

Mikhail Tavobilov

List of Publications by Year in descending order

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38
papers

24,288
citations

126907

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times ranked

33577
citing authors

#	ARTICLE	IF	CITATIONS
1	The first clinical experience with innovative HOT AXIOS stent in a patient with parapancreatic fluid collection. Eksperimental'naya I Klinicheskaya Gastroenterologiya, 2021, , 114-119.	0.4	0
2	Acinar cell cystadenoma of the pancreatic head. Annals of HPB Surgery, 2020, 25, 145-151.	0.5	1
3	Combination of minimally invasive approaches of the debridement in treatment of the patient with infected necrotizing pancreatitis. Eksperimental'naya I Klinicheskaya Gastroenterologiya, 2020, , 119-123.	0.4	0
4	THE CHOICE OF THE METHOD OF ENDOSCOPIC STENTING IN PATIENTS WITH UNRESECTABLE ADENOCARCINOMA OF THE PANCREATIC HEAD. Siberian Journal of Oncology, 2020, 19, 116-121.	0.3	0
5	SOLID PSEUDOPAPILLARY TUMOR OF THE PANCREAS IN A MAN: A CASE REPORT. Siberian Journal of Oncology, 2019, 18, 113-117.	0.3	1
6	EXPERIENCE OF DOUBLE STENTING IN MALIGNANT BILIARY AND DUODENAL OBSTRUCTION IN PATIENTS WITH ADENOCARCINOMA OF THE PANCREATIC HEAD. Voprosy Onkologii, 2019, 65, 404-408.	0.2	0
7	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
8	Comprehensive Characterization of Cancer Driver Genes and Mutations. Cell, 2018, 173, 371-385.e18.	28.9	1,670
9	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
10	A Pan-Cancer Analysis of Enhancer Expression in Nearly 9000 Patient Samples. Cell, 2018, 173, 386-399.e12.	28.9	228
11	Perspective on Oncogenic Processes at the End of the Beginning of Cancer Genomics. Cell, 2018, 173, 305-320.e10.	28.9	272
12	Machine Learning Identifies Stemness Features Associated with Oncogenic Dedifferentiation. Cell, 2018, 173, 338-354.e15.	28.9	1,417
13	Oncogenic Signaling Pathways in The Cancer Genome Atlas. Cell, 2018, 173, 321-337.e10.	28.9	2,111
14	Pathogenic Germline Variants in 10,389 Adult Cancers. Cell, 2018, 173, 355-370.e14.	28.9	620
15	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
16	Driver Fusions and Their Implications in the Development and Treatment of Human Cancers. Cell Reports, 2018, 23, 227-238.e3.	6.4	407
17	Genomic, Pathway Network, and Immunologic Features Distinguishing Squamous Carcinomas. Cell Reports, 2018, 23, 194-212.e6.	6.4	245
18	Pan-Cancer Analysis of lncRNA Regulation Supports Their Targeting of Cancer Genes in Each Tumor Context. Cell Reports, 2018, 23, 297-312.e12.	6.4	205

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19	The Cancer Genome Atlas Comprehensive Molecular Characterization of Renal Cell Carcinoma. Cell Reports, 2018, 23, 313-326.e5.	6.4	523
20	Spatial Organization and Molecular Correlation of Tumor-Infiltrating Lymphocytes Using Deep Learning on Pathology Images. Cell Reports, 2018, 23, 181-193.e7.	6.4	683
21	The Immune Landscape of Cancer. Immunity, 2018, 48, 812-830.e14.	14.3	3,706
22	Machine Learning Detects Pan-cancer Ras Pathway Activation in The Cancer Genome Atlas. Cell Reports, 2018, 23, 172-180.e3.	6.4	119
23	Integrated Genomic Analysis of the Ubiquitin Pathway across Cancer Types. Cell Reports, 2018, 23, 213-226.e3.	6.4	83
24	Genomic and Molecular Landscape of DNA Damage Repair Deficiency across The Cancer Genome Atlas. Cell Reports, 2018, 23, 239-254.e6.	6.4	801
25	Molecular Characterization and Clinical Relevance of Metabolic Expression Subtypes in Human Cancers. Cell Reports, 2018, 23, 255-269.e4.	6.4	204
26	Systematic Analysis of Splice-Site-Creating Mutations in Cancer. Cell Reports, 2018, 23, 270-281.e3.	6.4	177
27	Scalable Open Science Approach for Mutation Calling of Tumor Exomes Using Multiple Genomic Pipelines. Cell Systems, 2018, 6, 271-281.e7.	6.2	605
28	Pan-cancer Alterations of the MYC Oncogene and Its Proximal Network across the Cancer Genome Atlas. Cell Systems, 2018, 6, 282-300.e2.	6.2	284
29	lncRNA Epigenetic Landscape Analysis Identifies EPIC1 as an Oncogenic lncRNA that Interacts with MYC and Promotes Cell-Cycle Progression in Cancer. Cancer Cell, 2018, 33, 706-720.e9.	16.8	400
30	Genomic and Functional Approaches to Understanding Cancer Aneuploidy. Cancer Cell, 2018, 33, 676-689.e3.	16.8	750
31	Comparative Molecular Analysis of Gastrointestinal Adenocarcinomas. Cancer Cell, 2018, 33, 721-735.e8.	16.8	396
32	A Comprehensive Pan-Cancer Molecular Study of Gynecologic and Breast Cancers. Cancer Cell, 2018, 33, 690-705.e9.	16.8	478
33	A Pan-Cancer Analysis Reveals High-Frequency Genetic Alterations in Mediators of Signaling by the TGF- β Superfamily. Cell Systems, 2018, 7, 422-437.e7.	6.2	134
34	Comprehensive Molecular Characterization of the Hippo Signaling Pathway in Cancer. Cell Reports, 2018, 25, 1304-1317.e5.	6.4	329
35	Comprehensive Analysis of Alternative Splicing Across Tumors from 8,705 Patients. Cancer Cell, 2018, 34, 211-224.e6.	16.8	623
36	COMBINED MODALITY TREATMENT FOR PATIENTS WITH INOPERABLE COLORECTAL LIVER METASTASES. Siberian Journal of Oncology, 2018, 17, 34-40.	0.3	4

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37	Integrated Genomic Characterization of Pancreatic Ductal Adenocarcinoma. <i>Cancer Cell</i> , 2017, 32, 185-203.e13.	16.8	1,428
38	ENDOSCOPIC ULTRASOUND-GUIDED RESECTION OF SOMATOSTATINOMA OF THE AMPULLA OF THE MAJOR DUODENAL PAPILLA IN THE PATIENT WITH INHERITED NEUROFIBROMATOSIS. <i>Siberian Journal of Oncology</i> , 2017, 16, 98-103.	0.3	0