 1439.	3.7	19
14	Establishment of in vitro 3D spheroid cell cultivation from human gynecologic cancer tissues. STAR Protocols, 2021, 2, 100354.	1.2	4
15	Biological significance of KRAS mutant allele expression in ovarian endometriosis. Cancer Science, 2021, 112, 2020-2032.	3.9	18
16	Clinical Significance of Mesenteric Lymph Node Involvement in the Pattern of Liver Metastasis in Patients with Ovarian Cancer. Annals of Surgical Oncology, 2021, 28, 7606-7613.	1.5	5
17	PET/MR imaging for the evaluation of cervical cancer during pregnancy. BMC Pregnancy and Childbirth, 2021, 21, 288.	2.4	6
18	Three-dimensional understanding of the morphological complexity of the human uterine endometrium. IScience, 2021, 24, 102258.	4.1	59

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19	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. Human Genetics, 2021, 140, 1353-1365.	3.8	18
20	The New Era of Three-Dimensional Histoarchitecture of the Human Endometrium. Journal of Personalized Medicine, 2021, 11, 713.	2.5	8
21	Frequency of homologous recombination associated gene mutations in Japanese patients with ovarian cancer. Gynecologic Oncology, 2021, 162, S163-S164.	1.4	0
22	Proposing a molecular classification associated with hypercoagulation in ovarian clear cell carcinoma. Gynecologic Oncology, 2021, 163, 327-333.	1.4	5
23	Sleeping Beauty Transposon Mutagenesis Identifies Genes Driving the Initiation and Metastasis of Uterine Leiomyosarcoma. Cancer Research, 2021, 81, 5413-5424.	0.9	2
24	Integrative analyses of gene expression and chemosensitivity of patient-derived ovarian cancer spheroids link G6PD-driven redox metabolism to cisplatin chemoresistance. Cancer Letters, 2021, 521, 29-38.	7.2	23
25	Clinical relevance of TP53 hotspot mutations in high-grade serous ovarian cancers. British Journal of Cancer, 2020, 122, 405-412.	6.4	53
26	ARID1A protein expression is retained in ovarian endometriosis with ARID1A loss-of-function mutations: implication for the two-hit hypothesis. Scientific Reports, 2020, 10, 14260.	3.3	18
27	Temporal trends of subsequent breast cancer among women with ovarian cancer: a population-based study. Archives of Gynecology and Obstetrics, 2020, 301, 1235-1245.	1.7	6
28	Shared Molecular Genetic Mechanisms Underlie Endometriosis and Migraine Comorbidity. Genes, 2020, 11, 268.	2.4	53
29	XCL1 expression correlates with CD8-positive T cells infiltration and PD-L1 expression in squamous cell carcinoma arising from mature cystic teratoma of the ovary. Oncogene, 2020, 39, 3541-3554.	5.9	26
30	The rapid adoption of opportunistic salpingectomy at the time of hysterectomy for benign gynecologic disease in the United States. American Journal of Obstetrics and Gynecology, 2020, 223, 721.e1-721.e18.	1.3	28
31	Clonal lineage from normal endometrium to ovarian clear cell carcinoma through ovarian endometriosis. Cancer Science, 2020, 111, 3000-3009.	3.9	34
32	Association of g BRCA1/2 mutation locations with ovarian cancer risk in Japanese patients from the CHARLOTTE study. Cancer Science, 2020, 111, 3350-3358.	3.9	7
33	The first Japanese nationwide multicenter study of <i>BRCA</i> mutation testing in ovarian cancer: CHARacterizing the cross-sectionaL approach to Ovarian cancer geneTic TEsting of <i>BRCA</i> (CHARLOTTE). International Journal of Gynecological Cancer, 2019, 29, 1043-1049.	2.5	80
34	ALDH-Dependent Glycolytic Activation Mediates Stemness and Paclitaxel Resistance in Patient-Derived Spheroid Models of Uterine Endometrial Cancer. Stem Cell Reports, 2019, 13, 730-746.	4.8	59
35	Different mutation profiles between epithelium and stroma in endometriosis and normal endometrium. Human Reproduction, 2019, 34, 1899-1905.	0.9	37
36	Recurrence, death, and secondary malignancy after ovarian conservation for young women with early-stage low-grade endometrial cancer. Gynecologic Oncology, 2019, 155, 39-50.	1.4	16

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37	Decreasing Trends of Secondary Primary Colorectal Cancer among Women with Uterine Cancer: A Population-Based Analysis. Journal of Clinical Medicine, 2019, 8, 714.	2.4	3
38	Decreasing secondary primary uterine cancer after breast cancer: A population-based analysis. Gynecologic Oncology, 2019, 154, 169-176.	1.4	8
39	Concurrent isolated retroperitoneal HGSC and STIC defined by somatic mutation analysis: a case report. Diagnostic Pathology, 2019, 14, 17.	2.0	4
40	Germline and somatic mutations of homologous recombination-associated genes in Japanese ovarian cancer patients. Scientific Reports, 2019, 9, 17808.	3.3	38
41	The 61st Annual Meeting of the Japanese Society for Gynecologic Oncology (JSGO). Journal of Gynecologic Oncology, 2019, 30, e114.	2.2	0
42	The Safety and Effectiveness of Abdominal Radical Trachelectomy for Early-Stage Cervical Cancer During Pregnancy. International Journal of Gynecological Cancer, 2018, 28, 782-787.	2.5	17
43	Novel therapeutic strategy for cervical cancer harboring FGFR3-TACC3 fusions. Oncogenesis, 2018, 7, 4.	4.9	41
44	TumorFusions: an integrative resource for cancer-associated transcript fusions. Nucleic Acids Research, 2018, 46, D1144-D1149.	14.5	179
45	Effectiveness of fetal cardiac screening for congenital heart disease using a combination of the fourâ€chamber view and threeâ€vessel view during the second trimester scan. Journal of Obstetrics and Gynaecology Research, 2018, 44, 49-53.	1.3	14
46	The impact of stromal Hic-5 on the tumorigenesis of colorectal cancer through lysyl oxidase induction and stromal remodeling. Oncogene, 2018, 37, 1205-1219.	5.9	27
47	Novel <i>MXD4–NUTM1</i> fusion transcript identified in primary ovarian undifferentiated small round cell sarcoma. Genes Chromosomes and Cancer, 2018, 57, 557-563.	2.8	28
48	Clonal Expansion and Diversification of Cancer-Associated Mutations in Endometriosis and Normal Endometrium. Cell Reports, 2018, 24, 1777-1789.	6.4	296
49	Meta-analysis identifies five novel loci associated with endometriosis highlighting key genes involved in hormone metabolism. Nature Communications, 2017, 8, 15539.	12.8	230
50	Sox2â€dependent inhibition of p21 is associated with poor prognosis of endometrial cancer. Cancer Science, 2017, 108, 632-640.	3.9	29
51	Whole-genome sequencing revealed novel prognostic biomarkers and promising targets for therapy of ovarian clear cell carcinoma. British Journal of Cancer, 2017, 117, 717-724.	6.4	78
52	In vivo loss-of-function screens identify KPNB1 as a new druggable oncogene in epithelial ovarian cancer. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E7301-E7310.	7.1	88
53	Abstract 3383: Exome sequencing in dedifferentiated ovarian mucinous carcinoma. , 2017, , .		0
54	Abstract 411:In vivopooled shRNA library identifies KPNB1 as a new drug target for epithelial ovarian cancer. , 2017, , .		0

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55	Abstract 529: The significance of activated PI3K/AKT pathway in FGFR3-TACC3 fusion positive cervical cancer., 2017, , .		O
56	Allelic Imbalance in Regulation of ANRIL through Chromatin Interaction at 9p21 Endometriosis Risk Locus. PLoS Genetics, 2016, 12, e1005893.	3.5	40
57	Association of <i>NR3C1</i> / <i>Glucocorticoid Receptor</i> gene SNP with azoospermia in Japanese men. Journal of Obstetrics and Gynaecology Research, 2016, 42, 59-66.	1.3	6
58	Two-Step Forward Genetic Screen in Mice Identifies RalÂGTPase-Activating Proteins as Suppressors of HepatocellularÂCarcinoma. Gastroenterology, 2016, 151, 324-337.e12.	1.3	27
59	Novel kinase fusion transcripts found in endometrial cancer. Scientific Reports, 2016, 5, 18657.	3.3	11
60	Association of Low-Dose Aspirin and Survival of Women With Endometrial Cancer. Obstetrics and Gynecology, 2016, 128, 127-137.	2.4	39
61	Transposon mutagenesis identifies genes and cellular processes driving epithelial-mesenchymal transition in hepatocellular carcinoma. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E3384-93.	7.1	56
62	Establishment of a Novel Histopathological Classification of High-Grade Serous Ovarian Carcinoma Correlated with Prognostically Distinct Gene Expression Subtypes. American Journal of Pathology, 2016, 186, 1103-1113.	3.8	71
63	Similar protein expression profiles of ovarian and endometrial high-grade serous carcinomas. British Journal of Cancer, 2016, 114, 554-561.	6.4	22
64	Abstract 1504: Identification of novel kinase fusion transcripts in endometrial cancer., 2016,,.		0
65	Abstract 99: Ovarian cancer specific therapeutic vulnerability. , 2016, , .		0
66	Identification of novel exonic mobile element insertions in epithelial ovarian cancers. Human Genome Variation, 2015, 2, 15030.	0.7	2
67	Somatic Copy Number Alterations Associated with Japanese or Endometriosis in Ovarian Clear Cell Adenocarcinoma. PLoS ONE, 2015, 10, e0116977.	2.5	28
68	Comprehensive, Integrative Genomic Analysis of Diffuse Lower-Grade Gliomas. New England Journal of Medicine, 2015, 372, 2481-2498.	27.0	2,582
69	Susceptibility to male infertility: replication study in Japanese men looking for an association with four GWAS-derived loci identified in European men. Journal of Assisted Reproduction and Genetics, 2015, 32, 903-908.	2.5	3
70	The landscape and therapeutic relevance of cancer-associated transcript fusions. Oncogene, 2015, 34, 4845-4854.	5.9	398
71	Comparison of gene expression patterns across 12 tumor types identifies a cancer supercluster characterized by TP53 mutations and cell cycle defects. Oncogene, 2015, 34, 2732-2740.	5.9	46
72	Histone demethylase JARID1C inactivation triggers genomic instability in sporadic renal cancer. Journal of Clinical Investigation, 2015, 125, 4625-4637.	8.2	62

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73	Abstract 4795: A novel gene fusion in glioblastoma and a radiation response methylation signature identified by genomic characterization of glioma sphere-forming cells. , 2015, , .		O
74	Abstract 3762: The landscape of therapeutic targetable fusions. , 2015, , .		0
75	Molecular Characterization of an Intact p53 Pathway Subtype in High-Grade Serous Ovarian Cancer. PLoS ONE, 2014, 9, e114491.	2.5	17
76	Hiding in the dark: uncovering cancer drivers through image-guided genomics. Genome Biology, 2014, 15, 563.	8.8	6
77	Promoter methylation of DAPK1, FHIT, MGMT, and CDKN2A genes in cervical carcinoma. International Journal of Clinical Oncology, 2014, 19, 127-132.	2.2	43
78	Multiplatform Analysis of 12 Cancer Types Reveals Molecular Classification within and across Tissues of Origin. Cell, 2014, 158, 929-944.	28.9	1,242
79	A pan-cancer proteomic perspective on The Cancer Genome Atlas. Nature Communications, 2014, 5, 3887.	12.8	456
80	A nonsynonymous variant of IL1A is associated with endometriosis in Japanese population. Journal of Human Genetics, 2013, 58, 517-520.	2.3	25
81	Inferring tumour purity and stromal and immune cell admixture from expression data. Nature Communications, 2013, 4, 2612.	12.8	5,788
82	Increased incidence of brain metastases in <i>BRCA1</i> â€related ovarian cancers. Journal of Obstetrics and Gynaecology Research, 2013, 39, 292-296.	1.3	30
83	Predicting time to ovarian carcinoma recurrence using protein markers. Journal of Clinical Investigation, 2013, 123, 3740-50.	8.2	46
84	Predicting time to ovarian carcinoma recurrence using protein markers. Journal of Clinical Investigation, 2013, 123, 5410-5410.	8.2	24
85	Possible involvement of the E-cadherin gene in genetic susceptibility to endometriosis. Human Reproduction, 2012, 27, 1685-1689.	0.9	18
86	High-Risk Ovarian Cancer Based on 126-Gene Expression Signature Is Uniquely Characterized by Downregulation of Antigen Presentation Pathway. Clinical Cancer Research, 2012, 18, 1374-1385.	7.0	165
87	Germline copy number variations in <i>BRCA1</i> â€associated ovarian cancer patients. Genes Chromosomes and Cancer, 2011, 50, 167-177.	2.8	37
88	Changes in Fetal Circulation Associated with Congenital Heart Disease and Their Effects on Fetal Growth. Fetal Diagnosis and Therapy, 2011, 30, 219-224.	1.4	28
89	Identification of Receptor Tyrosine Kinase, Discoidin Domain Receptor 1 (DDR1), as a Potential Biomarker for Serous Ovarian Cancer. International Journal of Molecular Sciences, 2011, 12, 971-982.	4.1	58
90	Serum leptin–adiponectin ratio and endometrial cancer risk in postmenopausal female subjects. Gynecologic Oncology, 2010, 119, 65-69.	1.4	97

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91	Meta-analysis of genome-wide association scans for genetic susceptibility to endometriosis in Japanese population. Journal of Human Genetics, 2010, 55, 816-821.	2.3	87
92	Gene Expression Profile for Predicting Survival in Advanced-Stage Serous Ovarian Cancer Across Two Independent Datasets. PLoS ONE, 2010, 5, e9615.	2.5	124
93	Abstract 2162: Gene expression profile for predicting survival in ovarian cancer across two independent datasets., 2010,,.		O
94	Gene expression profiling of advancedâ€stage serous ovarian cancers distinguishes novel subclasses and implicates <i>ZEB2</i> in tumor progression and prognosis. Cancer Science, 2009, 100, 1421-1428.	3.9	168
95	Association of single nucleotide polymorphisms in adiponectin and its receptor genes with polycystic ovary syndrome. Journal of reproductive medicine, The, 2009, 54, 669-74.	0.2	7