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	Inferring tumour purity and stromal and immune cell admixture from expression data. Nature Communications, 2013, 4, 2612.	12.8	5,788
2	Comprehensive, Integrative Genomic Analysis of Diffuse Lower-Grade Gliomas. New England Journal of Medicine, 2015, 372, 2481-2498.	27.0	2,582
3	Multiplatform Analysis of 12 Cancer Types Reveals Molecular Classification within and across Tissues of Origin. Cell, 2014, 158, 929-944.	28.9	1,242
4	A pan-cancer proteomic perspective on The Cancer Genome Atlas. Nature Communications, 2014, 5, 3887.	12.8	456
5	The landscape and therapeutic relevance of cancer-associated transcript fusions. Oncogene, 2015, 34, 4845-4854.	5.9	398
6	Clonal Expansion and Diversification of Cancer-Associated Mutations in Endometriosis and Normal Endometrium. Cell Reports, 2018, 24, 1777-1789.	6.4	296
7	Meta-analysis identifies five novel loci associated with endometriosis highlighting key genes involved in hormone metabolism. Nature Communications, 2017, 8, 15539.	12.8	230
8	TumorFusions: an integrative resource for cancer-associated transcript fusions. Nucleic Acids Research, 2018, 46, D1144-D1149.	14.5	179
9	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
10	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
11	Gene Expression Profile for Predicting Survival in Advanced-Stage Serous Ovarian Cancer Across Two Independent Datasets. PLoS ONE, 2010, 5, e9615.	2.5	124
12	Serum leptin–adiponectin ratio and endometrial cancer risk in postmenopausal female subjects. Gynecologic Oncology, 2010, 119, 65-69.	1.4	97
13	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
14	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
15	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
16	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
17	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
18	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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19	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
20	Three-dimensional understanding of the morphological complexity of the human uterine endometrium. IScience, 2021, 24, 102258.	4.1	59
21	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
22	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
23	Clinical relevance of TP53 hotspot mutations in high-grade serous ovarian cancers. British Journal of Cancer, 2020, 122, 405-412.	6.4	53
24	Shared Molecular Genetic Mechanisms Underlie Endometriosis and Migraine Comorbidity. Genes, 2020, 11, 268.	2.4	53
25	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
26	Predicting time to ovarian carcinoma recurrence using protein markers. Journal of Clinical Investigation, 2013, 123, 3740-50.	8.2	46
27	Promoter methylation of DAPK1, FHIT, MGMT, and CDKN2A genes in cervical carcinoma. International Journal of Clinical Oncology, 2014, 19, 127-132.	2.2	43
28	Novel therapeutic strategy for cervical cancer harboring FGFR3-TACC3 fusions. Oncogenesis, 2018, 7, 4.	4.9	41
29	Allelic Imbalance in Regulation of ANRIL through Chromatin Interaction at 9p21 Endometriosis Risk Locus. PLoS Genetics, 2016, 12, e1005893.	3.5	40
30	Association of Low-Dose Aspirin and Survival of Women With Endometrial Cancer. Obstetrics and Gynecology, 2016, 128, 127-137.	2.4	39
31	Germline and somatic mutations of homologous recombination-associated genes in Japanese ovarian cancer patients. Scientific Reports, 2019, 9, 17808.	3.3	38
32	Germline copy number variations in <i>BRCA1</i> â€associated ovarian cancer patients. Genes Chromosomes and Cancer, 2011, 50, 167-177.	2.8	37
33	Different mutation profiles between epithelium and stroma in endometriosis and normal endometrium. Human Reproduction, 2019, 34, 1899-1905.	0.9	37
34	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
35	Clonal lineage from normal endometrium to ovarian clear cell carcinoma through ovarian endometriosis. Cancer Science, 2020, 111, 3000-3009.	3.9	34
36	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

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37	Sox2â€dependent inhibition of p21 is associated with poor prognosis of endometrial cancer. Cancer Science, 2017, 108, 632-640.	3.9	29
38	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
39	Somatic Copy Number Alterations Associated with Japanese or Endometriosis in Ovarian Clear Cell Adenocarcinoma. PLoS ONE, 2015, 10, e0116977.	2.5	28
40	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
41	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
42	Two-Step Forward Genetic Screen in Mice Identifies RalÂGTPase-Activating Proteins as Suppressors of HepatocellularÂCarcinoma. Gastroenterology, 2016, 151, 324-337.e12.	1.3	27
43	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
44	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
45	A nonsynonymous variant of IL1A is associated with endometriosis in Japanese population. Journal of Human Genetics, 2013, 58, 517-520.	2.3	25
46	Predicting time to ovarian carcinoma recurrence using protein markers. Journal of Clinical Investigation, 2013, 123, 5410-5410.	8.2	24
47	Spatiotemporal dynamics of clonal selection and diversification in normal endometrial epithelium. Nature Communications, 2022, 13, 943.	12.8	24
48	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
49	Similar protein expression profiles of ovarian and endometrial high-grade serous carcinomas. British Journal of Cancer, 2016, 114, 554-561.	6.4	22
50	How Does Endometriosis Lead to Ovarian Cancer? The Molecular Mechanism of Endometriosis-Associated Ovarian Cancer Development. Cancers, 2021, 13, 1439.	3.7	19
51	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
52	Possible involvement of the E-cadherin gene in genetic susceptibility to endometriosis. Human Reproduction, 2012, 27, 1685-1689.	0.9	18
53	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
54	Biological significance of KRAS mutant allele expression in ovarian endometriosis. Cancer Science, 2021, 112, 2020-2032.	3.9	18

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55	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
56	Molecular Characterization of an Intact p53 Pathway Subtype in High-Grade Serous Ovarian Cancer. PLoS ONE, 2014, 9, e114491.	2.5	17
57	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
58	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
59	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
60	Novel kinase fusion transcripts found in endometrial cancer. Scientific Reports, 2016, 5, 18657.	3.3	11
61	Decreasing secondary primary uterine cancer after breast cancer: A population-based analysis. Gynecologic Oncology, 2019, 154, 169-176.	1.4	8
62	The New Era of Three-Dimensional Histoarchitecture of the Human Endometrium. Journal of Personalized Medicine, 2021, 11, 713.	2.5	8
63	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
64	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
65	Therapeutic Strategies Focused on Cancer-Associated Hypercoagulation for Ovarian Clear Cell Carcinoma. Cancers, 2022, 14, 2125.	3.7	7
66	Hiding in the dark: uncovering cancer drivers through image-guided genomics. Genome Biology, 2014, 15, 563.	8.8	6
67	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
68	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
69	PET/MR imaging for the evaluation of cervical cancer during pregnancy. BMC Pregnancy and Childbirth, 2021, 21, 288.	2.4	6
70	Sentinel node navigation surgery in cervical cancer: a systematic review and metaanalysis. International Journal of Clinical Oncology, 2022, 27, 1247-1255.	2.2	6
71	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
72	Proposing a molecular classification associated with hypercoagulation in ovarian clear cell carcinoma. Gynecologic Oncology, 2021, 163, 327-333.	1.4	5

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73	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
74	Concurrent isolated retroperitoneal HGSC and STIC defined by somatic mutation analysis: a case report. Diagnostic Pathology, 2019, 14, 17.	2.0	4
75	Establishment of in vitro 3D spheroid cell cultivation from human gynecologic cancer tissues. STAR Protocols, 2021, 2, 100354.	1.2	4
76	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
77	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
78	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
79	APOBEC mediated mutagenesis drives genomic heterogeneity in endometriosis. Journal of Human Genetics, 2022, 67, 323-329.	2.3	3
80	Identification of novel exonic mobile element insertions in epithelial ovarian cancers. Human Genome Variation, 2015, 2, 15030.	0.7	2
81	Sleeping Beauty Transposon Mutagenesis Identifies Genes Driving the Initiation and Metastasis of Uterine Leiomyosarcoma. Cancer Research, 2021, 81, 5413-5424.	0.9	2
82	Trends in Pregnancy-Associated Cervical Cancer in Japan between 2012 and 2017: A Multicenter Survey. Cancers, 2022, 14, 3072.	3.7	1
83	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
84	Frequency of homologous recombination associated gene mutations in Japanese patients with ovarian cancer. Gynecologic Oncology, 2021, 162, S163-S164.	1.4	0
85	Abstract 2162: Gene expression profile for predicting survival in ovarian cancer across two independent datasets. , 2010, , .		0
86	Abstract 4795: A novel gene fusion in glioblastoma and a radiation response methylation signature identified by genomic characterization of glioma sphere-forming cells. , 2015, , .		0
87	Abstract 3762: The landscape of therapeutic targetable fusions. , 2015, , .		0
88	Abstract 1504: Identification of novel kinase fusion transcripts in endometrial cancer., 2016,,.		0
89	Abstract 99: Ovarian cancer specific therapeutic vulnerability. , 2016, , .		0
90	Abstract 3383: Exome sequencing in dedifferentiated ovarian mucinous carcinoma., 2017,,.		0

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91	Abstract 411 :In vivopooled shRNA library identifies KPNB1 as a new drug target for epithelial ovarian cancer., $2017,$		O
92	Abstract 529: The significance of activated PI3K/AKT pathway in FGFR3-TACC3 fusion positive cervical cancer. , 2017, , .		0
93	The 61st Annual Meeting of the Japanese Society for Gynecologic Oncology (JSGO). Journal of Gynecologic Oncology, 2019, 30, e114.	2.2	O
94	Population incidence and characteristics of secondary breast cancer after uterine cancer: a competing risk analysis. Archives of Gynecology and Obstetrics, 2022, , 1.	1.7	0
95	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	O