

# Toshio F Yoshimatsu

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8461657/publications.pdf>

Version: 2024-02-01

12  
papers

654  
citations

933447

10  
h-index

1199594

12  
g-index

14  
all docs

14  
docs citations

14  
times ranked

1390  
citing authors

#	ARTICLE	IF	CITATIONS
1	Racial disparities in survival outcomes among breast cancer patients by molecular subtypes. Breast Cancer Research and Treatment, 2021, 185, 841-849.	2.5	25
2	The <i>BRCA1</i> Pseudogene Negatively Regulates Antitumor Responses through Inhibition of Innate Immune Defense Mechanisms. Cancer Research, 2021, 81, 1540-1551.	0.9	6
3	Abstract PS18-12: Comparative analysis of differential gene expression by ancestry using primary breast cancers from Nigeria and the cancer genome atlas (TCGA). , 2021, , .		0
4	Whole-genome analysis of Nigerian patients with breast cancer reveals ethnic-driven somatic evolution and distinct genomic subtypes. Nature Communications, 2021, 12, 6946.	12.8	22
5	Prevalence of Inherited Mutations in Breast Cancer Predisposition Genes among Women in Uganda and Cameroon. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 359-367.	2.5	36
6	Germline variants and somatic mutation signatures of breast cancer across populations of African and European ancestry in the US and Nigeria. International Journal of Cancer, 2019, 145, 3321-3333.	5.1	16
7	Intensive Surveillance with Biannual Dynamic Contrast-Enhanced Magnetic Resonance Imaging Downstages Breast Cancer in <i>BRCA1</i> Mutation Carriers. Clinical Cancer Research, 2019, 25, 1786-1794.	7.0	44
8	Inherited Breast Cancer in Nigerian Women. Journal of Clinical Oncology, 2018, 36, 2820-2825.	1.6	80
9	Characterization of Nigerian breast cancer reveals prevalent homologous recombination deficiency and aggressive molecular features. Nature Communications, 2018, 9, 4181.	12.8	77
10	Comparison of Breast Cancer Molecular Features and Survival by African and European Ancestry in The Cancer Genome Atlas. JAMA Oncology, 2017, 3, 1654.	7.1	208
11	Identification of a circulating MicroRNA signature to distinguish recurrence in breast cancer patients. Oncotarget, 2016, 7, 55231-55248.	1.8	70
12	Naturally occurring <i>BRCA2</i> alternative mRNA splicing events in clinically relevant samples. Journal of Medical Genetics, 2016, 53, 548-558.	3.2	69