

# Peter S Choi

## List of Publications by Year in Descending Order

**Source:** <https://exaly.com/author-pdf/3745907/peter-s-choi-publications-by-year.pdf>

**Version:** 2024-04-23

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

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

17  
papers

1,512  
citations

12  
h-index

20  
g-index

20  
ext. papers

2,012  
ext. citations

17.2  
avg, IF

4.5  
L-index

#	Paper	IF	Citations
17	A genome-scale CRISPR screen reveals PRMT1 as a critical regulator of androgen receptor signaling in prostate cancer.. <i>Cell Reports</i> , <b>2022</b> , 38, 110417	10.6	0
16	RNA-binding proteins of COSMIC importance in cancer. <i>Journal of Clinical Investigation</i> , <b>2021</b> , 131,	15.9	2
15	A predominant enhancer co-amplified with the SOX2 oncogene is necessary and sufficient for its expression in squamous cancer. <i>Nature Communications</i> , <b>2021</b> , 12, 7139	17.4	3
14	Multi-Omics Analysis Identifies MGA as a Negative Regulator of the MYC Pathway in Lung Adenocarcinoma. <i>Molecular Cancer Research</i> , <b>2020</b> , 18, 574-584	6.6	13
13	Chromatin Looping Shapes KLF5-Dependent Transcriptional Programs in Human Epithelial Cancers. <i>Cancer Research</i> , <b>2020</b> , 80, 5464-5477	10.1	7
12	Identification and Characterization of Oncogenic Mutations in Lung Adenocarcinoma. <i>Molecular Cancer Research</i> , <b>2019</b> , 17, 1002-1012	6.6	18
11	Somatic Superenhancer Duplications and Hotspot Mutations Lead to Oncogenic Activation of the KLF5 Transcription Factor. <i>Cancer Discovery</i> , <b>2018</b> , 8, 108-125	24.4	67
10	An alternative splicing switch in FLNB promotes the mesenchymal cell state in human breast cancer. <i>ELife</i> , <b>2018</b> , 7,	8.9	47
9	Identification of ADAR1 adenosine deaminase dependency in a subset of cancer cells. <i>Nature Communications</i> , <b>2018</b> , 9, 5450	17.4	83
8	Genome-scale analysis identifies paralog lethality as a vulnerability of chromosome 1p loss in cancer. <i>Nature Genetics</i> , <b>2018</b> , 50, 937-943	36.3	35
7	Human Kidney Tubule-Specific Gene Expression Based Dissection of Chronic Kidney Disease Traits. <i>EBioMedicine</i> , <b>2017</b> , 24, 267-276	8.8	43
6	Identification of focally amplified lineage-specific super-enhancers in human epithelial cancers. <i>Nature Genetics</i> , <b>2016</b> , 48, 176-82	36.3	210
5	Defective fatty acid oxidation in renal tubular epithelial cells has a key role in kidney fibrosis development. <i>Nature Medicine</i> , <b>2015</b> , 21, 37-46	50.5	628
4	Functional genomic annotation of genetic risk loci highlights inflammation and epithelial biology networks in CKD. <i>Journal of the American Society of Nephrology: JASN</i> , <b>2015</b> , 26, 692-714	12.7	39
3	Targeted genomic rearrangements using CRISPR/Cas technology. <i>Nature Communications</i> , <b>2014</b> , 5, 3728	17.4	203
2	A pan-cancer analysis of transcriptome changes associated with somatic mutations in U2AF1 reveals commonly altered splicing events. <i>PLoS ONE</i> , <b>2014</b> , 9, e87361	3.7	112
1	A genome-scale CRISPR screen reveals PRMT1 as a critical regulator of androgen receptor signaling in prostate cancer		1

