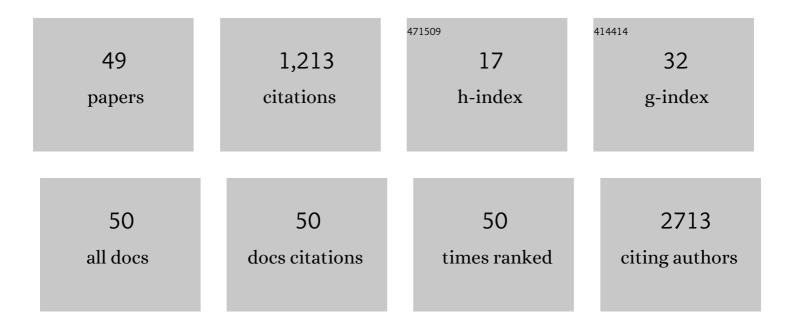
## Shinji Kohsaka

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Recurrent DUX4 fusions in B cell acute lymphoblastic leukemia of adolescents and young adults. Nature Genetics, 2016, 48, 569-574.	21.4	198
2	A method of high-throughput functional evaluation of <i>EGFR</i> gene variants of unknown significance in cancer. Science Translational Medicine, 2017, 9, .	12.4	168
3	Tumor clonality and resistance mechanisms in <i>EGFR</i> mutation-positive non-small-cell lung cancer: implications for therapeutic sequencing. Future Oncology, 2019, 15, 637-652.	2.4	80
4	High-Throughput Functional Evaluation of Variants of Unknown Significance in <i>ERBB2</i> . Clinical Cancer Research, 2018, 24, 5112-5122.	7.0	60
5	DNA mismatch repair deficiency in surgically resected lung adenocarcinoma: Microsatellite instability analysis using the Promega panel. Lung Cancer, 2017, 110, 26-31.	2.0	59
6	Fusion Kinases Identified by Genomic Analyses of Sporadic Microsatellite Instability–High Colorectal Cancers. Clinical Cancer Research, 2019, 25, 378-389.	7.0	49
7	Clinical practice guidance for next-generation sequencing in cancer diagnosis and treatment (edition) Tj ETQq1 1	0,784314 2.2	rgBT /Over
8	Comprehensive assay for the molecular profiling of cancer by target enrichment from formalinâ€fixed paraffinâ€embedded specimens. Cancer Science, 2019, 110, 1464-1479.	3.9	48
9	Integrative analysis of genomic alterations in triple-negative breast cancer in association with homologous recombination deficiency. PLoS Genetics, 2017, 13, e1006853.	3.5	39
10	Prognostic and histogenetic roles of gene alteration and the expression of key potentially actionable targets in salivary duct carcinomas. Oncotarget, 2018, 9, 1852-1867.	1.8	39
11	High-throughput functional evaluation of BRCA2 variants of unknown significance. Nature Communications, 2020, 11, 2573.	12.8	38
12	Small lung tumor biopsy samples are feasible for high quality targeted next generation sequencing. Cancer Science, 2019, 110, 2652-2657.	3.9	37
13	Identification of Novel CD74-NRG2α Fusion From Comprehensive Profiling of Lung Adenocarcinoma in Japanese Never or Light Smokers. Journal of Thoracic Oncology, 2020, 15, 948-961.	1.1	30
14	Highly sensitive fusion detection using plasma cellâ€free RNA in nonâ€smallâ€cell lung cancers. Cancer Science, 2021, 112, 4393-4403.	3.9	21
15	Chorionic Gonadotropin-Î <sup>2</sup> Modulates Epithelial-Mesenchymal Transition in Colorectal Carcinoma Metastasis. American Journal of Pathology, 2018, 188, 204-215.	3.8	19
16	Comprehensive functional evaluation of variants of fibroblast growth factor receptor genes in cancer. Npj Precision Oncology, 2021, 5, 66.	5.4	19
17	Diffuse expression of MUC6 defines a distinct clinicopathological subset of pulmonary invasive mucinous adenocarcinoma. Modern Pathology, 2021, 34, 786-797.	5.5	18
18	IRE1α-XBP1 inhibitors exerted anti-tumor activities in Ewing's sarcoma. Oncotarget, 2018, 9, 14428-14443.	1.8	18

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#	Article	IF	CITATIONS
19	Cabozantinib and dastinib exert anti-tumor activity in alveolar soft part sarcoma. PLoS ONE, 2017, 12, e0185321.	2.5	17
20	Appropriate use of cancer comprehensive genome profiling assay using circulating tumor DNA. Cancer Science, 2021, 112, 3911-3917.	3.9	17
21	Clinicopathological effects of protein phosphatase 2, regulatory subunit A, alpha mutations in gastrointestinal stromal tumors. Modern Pathology, 2016, 29, 1424-1432.	5.5	16
22	Extensive functional evaluation of exon 20 insertion mutations of EGFR. Lung Cancer, 2021, 152, 135-142.	2.0	15
23	Pediatric soft tissue tumor of the upper arm with LMNA-NTRK1 fusion. Human Pathology, 2018, 72, 167-173.	2.0	13
24	Mesenchymal tumours with RREB1–MRTFB fusion involving the mediastinum: extraâ€glossal ectomesenchymal chondromyxoid tumours?. Histopathology, 2020, 76, 1023-1031.	2.9	12
25	Feasibility and clinical utility of comprehensive genomic profiling of hematological malignancies. Cancer Science, 2022, 113, 2763-2777.	3.9	11
26	Expression of F-actin-capping protein subunit beta, CAPZB, is associated with cell growth and motility in epithelioid sarcoma. BMC Cancer, 2016, 16, 206.	2.6	10
27	Detection of circulating sarcoma tumor cells using a microfluidic chip-type cell sorter. Scientific Reports, 2019, 9, 20047.	3.3	10
28	Comprehensive molecular and clinicopathological profiling of desmoid tumours. European Journal of Cancer, 2021, 145, 109-120.	2.8	10
29	Assessment of Predictive Biomarkers of the Response to Pazopanib Based on an Integrative Analysis of High-grade Soft-tissue Sarcomas: Analysis of a Tumor Sample from a Responder and Patients with Other Soft-tissue Sarcomas. Clinical Orthopaedics and Related Research, 2020, 478, 2461-2476.	1.5	9
30	Comprehensive clinicopathological and molecular analysis of primary malignant melanoma of the oesophagus. Histopathology, 2021, 78, 240-251.	2.9	9
31	Genomic profiling of multiple primary cancers including synchronous lung adenocarcinoma and bilateral malignant mesotheliomas: Identification of a novel <i>BAP1</i> germline variant. Pathology International, 2020, 70, 775-780.	1.3	8
32	LSD1/KDM1 isoform LSD1+8a contributes to neural differentiation in small cell lung cancer. Biochemistry and Biophysics Reports, 2017, 9, 86-94.	1.3	7
33	Clinicopathological characteristics of lung adenocarcinoma with compound EGFR mutations. Human Pathology, 2020, 103, 42-51.	2.0	7
34	Development of an optimal protocol for molecular profiling of tumor cells in pleural effusions at singleâ€cell level. Cancer Science, 2021, 112, 2006-2019.	3.9	7
35	<i>PPP2R1A</i> regulated by PAX3/FOXO1 fusion contributes to the acquisition of aggressive behavior in PAX3/FOXO1-positive alveolar rhabdomyosarcoma. Oncotarget, 2018, 9, 25206-25215.	1.8	7
36	Preclinical assessment of combination therapy of EGFR tyrosine kinase inhibitors in a highly heterogeneous tumor model. Oncogene, 2022, 41, 2470-2479.	5.9	7

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#	Article	IF	CITATIONS
37	Comprehensive molecular profiling of pulmonary pleomorphic carcinoma. Npj Precision Oncology, 2021, 5, 57.	5.4	6
38	Spinal solitary fibrous tumor of the neck: Next-generation sequencing-based analysis of genomic aberrations. Auris Nasus Larynx, 2020, 47, 1058-1063.	1.2	5
39	Tumor budding and human chorionic gonadotropin-β expression correlate with unfavorable patient outcome in colorectal carcinoma. Medical Oncology, 2018, 35, 104.	2.5	4
40	Transformation from EGFR/PTEN coâ€mutated lung adenocarcinoma to small cell carcinoma in lymph node metastasis. Pathology International, 2020, 70, 295-299.	1.3	4
41	Histological characteristics of lung adenocarcinoma with uncommon actionable alterations: special emphasis on <i>MET</i> exon 14 skipping alterations. Histopathology, 2021, 78, 987-999.	2.9	3
42	Proteomic signatures corresponding to the SS18/SSX fusion gene in synovial sarcoma. Oncotarget, 2018, 9, 37509-37519.	1.8	2
43	Scapular Angiomatoid Fibrous Histiocytoma with EWSR1-CREB1 Fusion in an Adult Patient. Case Reports in Orthopedics, 2021, 2021, 1-6.	0.3	2
44	MicroSEC filters sequence errors for formalin-fixed and paraffin-embedded samples. Communications Biology, 2021, 4, 1396.	4.4	2
45	Genetic Testing Distinguishes Multiple Chondroid Chordomas with Neuraxial Bone Metastases from Multicentric Tumors. Case Reports in Genetics, 2020, 2020, 1-6.	0.2	1
46	Transcription start siteâ€level expression of thyroid transcription factor 1 isoforms in lung adenocarcinoma and its clinicopathological significance. Journal of Pathology: Clinical Research, 2021, 7, 361-374.	3.0	1
47	Genetic diagnosis of pseudomyxoma peritonei originating from mucinous borderline tumor inside an ovarian teratoma. BMC Medical Genomics, 2022, 15, 51.	1.5	1
48	OUP accepted manuscript. Japanese Journal of Clinical Oncology, 2022, , .	1.3	1
49	Cancer Genomic Profiling of Gynecological Malignancies by Todai OncoPanel, a Twin DNA and RNA Panel. Current Human Cell Research and Applications, 2021, , 27-39.	0.1	0