

# Mauricio Oberti

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1008546/publications.pdf>

Version: 2024-02-01

9  
papers

1,718  
citations

1478505

6  
h-index

1720034

7  
g-index

9  
all docs

9  
docs citations

9  
times ranked

3790  
citing authors

#	ARTICLE	IF	CITATIONS
1	Integrated Proteomic and Glycoproteomic Characterization of Human High-Grade Serous Ovarian Carcinoma. Cell Reports, 2020, 33, 108276.	6.4	83
2	<scp>cnnAlpha</scp>: Protein disordered regions prediction by reduced amino acid alphabets and convolutional neural networks. Proteins: Structure, Function and Bioinformatics, 2020, 88, 1472-1481.	2.6	5
3	Li-Fraumeni Exploration Consortium Data Coordinating Center: Building an Interactive Web-Based Resource for Collaborative International Cancer Epidemiology Research for a Rare Condition. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 927-935.	2.5	7
4	Proteogenomic Analysis of Human Colon Cancer Reveals New Therapeutic Opportunities. Cell, 2019, 177, 1035-1049.e19.	28.9	498
5	Identification and Prediction of Intrinsically Disordered Regions in Proteins Using n-grams. , 2017, , .		2
6	Integrated Proteogenomic Characterization of Human High-Grade Serous Ovarian Cancer. Cell, 2016, 166, 755-765.	28.9	804
7	The CPTAC Data Portal: A Resource for Cancer Proteomics Research. Journal of Proteome Research, 2015, 14, 2707-2713.	3.7	309
8	Informatics and data quality at collaborative multicenter Breast and Colon Cancer Family Registries. Journal of the American Medical Informatics Association: JAMIA, 2012, 19, e125-e128.	4.4	10
9	shiny-pred: a server for the prediction of protein disordered regions. F1000Research, 0, 8, 230.	1.6	0