

# Gui-Bo Li

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5334906/publications.pdf>

Version: 2024-02-01

10  
papers

537  
citations

1307594

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h-index

1372567

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g-index

11  
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docs citations

11  
times ranked

879  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic alterations and evolution of cell clusters in metastatic invasive micropapillary carcinoma of the breast. <i>Nature Communications</i> , 2022, 13, 111.	12.8	13
2	Chord: an ensemble machine learning algorithm to identify doublets in single-cell RNA sequencing data. <i>Communications Biology</i> , 2022, 5, .	4.4	6
3	Tumor heterogeneity assessed by sequencing and fluorescence <i>in situ</i> hybridization (FISH) data. <i>Bioinformatics</i> , 2021, 37, 4704-4711.	4.1	5
4	Dissecting spatial heterogeneity and the immune-evasion mechanism of CTCs by single-cell RNA-seq in hepatocellular carcinoma. <i>Nature Communications</i> , 2021, 12, 4091.	12.8	90
5	Single-cell exome sequencing reveals multiple subclones in metastatic colorectal carcinoma. <i>Genome Medicine</i> , 2021, 13, 148.	8.2	14
6	Neoadjuvant chemotherapy alters peripheral and tumour-infiltrating immune cells in breast cancer revealed by single-cell RNA sequencing. <i>Clinical and Translational Medicine</i> , 2021, 11, e621.	4.0	9
7	Single-cell transcriptomics reveals regulators underlying immune cell diversity and immune subtypes associated with prognosis in nasopharyngeal carcinoma. <i>Cell Research</i> , 2020, 30, 1024-1042.	12.0	182
8	Deconvolution of single-cell multi-omics layers reveals regulatory heterogeneity. <i>Nature Communications</i> , 2019, 10, 470.	12.8	156
9	Population-wide sampling of retrotransposon insertion polymorphisms using deep sequencing and efficient detection. <i>GigaScience</i> , 2017, 6, 1-11.	6.4	11
10	Full-length single-cell RNA-seq applied to a viral human cancer: applications to HPV expression and splicing analysis in HeLa S3 cells. <i>GigaScience</i> , 2015, 4, 51.	6.4	51