

# Ronak H Shah Bs

## List of Publications by Year in Descending Order

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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.

23  
papers

5,434  
citations

20  
h-index

30  
g-index

30  
ext. papers

6,849  
ext. citations

14.8  
avg, IF

4.43  
L-index

#	Paper	IF	Citations
23	Novel insights into breast cancer copy number genetic heterogeneity revealed by single-cell genome sequencing. <i>ELife</i> , <b>2020</b> , 9,	8.9	20
22	Tracking tumour evolution in glioma through liquid biopsies of cerebrospinal fluid. <i>Nature</i> , <b>2019</b> , 565, 654-658	50.4	214
21	Precision oncology: lessons learned and challenges for the future. <i>Cancer Management and Research</i> , <b>2019</b> , 11, 7525-7536	3.6	4
20	HMGB1 Causes Anemia of Inflammation By Modulating Erythropoietin Signal Transduction. <i>Blood</i> , <b>2018</b> , 132, 628-628	2.2	
19	Real-Time Genomic Characterization of Metastatic Pancreatic Neuroendocrine Tumors Has Prognostic Implications and Identifies Potential Germline Actionability. <i>JCO Precision Oncology</i> , <b>2018</b> , 2018,	3.6	21
18	Mutational landscape of metastatic cancer revealed from prospective clinical sequencing of 10,000 patients. <i>Nature Medicine</i> , <b>2017</b> , 23, 703-713	50.5	1638
17	Mutation Detection in Patients With Advanced Cancer by Universal Sequencing of Cancer-Related Genes in Tumor and Normal DNA vs Guideline-Based Germline Testing. <i>JAMA - Journal of the American Medical Association</i> , <b>2017</b> , 318, 825-835	27.4	235
16	Clinical and genetic determinants of ovarian metastases from colorectal cancer. <i>Cancer</i> , <b>2017</b> , 123, 11346-1143	14.3	30
15	Local recurrences at the anastomotic area are clonally related to the primary tumor in sporadic colorectal carcinoma. <i>Oncotarget</i> , <b>2017</b> , 8, 42487-42494	3.3	6
14	Next-generation sequencing (NGS) in pancreatic neuroendocrine tumors (panNETs): Defining differentiation and grade genetically.. <i>Journal of Clinical Oncology</i> , <b>2017</b> , 35, 291-291	2.2	
13	The oncocytic subtype is genetically distinct from other pancreatic intraductal papillary mucinous neoplasm subtypes. <i>Modern Pathology</i> , <b>2016</b> , 29, 1058-69	9.8	54
12	Genomic and transcriptomic hallmarks of poorly differentiated and anaplastic thyroid cancers. <i>Journal of Clinical Investigation</i> , <b>2016</b> , 126, 1052-66	15.9	576
11	Evaluating Cancer of the Central Nervous System Through Next-Generation Sequencing of Cerebrospinal Fluid. <i>Journal of Clinical Oncology</i> , <b>2016</b> , 34, 2404-15	2.2	208
10	Sequencing of 279 cancer genes in ampullary carcinoma reveals trends relating to histologic subtypes and frequent amplification and overexpression of ERBB2 (HER2). <i>Modern Pathology</i> , <b>2015</b> , 28, 1123-9	9.8	37
9	Memorial Sloan Kettering-Integrated Mutation Profiling of Actionable Cancer Targets (MSK-IMPACT): A Hybridization Capture-Based Next-Generation Sequencing Clinical Assay for Solid Tumor Molecular Oncology. <i>Journal of Molecular Diagnostics</i> , <b>2015</b> , 17, 251-64	5.1	1066
8	Alternative transcription initiation leads to expression of a novel ALK isoform in cancer. <i>Nature</i> , <b>2015</b> , 526, 453-7	50.4	144
7	Genomic Characterization of Upper Tract Urothelial Carcinoma. <i>European Urology</i> , <b>2015</b> , 68, 970-7	10.2	147

6	Convergent loss of PTEN leads to clinical resistance to a PI(3)K inhibitor. <i>Nature</i> , <b>2015</b> , 518, 240-4	50.4	366
5	Comprehensive mutation profiling by next-generation sequencing of effusion fluids from patients with high-grade serous ovarian carcinoma. <i>Cancer Cytopathology</i> , <b>2015</b> , 123, 289-97	3.9	21
4	Genomic predictors of survival in patients with high-grade urothelial carcinoma of the bladder. <i>European Urology</i> , <b>2015</b> , 67, 198-201	10.2	105
3	Comparative sequencing analysis reveals high genomic concordance between matched primary and metastatic colorectal cancer lesions. <i>Genome Biology</i> , <b>2014</b> , 15, 454	18.3	224
2	Recurrent SMARCA4 mutations in small cell carcinoma of the ovary. <i>Nature Genetics</i> , <b>2014</b> , 46, 424-6	36.3	231
1	Detecting somatic genetic alterations in tumor specimens by exon capture and massively parallel sequencing. <i>Journal of Visualized Experiments</i> , <b>2013</b> , e50710	1.6	53