

# Anke Busch

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3464262/publications.pdf>

Version: 2024-02-01

10  
papers

700  
citations

1040056

9  
h-index

1474206

9  
g-index

10  
all docs

10  
docs citations

10  
times ranked

1516  
citing authors

#	ARTICLE	IF	CITATIONS
1	The RNA-binding profile of the splicing factor SRSF6 in immortalized human pancreatic Î²-cells. <i>Life Science Alliance</i> , 2021, 4, e202000825.	2.8	14
2	iCLIP data analysis: A complete pipeline from sequencing reads to RBP binding sites. <i>Methods</i> , 2020, 178, 49-62.	3.8	45
3	Coupling between alternative polyadenylation and alternative splicing is limited to terminal introns. <i>RNA Biology</i> , 2016, 13, 646-655.	3.1	34
4	Splicing predictions reliably classify different types of alternative splicing. <i>Rna</i> , 2015, 21, 813-823.	3.5	22
5	The TCF C-clamp DNA binding domain expands the Wnt transcriptome via alternative target recognition. <i>Nucleic Acids Research</i> , 2014, 42, 13615-13632.	14.5	34
6	The chromatin regulator Brg1 suppresses formation of intraductal papillary mucinous neoplasm and pancreatic ductal adenocarcinoma. <i>Nature Cell Biology</i> , 2014, 16, 255-267.	10.3	172
7	HEXEvent: a database of Human EXon splicing Events. <i>Nucleic Acids Research</i> , 2012, 41, D118-D124.	14.5	64
8	Extensive regulation of NAGNAG alternative splicing: new tricks for the spliceosome?. <i>Genome Biology</i> , 2012, 13, 143.	9.6	14
9	Evolution of SR protein and hnRNP splicing regulatory factors. <i>Wiley Interdisciplinary Reviews RNA</i> , 2012, 3, 1-12.	6.4	301
10	Computational prediction of alternative splice site selection. <i>FASEB Journal</i> , 2012, 26, 748.2.	0.5	0