

# Christopher Douville

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3597665/publications.pdf>

Version: 2024-02-01

25  
papers

4,156  
citations

471509

17  
h-index

610901

24  
g-index

31  
all docs

31  
docs citations

31  
times ranked

8106  
citing authors

#	ARTICLE	IF	CITATIONS
1	Detection and localization of surgically resectable cancers with a multi-analyte blood test. <i>Science</i> , 2018, 359, 926-930.	12.6	1,872
2	A Combination of Molecular Markers and Clinical Features Improve the Classification of Pancreatic Cysts. <i>Gastroenterology</i> , 2015, 149, 1501-1510.	1.3	376
3	Feasibility of blood testing combined with PET-CT to screen for cancer and guide intervention. <i>Science</i> , 2020, 369, .	12.6	351
4	Whole Genome Sequencing Defines the Genetic Heterogeneity of Familial Pancreatic Cancer. <i>Cancer Discovery</i> , 2016, 6, 166-175.	9.4	282
5	Mutational Signature of Aristolochic Acid Exposure as Revealed by Whole-Exome Sequencing. <i>Science Translational Medicine</i> , 2013, 5, 197ra102.	12.4	220
6	Evaluation of liquid from the Papanicolaou test and other liquid biopsies for the detection of endometrial and ovarian cancers. <i>Science Translational Medicine</i> , 2018, 10, .	12.4	178
7	CRAVAT: cancer-related analysis of variants toolkit. <i>Bioinformatics</i> , 2013, 29, 647-648.	4.1	140
8	A multimodality test to guide the management of patients with a pancreatic cyst. <i>Science Translational Medicine</i> , 2019, 11, .	12.4	129
9	Non-invasive detection of urothelial cancer through the analysis of driver gene mutations and aneuploidy. <i>ELife</i> , 2018, 7, .	6.0	118
10	Assessing the Pathogenicity of Insertion and Deletion Variants with the Variant Effect Scoring Tool (VEST4). <i>Human Mutation</i> , 2016, 37, 28-35.	2.5	101
11	CRAVAT 4: Cancer-Related Analysis of Variants Toolkit. <i>Cancer Research</i> , 2017, 77, e35-e38.	0.9	51
12	Assessing aneuploidy with repetitive element sequencing. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 4858-4863.	7.1	50
13	Detection of aneuploidy in patients with cancer through amplification of long interspersed nucleotide elements (LINEs). <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 1871-1876.	7.1	48
14	Proteins linked to autosomal dominant and autosomal recessive disorders harbor characteristic rare missense mutation distribution patterns. <i>Human Molecular Genetics</i> , 2015, 24, 5995-6002.	2.9	40
15	Detection of low-frequency DNA variants by targeted sequencing of the Watson and Crick strands. <i>Nature Biotechnology</i> , 2021, 39, 1220-1227.	17.5	40
16	Pembrolizumab for patients with leptomeningeal metastasis from solid tumors: efficacy, safety, and cerebrospinal fluid biomarkers. , 2021, 9, e002473.		33
17	Eliminating accidental deviations to minimize generalization error and maximize replicability: Applications in connectomics and genomics. <i>PLoS Computational Biology</i> , 2021, 17, e1009279.	3.2	28
18	Integration of genetic and metabolic features related to sialic acid metabolism distinguishes human breast cell subtypes. <i>PLoS ONE</i> , 2018, 13, e0195812.	2.5	24

#	ARTICLE	IF	CITATIONS
19	Massively Parallel Sequencing of Esophageal Brushings Enables an Aneuploidy-Based Classification of Patients With Barrett's Esophagus. <i>Gastroenterology</i> , 2021, 160, 2043-2054.e2.	1.3	17
20	Matching phenotypes to whole genomes: Lessons learned from four iterations of the personal genome project community challenges. <i>Human Mutation</i> , 2017, 38, 1266-1276.	2.5	14
21	A Probabilistic Model to Predict Clinical Phenotypic Traits from Genome Sequencing. <i>PLoS Computational Biology</i> , 2014, 10, e1003825.	3.2	10
22	Use of a Polygenic Risk Score Improves Prediction of Myocardial Injury After Non-Cardiac Surgery. <i>Circulation Genomic and Precision Medicine</i> , 2020, 13, e002817.	3.6	9
23	The mutational landscape of spinal chordomas and their sensitive detection using circulating tumor DNA. <i>Neuro-Oncology Advances</i> , 2021, 3, vdaa173.	0.7	6
24	Detection of malignant peripheral nerve sheath tumors in patients with neurofibromatosis using aneuploidy and mutation identification in plasma. <i>ELife</i> , 2022, 11, .	6.0	4
25	BIOM-58. ASSESSMENT OF ANEUPLOIDY BY REPETITIVE ELEMENT SEQUENCING TO DETECT PEDIATRIC AND ADULT BRAIN CANCERS IN PLASMA. <i>Neuro-Oncology</i> , 2020, 22, ii14-ii14.	1.2	0