

# Chong Sun

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1920400/publications.pdf>

Version: 2024-02-01

13  
papers

5,771  
citations

759233

12  
h-index

1125743

13  
g-index

14  
all docs

14  
docs citations

14  
times ranked

11148  
citing authors

#	ARTICLE	IF	CITATIONS
1	Allelic expression imbalance of PIK3CA mutations is frequent in breast cancer and prognostically significant. <i>Npj Breast Cancer</i> , 2022, 8, .	5.2	1
2	EGFR activation limits the response of liver cancer to lenvatinib. <i>Nature</i> , 2021, 595, 730-734.	27.8	183
3	Regulation and Function of the PD-L1 Checkpoint. <i>Immunity</i> , 2018, 48, 434-452.	14.3	1,437
4	Identification of CMTM6 and CMTM4 as PD-L1 protein regulators. <i>Nature</i> , 2017, 549, 106-110.	27.8	501
5	BRAF V600E Kinase Domain Duplication Identified in Therapy-Refractory Melanoma Patient-Derived Xenografts. <i>Cell Reports</i> , 2016, 16, 263-277.	6.4	61
6	SMARCE1 suppresses EGFR expression and controls responses to MET and ALK inhibitors in lung cancer. <i>Cell Research</i> , 2015, 25, 445-458.	12.0	36
7	RAF Suppression Synergizes with MEK Inhibition in KRAS Mutant Cancer Cells. <i>Cell Reports</i> , 2014, 8, 1475-1483.	6.4	103
8	Reversible and adaptive resistance to BRAF(V600E) inhibition in melanoma. <i>Nature</i> , 2014, 508, 118-122.	27.8	702
9	Feedback and redundancy in receptor tyrosine kinase signaling: relevance to cancer therapies. <i>Trends in Biochemical Sciences</i> , 2014, 39, 465-474.	7.5	134
10	Intrinsic Resistance to MEK Inhibition in KRAS Mutant Lung and Colon Cancer through Transcriptional Induction of ERBB3. <i>Cell Reports</i> , 2014, 7, 86-93.	6.4	266
11	MED12 Controls the Response to Multiple Cancer Drugs through Regulation of TGF- $\beta$ 2 Receptor Signaling. <i>Cell</i> , 2012, 151, 937-950.	28.9	371
12	Unresponsiveness of colon cancer to BRAF(V600E) inhibition through feedback activation of EGFR. <i>Nature</i> , 2012, 483, 100-103.	27.8	1,769
13	Global gene disruption in human cells to assign genes to phenotypes by deep sequencing. <i>Nature Biotechnology</i> , 2011, 29, 542-546.	17.5	207