

VÃ©steinn Thorsson

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4070441/publications.pdf>

Version: 2024-02-01

24
papers

13,132
citations

471509

17
h-index

677142

22
g-index

28
all docs

28
docs citations

28
times ranked

20091
citing authors

#	ARTICLE	IF	CITATIONS
1	The Immune Landscape of Cancer. Immunity, 2018, 48, 812-830.e14.	14.3	3,706
2	An Integrated TCGA Pan-Cancer Clinical Data Resource to Drive High-Quality Survival Outcome Analytics. Cell, 2018, 173, 400-416.e11.	28.9	2,277
3	Integrated Genomic and Proteomic Analyses of a Systematically Perturbed Metabolic Network. Science, 2001, 292, 929-934.	12.6	1,921
4	Cell-of-Origin Patterns Dominate the Molecular Classification of 10,000 Tumors from 33 Types of Cancer. Cell, 2018, 173, 291-304.e6.	28.9	1,718
5	Spatial Organization and Molecular Correlation of Tumor-Infiltrating Lymphocytes Using Deep Learning on Pathology Images. Cell Reports, 2018, 23, 181-193.e7.	6.4	683
6	Integrative Analysis Identifies Four Molecular and Clinical Subsets in Uveal Melanoma. Cancer Cell, 2017, 32, 204-220.e15.	16.8	642
7	Integrative Molecular Characterization of Malignant Pleural Mesothelioma. Cancer Discovery, 2018, 8, 1548-1565.	9.4	422
8	Comparative Molecular Analysis of Gastrointestinal Adenocarcinomas. Cancer Cell, 2018, 33, 721-735.e8.	16.8	396
9	Integrated Molecular Characterization of Testicular Germ Cell Tumors. Cell Reports, 2018, 23, 3392-3406.	6.4	324
10	Perspective on Oncogenic Processes at the End of the Beginning of Cancer Genomics. Cell, 2018, 173, 305-320.e10.	28.9	272
11	Meta-analysis of five genome-wide association studies identifies multiple new loci associated with testicular germ cell tumor. Nature Genetics, 2017, 49, 1141-1147.	21.4	105
12	Germline genetic contribution to the immune landscape of cancer. Immunity, 2021, 54, 367-386.e8.	14.3	95
13	MITF has a central role in regulating starvation-induced autophagy in melanoma. Scientific Reports, 2019, 9, 1055.	3.3	66
14	Toward a comprehensive view of cancer immune responsiveness: a synopsis from the SITC workshop. , 2019, 7, 131.		64
15	CRI iAtlas: an interactive portal for immuno-oncology research. F1000Research, 2020, 9, 1028.	1.6	39
16	MITI minimum information guidelines for highly multiplexed tissue images. Nature Methods, 2022, 19, 262-267.	19.0	37
17	Mutational Landscapes of Smoking-Related Cancers in Caucasians and African Americans: Precision Oncology Perspectives at Wake Forest Baptist Comprehensive Cancer Center. Theranostics, 2017, 7, 2914-2923.	10.0	31
18	Validation and calibration of next-generation sequencing to identify Epstein-Barr virus-positive gastric cancer in The Cancer Genome Atlas. Gastric Cancer, 2016, 19, 676-681.	5.3	15

#	ARTICLE	IF	CITATIONS
19	Reverse Engineering Galactose Regulation in Yeast through Model Selection. Statistical Applications in Genetics and Molecular Biology, 2005, 4, Article28.	0.6	9
20	Society for Immunotherapy of Cancer clinical and biomarkers data sharing resource document: Volume IIâ€”conceptual challenges. , 2020, 8, e001389.		7
21	Patient-Specific Cell Communication Networks Associate With Disease Progression in Cancer. Frontiers in Genetics, 2021, 12, 667382.	2.3	5
22	Society for Immunotherapy of Cancer clinical and biomarkers data sharing resource document: Volume IIâ€”practical challenges. , 2020, 8, e001472.		4
23	Multiplatform Integrative Analysis of Immunogenomic Data for Biomarker Discovery. Methods in Molecular Biology, 2020, 2055, 679-698.	0.9	1
24	P854â€”Construction of the immune landscape of durable response to checkpoint blockade therapy by integrating publicly available datasets. , 2020, , .		0