

Robert K Hastings

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1774430/publications.pdf>

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16
papers

1,915
citations

840776

11
h-index

996975

15
g-index

17
all docs

17
docs citations

17
times ranked

5388
citing authors

#	ARTICLE	IF	CITATIONS
1	Inclusion of multiple high-risk histopathological criteria improves the prediction of adjuvant chemotherapy efficacy in lung adenocarcinoma. <i>Histopathology</i> , 2021, 78, 838-848.	2.9	9
2	Comparison of two targeted ultra-deep sequencing technologies for analysis of plasma circulating tumour DNA in endocrine-therapy-resistant breast cancer patients. <i>Breast Cancer Research and Treatment</i> , 2021, 188, 465-476.	2.5	1
3	Circulating Tumor DNA Profiling From Breast Cancer Screening Through to Metastatic Disease. <i>JCO Precision Oncology</i> , 2021, 5, 1768-1776.	3.0	12
4	The Circulating Nucleic Acid Characteristics of Non-Metastatic Soft Tissue Sarcoma Patients. <i>International Journal of Molecular Sciences</i> , 2020, 21, 4483.	4.1	8
5	Personalized Detection of Circulating Tumor DNA Antedates Breast Cancer Metastatic Recurrence. <i>Clinical Cancer Research</i> , 2019, 25, 4255-4263.	7.0	281
6	Plasma cell-free DNA (cfDNA) as a predictive and prognostic marker in patients with metastatic breast cancer. <i>Breast Cancer Research</i> , 2019, 21, 149.	5.0	89
7	Circulating tumour-derived DNA in metastatic soft tissue sarcoma. <i>Oncotarget</i> , 2018, 9, 10549-10560.	1.8	29
8	Mutation Analysis of Cell-Free DNA and Single Circulating Tumor Cells in Metastatic Breast Cancer Patients with High Circulating Tumor Cell Counts. <i>Clinical Cancer Research</i> , 2017, 23, 88-96.	7.0	186
9	Next Generation Sequencing of Circulating Cell-Free DNA for Evaluating Mutations and Gene Amplification in Metastatic Breast Cancer. <i>Clinical Chemistry</i> , 2017, 63, 532-541.	3.2	81
10	Profiling tumour heterogeneity through circulating tumour DNA in patients with pancreatic cancer. <i>Oncotarget</i> , 2017, 8, 87221-87233.	1.8	38
11	Noninvasive Detection of Activating Estrogen Receptor 1 (ESR1) Mutations in Estrogen Receptor-Positive Metastatic Breast Cancer. <i>Clinical Chemistry</i> , 2015, 61, 974-982.	3.2	155
12	The BioMart community portal: an innovative alternative to large, centralized data repositories. <i>Nucleic Acids Research</i> , 2015, 43, W589-W598.	14.5	682
13	GWAS Central: a comprehensive resource for the comparison and interrogation of genome-wide association studies. <i>European Journal of Human Genetics</i> , 2014, 22, 949-952.	2.8	155
14	Extension to Distributed Annotation System: Summary command. , 2014, , .		1
15	Epigenetic Control of Retrotransposon Expression in Human Embryonic Stem Cells. <i>Molecular and Cellular Biology</i> , 2011, 31, 300-316.	2.3	128
16	HGVbaseG2P: a central genetic association database. <i>Nucleic Acids Research</i> , 2009, 37, D797-D802.	14.5	60