

# John R Mcpherson

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10929334/publications.pdf>

Version: 2024-02-01

17  
papers

5,269  
citations

567281

15  
h-index

888059

17  
g-index

18  
all docs

18  
docs citations

18  
times ranked

10568  
citing authors

#	ARTICLE	IF	CITATIONS
1	The repertoire of mutational signatures in human cancer. <i>Nature</i> , 2020, 578, 94-101.	27.8	2,104
2	Functional genomics identifies specific vulnerabilities in PTEN-deficient breast cancer. <i>Breast Cancer Research</i> , 2018, 20, 22.	5.0	15
3	A formalin-fixed paraffin-embedded (FFPE)-based prognostic signature to predict metastasis in clinically low risk stage I/II microsatellite stable colorectal cancer. <i>Cancer Letters</i> , 2017, 403, 13-20.	7.2	16
4	Whole-Genome and Epigenomic Landscapes of Etiologically Distinct Subtypes of Cholangiocarcinoma. <i>Cancer Discovery</i> , 2017, 7, 1116-1135.	9.4	637
5	Exome sequencing reveals recurrent REV3L mutations in cisplatin-resistant squamous cell carcinoma of head and neck. <i>Scientific Reports</i> , 2016, 6, 19552.	3.3	26
6	Transcription-associated mutation of lasR in <i>Pseudomonas aeruginosa</i> . <i>DNA Repair</i> , 2016, 46, 9-19.	2.8	11
7	<i>SETD2</i> histone modifier loss in aggressive GI stromal tumours. <i>Gut</i> , 2016, 65, 1960-1972.	12.1	49
8	MSIseq: Software for Assessing Microsatellite Instability from Catalogs of Somatic Mutations. <i>Scientific Reports</i> , 2015, 5, 13321.	3.3	113
9	Mutational landscapes of tongue carcinoma reveal recurrent mutations in genes of therapeutic and prognostic relevance. <i>Genome Medicine</i> , 2015, 7, 98.	8.2	74
10	Genomic landscapes of breast fibroepithelial tumors. <i>Nature Genetics</i> , 2015, 47, 1341-1345.	21.4	167
11	Mutation signatures implicate aristolochic acid in bladder cancer development. <i>Genome Medicine</i> , 2015, 7, 38.	8.2	87
12	Mutation signatures of carcinogen exposure: genome-wide detection and new opportunities for cancer prevention. <i>Genome Medicine</i> , 2014, 6, 24.	8.2	75
13	Exome sequencing identifies highly recurrent MED12 somatic mutations in breast fibroadenoma. <i>Nature Genetics</i> , 2014, 46, 877-880.	21.4	172
14	Exome sequencing identifies distinct mutational patterns in liver fluke-associated and non-infection-related bile duct cancers. <i>Nature Genetics</i> , 2013, 45, 1474-1478.	21.4	426
15	Genome-Wide Mutational Signatures of Aristolochic Acid and Its Application as a Screening Tool. <i>Science Translational Medicine</i> , 2013, 5, 197ra101.	12.4	233
16	Exome sequencing of gastric adenocarcinoma identifies recurrent somatic mutations in cell adhesion and chromatin remodeling genes. <i>Nature Genetics</i> , 2012, 44, 570-574.	21.4	560
17	Exome sequencing of liver fluke-associated cholangiocarcinoma. <i>Nature Genetics</i> , 2012, 44, 690-693.	21.4	412