

Ian Bosdet

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

3,752
citations

840776

11
h-index

794594

19
g-index

20
all docs

20
docs citations

20
times ranked

5994
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome sequence of the Brown Norway rat yields insights into mammalian evolution. <i>Nature</i> , 2004, 428, 493-521.	27.8	1,943
2	The Genome of the Sea Urchin <i>Strongylocentrotus purpuratus</i> . <i>Science</i> , 2006, 314, 941-952.	12.6	1,018
3	A physical map of the mouse genome. <i>Nature</i> , 2002, 418, 743-750.	27.8	316
4	Homologous Recombination Deficiency and Platinum-Based Therapy Outcomes in Advanced Breast Cancer. <i>Clinical Cancer Research</i> , 2017, 23, 7521-7530.	7.0	144
5	A set of BAC clones spanning the human genome. <i>Nucleic Acids Research</i> , 2004, 32, 3651-3660.	14.5	119
6	Integrated and Sequence-Ordered BAC- and YAC-Based Physical Maps for the Rat Genome. <i>Genome Research</i> , 2004, 14, 766-779.	5.5	44
7	Physical Maps for Genome Analysis of Serotype A and D Strains of the Fungal Pathogen <i>Cryptococcus neoformans</i> . <i>Genome Research</i> , 2002, 12, 1445-1453.	5.5	38
8	Data sharing as a national quality improvement program: reporting on BRCA1 and BRCA2 variant-interpretation comparisons through the Canadian Open Genetics Repository (COGR). <i>Genetics in Medicine</i> , 2018, 20, 294-302.	2.4	27
9	Internet Contig Explorer (ICE)—A Tool for Visualizing Clone Fingerprint Maps. <i>Genome Research</i> , 2003, 13, 1244-1249.	5.5	22
10	A BAC-based physical map of the <i>Drosophila buzzatii</i> genome. <i>Genome Research</i> , 2005, 15, 885-889.	5.5	21
11	MET exon 14 skipping mutation positive non-small cell lung cancer: Response to systemic therapy. <i>Lung Cancer</i> , 2021, 154, 142-145.	2.0	14
12	Data sharing to improve concordance in variant interpretation across laboratories: results from the Canadian Open Genetics Repository. <i>Journal of Medical Genetics</i> , 2022, 59, 571-578.	3.2	14
13	Sample Tracking Using Unique Sequence Controls. <i>Journal of Molecular Diagnostics</i> , 2020, 22, 141-146.	2.8	10
14	Assessing Limit of Detection in Clinical Sequencing. <i>Journal of Molecular Diagnostics</i> , 2021, 23, 455-466.	2.8	7
15	Establishing a Framework for the Clinical Translation of Germline Findings in Precision Oncology. <i>JNCI Cancer Spectrum</i> , 2020, 4, pkaa045.	2.9	6
16	EGFR circulating tumour DNA testing: identification of predictors of ctDNA detection and implications for survival outcomes. <i>Translational Lung Cancer Research</i> , 2020, 9, 1084-1092.	2.8	5
17	Use of Treatment-Focused Tumor Sequencing to Screen for Germline Cancer Predisposition. <i>Journal of Molecular Diagnostics</i> , 2021, 23, 1145-1158.	2.8	2
18	Integration of Whole-Genome Sequencing With Circulating Tumor DNA Analysis Captures Clonal Evolution and Tumor Heterogeneity in Non-V600 BRAF Mutant Colorectal Cancer. <i>Clinical Colorectal Cancer</i> , 2020, 19, 132-136.e3.	2.3	1

#	ARTICLE	IF	CITATIONS
19	Integrating Tumor Sequencing Into Clinical Practice for Patients With Mismatch Repair-Deficient Lynch Syndrome Spectrum Cancers. <i>Clinical and Translational Gastroenterology</i> , 2021, 12, e00397.	2.5	1
20	Towards the Human Cancer Genome Project: A Sequence-Ready Physical Map of a Follicular Lymphoma Genome.. <i>Blood</i> , 2005, 106, 605-605.	1.4	0