

# Yoshitsugu Mitani

## List of Publications by Year in Descending Order

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**Version:** 2024-04-28

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

44  
papers

2,403  
citations

24  
h-index

46  
g-index

46  
ext. papers

2,700  
ext. citations

7.4  
avg, IF

4.1  
L-index

#	Paper	IF	Citations
44	LMW cyclin E and its novel catalytic partner CDK5 are therapeutic targets and prognostic biomarkers in salivary gland cancers. <i>Oncogenesis</i> , <b>2021</b> , 10, 40	6.6	
43	Whole-Genome Sequencing of Common Salivary Gland Carcinomas: Subtype-Restricted and Shared Genetic Alterations. <i>Clinical Cancer Research</i> , <b>2021</b> , 27, 3960-3969	12.9	2
42	Proteogenomic Analysis of Salivary Adenoid Cystic Carcinomas Defines Molecular Subtypes and Identifies Therapeutic Targets. <i>Clinical Cancer Research</i> , <b>2021</b> , 27, 852-864	12.9	12
41	Reciprocal and Autonomous Glucocorticoid and Androgen Receptor Activation in Salivary Duct Carcinoma. <i>Clinical Cancer Research</i> , <b>2020</b> , 26, 1175-1184	12.9	5
40	Oncogenic Orphan Nuclear Receptor NR4A3 Interacts and Cooperates with MYB in Acinic Cell Carcinoma. <i>Cancers</i> , <b>2020</b> , 12,	6.6	3
39	N-Terminal Truncated Myb with New Transcriptional Activity Produced Through Use of an Alternative MYB Promoter in Salivary Gland Adenoid Cystic Carcinoma. <i>Cancers</i> , <b>2019</b> , 12,	6.6	3
38	Frequent and differential mutations of the CYLD gene in basal cell salivary neoplasms: linkage to tumor development and progression. <i>Modern Pathology</i> , <b>2018</b> , 31, 1064-1072	9.8	20
37	Transcriptomes define distinct subgroups of salivary gland adenoid cystic carcinoma with different driver mutations and outcomes. <i>Oncotarget</i> , <b>2018</b> , 9, 7341-7358	3.3	21
36	Frequent PTEN loss and differential HER2/PI3K signaling pathway alterations in salivary duct carcinoma: Implications for targeted therapy. <i>Cancer</i> , <b>2018</b> , 124, 3693-3705	6.4	14
35	Chromosomal abnormalities and molecular landscape of metastasizing mucinous salivary adenocarcinoma. <i>Oral Oncology</i> , <b>2017</b> , 66, 38-45	4.4	7
34	Establishment and genomic characterization of primary salivary duct carcinoma cell line. <i>Oral Oncology</i> , <b>2017</b> , 69, 108-114	4.4	4
33	Activating NOTCH1 Mutations Define a Distinct Subgroup of Patients With Adenoid Cystic Carcinoma Who Have Poor Prognosis, Propensity to Bone and Liver Metastasis, and Potential Responsiveness to Notch1 Inhibitors. <i>Journal of Clinical Oncology</i> , <b>2017</b> , 35, 352-360	2.2	104
32	Spatio-Temporal Genomic Heterogeneity, Phylogeny, and Metastatic Evolution in Salivary Adenoid Cystic Carcinoma. <i>Journal of the National Cancer Institute</i> , <b>2017</b> , 109,	9.7	17
31	Novel MYBL1 Gene Rearrangements with Recurrent MYBL1-NFIB Fusions in Salivary Adenoid Cystic Carcinomas Lacking t(6;9) Translocations. <i>Clinical Cancer Research</i> , <b>2016</b> , 22, 725-33	12.9	116
30	Notch1 mutations to define a subgroup of adenoid cystic carcinoma (ACC): Tumor stage, propensity to bone and liver metastasis, risk of relapse, and overall survival.. <i>Journal of Clinical Oncology</i> , <b>2015</b> , 33, 6081-6081	2.2	3
29	Alterations associated with androgen receptor gene activation in salivary duct carcinoma of both sexes: potential therapeutic ramifications. <i>Clinical Cancer Research</i> , <b>2014</b> , 20, 6570-81	12.9	55
28	A unifying gene signature for adenoid cystic cancer identifies parallel MYB-dependent and MYB-independent therapeutic targets. <i>Oncotarget</i> , <b>2014</b> , 5, 12528-42	3.3	31

27	Detailed genome-wide SNP analysis of major salivary carcinomas localizes subtype-specific chromosome sites and oncogenes of potential clinical significance. <i>American Journal of Pathology</i> , <b>2013</b> , 182, 2048-57	5.8	21
26	Whole exome sequencing of adenoid cystic carcinoma. <i>Journal of Clinical Investigation</i> , <b>2013</b> , 123, 2965-8	5.9	188
25	MicroRNA profiling of salivary adenoid cystic carcinoma: association of miR-17-92 upregulation with poor outcome. <i>PLoS ONE</i> , <b>2013</b> , 8, e66778	3.7	33
24	Expression and regulation of the $\beta$ and TAp63 isoforms in salivary gland tumorigenesis clinical and experimental findings. <i>American Journal of Pathology</i> , <b>2011</b> , 179, 391-9	5.8	16
23	Myxolipoma in the tongue - a clinical case report and review of the literature. <i>Head &amp; Neck Oncology</i> , <b>2011</b> , 3, 50		9
22	Novel chromosomal rearrangements and break points at the t(6;9) in salivary adenoid cystic carcinoma: association with MYB-NFIB chimeric fusion, MYB expression, and clinical outcome. <i>Clinical Cancer Research</i> , <b>2011</b> , 17, 7003-14	12.9	110
21	Comprehensive analysis of the MYB-NFIB gene fusion in salivary adenoid cystic carcinoma: Incidence, variability, and clinicopathologic significance. <i>Clinical Cancer Research</i> , <b>2010</b> , 16, 4722-31	12.9	204
20	DeltaNp63alpha-dependent expression of Id-3 distinctively suppresses the invasiveness of human squamous cell carcinoma. <i>International Journal of Cancer</i> , <b>2009</b> , 124, 2837-44	7.5	36
19	Increased Expression of CENP-H Gene in Human Salivary Gland Carcinomas. <i>Oral Science International</i> , <b>2008</b> , 5, 43-51	0.5	1
18	Expression of epiregulin, a novel epidermal growth factor ligand associated with prognosis in human oral squamous cell carcinomas. <i>Oncology Reports</i> , <b>2008</b> ,	3.5	1
17	DNA demethylation of vascular endothelial growth factor-C is associated with gene expression and its possible involvement of lymphangiogenesis in gastric cancer. <i>International Journal of Cancer</i> , <b>2007</b> , 120, 1689-95	7.5	20
16	Reg IV is a serum biomarker for gastric cancer patients and predicts response to 5-fluorouracil-based chemotherapy. <i>Oncogene</i> , <b>2007</b> , 26, 4383-93	9.2	96
15	UCN-01 (7-hydroxystaurosporine) induces apoptosis and G1 arrest of both primary and metastatic oral cancer cell lines in vitro. <i>Oral Surgery Oral Medicine Oral Pathology Oral Radiology and Endodontics</i> , <b>2007</b> , 103, 391-7		12
14	Accumulation of DNA methylation is associated with tumor stage in gastric cancer. <i>Cancer</i> , <b>2006</b> , 106, 1250-9	6.4	111
13	Down-regulation of the claudin-18 gene, identified through serial analysis of gene expression data analysis, in gastric cancer with an intestinal phenotype. <i>Journal of Pathology</i> , <b>2006</b> , 208, 633-42	9.4	104
12	Systematic search for gastric cancer-specific genes based on SAGE data: melanoma inhibitory activity and matrix metalloproteinase-10 are novel prognostic factors in patients with gastric cancer. <i>Oncogene</i> , <b>2006</b> , 25, 2546-57	9.2	128
11	Differential expression of claudin-2 in normal human tissues and gastrointestinal carcinomas. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , <b>2006</b> , 448, 428-34	5.1	54
10	Loss of heterozygosity and histone hypoacetylation of the PINX1 gene are associated with reduced expression in gastric carcinoma. <i>Oncogene</i> , <b>2005</b> , 24, 157-64	9.2	33

9	Histone H3 acetylation is associated with reduced p21(WAF1/CIP1) expression by gastric carcinoma. <i>Journal of Pathology</i> , <b>2005</b> , 205, 65-73	9.4	60
8	Expression and localization of Reg IV in human neoplastic and non-neoplastic tissues: Reg IV expression is associated with intestinal and neuroendocrine differentiation in gastric adenocarcinoma. <i>Journal of Pathology</i> , <b>2005</b> , 207, 185-98	9.4	101
7	Genes involved in invasion and metastasis of gastric cancer identified by array-based hybridization and serial analysis of gene expression. <i>Oncology</i> , <b>2005</b> , 69 Suppl 1, 17-22	3.6	46
6	Frequent loss of RUNX3 expression by promoter hypermethylation in gastric carcinoma. <i>Pathobiology</i> , <b>2004</b> , 71, 137-43	3.6	63
5	Gene expression profile of gastric carcinoma: identification of genes and tags potentially involved in invasion, metastasis, and carcinogenesis by serial analysis of gene expression. <i>Cancer Research</i> , <b>2004</b> , 64, 2397-405	10.1	241
4	Expression of POT1 is associated with tumor stage and telomere length in gastric carcinoma. <i>Cancer Research</i> , <b>2004</b> , 64, 523-9	10.1	97
3	Frequent epigenetic inactivation of RIZ1 by promoter hypermethylation in human gastric carcinoma. <i>International Journal of Cancer</i> , <b>2004</b> , 110, 212-8	7.5	51
2	DNA hypermethylation and histone hypoacetylation of the HLTF gene are associated with reduced expression in gastric carcinoma. <i>Cancer Science</i> , <b>2003</b> , 94, 692-8	6.9	34
1	Histone acetylation and gastrointestinal carcinogenesis. <i>Annals of the New York Academy of Sciences</i> , <b>2003</b> , 983, 220-31	6.5	116