## **Zhong Sheng Sun**

## List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/8134824/zhong-sheng-sun-publications-by-year.pdf

Version: 2024-04-25

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.

7	243	6	9
papers	citations	h-index	g-index
9	403 ext. citations	13.9	3.07
ext. papers		avg, IF	L-index

#	Paper	IF	Citations
7	OncoVar: an integrated database and analysis platform for oncogenic driver variants in cancers.  Nucleic Acids Research, 2021, 49, D1289-D1301	20.1	18
6	Low intratumor heterogeneity correlates with increased response to PD-1 blockade in renal cell carcinoma. <i>Therapeutic Advances in Medical Oncology</i> , <b>2020</b> , 12, 1758835920977117	5.4	6
5	Al-Driver: an ensemble method for identifying driver mutations in personal cancer genomes. <i>NAR Genomics and Bioinformatics</i> , <b>2020</b> , 2, lqaa084	3.7	3
4	OncoBase: a platform for decoding regulatory somatic mutations in human cancers. <i>Nucleic Acids Research</i> , <b>2019</b> , 47, D1044-D1055	20.1	18
3	VarCards: an integrated genetic and clinical database for coding variants in the human genome. <i>Nucleic Acids Research</i> , <b>2018</b> , 46, D1039-D1048	20.1	86
2	EpiDenovo: a platform for linking regulatory de novo mutations to developmental epigenetics and diseases. <i>Nucleic Acids Research</i> , <b>2018</b> , 46, D92-D99	20.1	11
1	Performance evaluation of pathogenicity-computation methods for missense variants. <i>Nucleic Acids Research</i> , <b>2018</b> , 46, 7793-7804	20.1	100