

# Philip C Zuzarte

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2810912/publications.pdf>

Version: 2024-02-01

10  
papers

1,902  
citations

1163117

8  
h-index

1372567

10  
g-index

12  
all docs

12  
docs citations

12  
times ranked

4633  
citing authors

#	ARTICLE	IF	CITATIONS
1	Cryptic genomic lesions in adverse-risk acute myeloid leukemia identified by integrated whole genome and transcriptome sequencing. <i>Leukemia</i> , 2020, 34, 306-311.	7.2	14
2	Detection of Structural Rearrangements in Embryos. <i>New England Journal of Medicine</i> , 2020, 382, 2472-2474.	27.0	17
3	Comparative toxicoproteogenomics of mouse and rat liver identifies TCDD-resistance genes. <i>Archives of Toxicology</i> , 2019, 93, 2961-2978.	4.2	3
4	Nanopore native RNA sequencing of a human poly(A) transcriptome. <i>Nature Methods</i> , 2019, 16, 1297-1305.	19.0	411
5	High Interpatient Variability in Molecular MRD Response to Consolidation Chemotherapy in Acute Myeloid Leukemia. <i>Blood</i> , 2019, 134, 1424-1424.	1.4	0
6	Sensitive tumour detection and classification using plasma cell-free DNA methylomes. <i>Nature</i> , 2018, 563, 579-583.	27.8	624
7	Prediction of acute myeloid leukaemia risk in healthy individuals. <i>Nature</i> , 2018, 559, 400-404.	27.8	617
8	Fine mapping of chromosome 5p15.33 based on a targeted deep sequencing and high density genotyping identifies novel lung cancer susceptibility loci. <i>Carcinogenesis</i> , 2016, 37, 96-105.	2.8	36
9	A Two-Dimensional Pooling Strategy for Rare Variant Detection on Next-Generation Sequencing Platforms. <i>PLoS ONE</i> , 2014, 9, e93455.	2.5	18
10	Monoubiquitylation of H2A.Z Distinguishes Its Association with Euchromatin or Facultative Heterochromatin. <i>Molecular and Cellular Biology</i> , 2007, 27, 6457-6468.	2.3	158