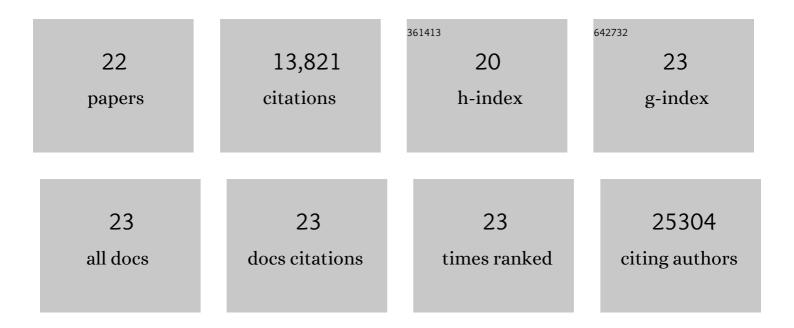
## Toshinori Hinoue

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11716076/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Molecular Features of Cancers Exhibiting Exceptional Responses to Treatment. Cancer Cell, 2021, 39, 38-53.e7.	16.8	65
2	Phase I trial of TRC102 (methoxyamine HCl) in combination with temozolomide in patients with relapsed solid tumors and lymphomas. Oncotarget, 2020, 11, 3959-3971.	1.8	8
3	An Integrated TCGA Pan-Cancer Clinical Data Resource to Drive High-Quality Survival Outcome Analytics. Cell, 2018, 173, 400-416.e11.	28.9	2,277
4	Cell-of-Origin Patterns Dominate the Molecular Classification of 10,000 Tumors from 33 Types of Cancer. Cell, 2018, 173, 291-304.e6.	28.9	1,718
5	Perspective on Oncogenic Processes at the End of the Beginning of Cancer Genomics. Cell, 2018, 173, 305-320.e10.	28.9	272
6	Somatic Mutational Landscape of Splicing Factor Genes and Their Functional Consequences across 33 Cancer Types. Cell Reports, 2018, 23, 282-296.e4.	6.4	333
7	Genomic and Molecular Landscape of DNA Damage Repair Deficiency across The Cancer Genome Atlas. Cell Reports, 2018, 23, 239-254.e6.	6.4	801
8	Genomic and Functional Approaches to Understanding Cancer Aneuploidy. Cancer Cell, 2018, 33, 676-689.e3.	16.8	750
9	Comparative Molecular Analysis of Gastrointestinal Adenocarcinomas. Cancer Cell, 2018, 33, 721-735.e8.	16.8	396
10	The chromatin accessibility landscape of primary human cancers. Science, 2018, 362, .	12.6	781
11	Integrative Epigenetic Analysis Reveals Therapeutic Targets to the DNA Methyltransferase Inhibitor Guadecitabine (SGIâ€110) in Hepatocellular Carcinoma. Hepatology, 2018, 68, 1412-1428.	7.3	48
12	Comprehensive and Integrative Genomic Characterization of Hepatocellular Carcinoma. Cell, 2017, 169, 1327-1341.e23.	28.9	1,794
13	Integrative Genomic Analysis of Cholangiocarcinoma Identifies Distinct IDH-Mutant Molecular Profiles. Cell Reports, 2017, 18, 2780-2794.	6.4	416
14	Effects of folylpolyglutamate synthase modulation on global and gene-specific DNA methylation and gene expression in human colon and breast cancer cells. Journal of Nutritional Biochemistry, 2016, 29, 27-35.	4.2	7
15	Î <sup>3</sup> -Glutamyl hydrolase modulation significantly influences global and gene-specific DNA methylation and gene expression in human colon and breast cancer cells. Genes and Nutrition, 2015, 10, 444.	2.5	10
16	Comprehensive, Integrative Genomic Analysis of Diffuse Lower-Grade Gliomas. New England Journal of Medicine, 2015, 372, 2481-2498.	27.0	2,582
17	Genome-scale analysis of aberrant DNA methylation in colorectal cancer. Genome Research, 2012, 22, 271-282.	5.5	527
18	Regions of focal DNA hypermethylation and long-range hypomethylation in colorectal cancer coincide with nuclear lamina–associated domains. Nature Genetics, 2012, 44, 40-46.	21.4	588

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#	Article	IF	CITATIONS
19	Genome-Scale Discovery of DNA-Methylation Biomarkers for Blood-Based Detection of Colorectal Cancer. PLoS ONE, 2012, 7, e50266.	2.5	103
20	Analysis of the Association between CIMP and BRAFV600E in Colorectal Cancer by DNA Methylation Profiling. PLoS ONE, 2009, 4, e8357.	2.5	133
21	PACAP mediates the neural proliferative pathway of Mastomys Enterochromaffin-like cell transformation. Regulatory Peptides, 2001, 102, 157-164.	1.9	22
22	Pituitary adenylate cyclase-activating polypeptide modulates gastric enterochromaffin-like cell proliferation in rats. Gastroenterology, 1999, 116, 623-635.	1.3	40