Gang Peng

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7327963/publications.pdf

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35	1,103	17 h-index	32
papers	citations		g-index
35	35	35	2217 citing authors
all docs	docs citations	times ranked	

#	Article	IF	Citations
1	Determining concordance of LFSPRO <i>TP53</i> germline carrier risk predictions to standard genetic counseling practice Journal of Clinical Oncology, 2022, 40, 10593-10593.	1.6	O
2	The role of the gut microbiome in cancer-related fatigue: pilot study on epigenetic mechanisms. Supportive Care in Cancer, 2021, 29, 3173-3182.	2.2	26
3	Periâ€implantation cytokine profile differs between singleton and twin IVF pregnancies. American Journal of Reproductive Immunology, 2021, 85, e13348.	1.2	4
4	Epigenetic age acceleration, fatigue, and inflammation in patients undergoing radiation therapy for head and neck cancer: A longitudinal study. Cancer, 2021, 127, 3361-3371.	4.1	28
5	Correlating genomic copy number alterations with clinicopathologic findings in 75 cases of hepatocellular carcinoma. BMC Medical Genomics, 2021, 14, 150.	1.5	5
6	Association of Epigenetic Age Acceleration With Risk Factors, Survival, and Quality of Life in Patients With Head and Neck Cancer. International Journal of Radiation Oncology Biology Physics, 2021, 111, 157-167.	0.8	18
7	ILâ€10 to TNFα ratios throughout early first trimester can discriminate healthy pregnancies from pregnancy losses. American Journal of Reproductive Immunology, 2020, 83, e13195.	1.2	32
8	A Class II-Restricted CD8γ13 T-Cell Clone Protects During Chlamydia muridarum Genital Tract Infection. Journal of Infectious Diseases, 2020, 221, 1895-1906.	4.0	7
9	Penetrance of Different Cancer Types in Families with Li-Fraumeni Syndrome: A Validation Study Using Multicenter Cohorts. Cancer Research, 2020, 80, 354-360.	0.9	22
10	Differential Protein Expression in Striatal D1- and D2-Dopamine Receptor-Expressing Medium Spiny Neurons. Proteomes, 2020, 8, 27.	3.5	6
11	Trophoblast-secreted soluble-PD-L1 modulates macrophage polarization and function. Journal of Leukocyte Biology, 2020, 108, 983-998.	3.3	45
12	Molecular and clinicopathologic characterization of intravenous leiomyomatosis. Modern Pathology, 2020, 33, 1844-1860.	5 . 5	16
13	Reducing False-Positive Results in Newborn Screening Using Machine Learning. International Journal of Neonatal Screening, 2020, 6, 16.	3.2	33
14	Ethnic variability in newborn metabolic screening markers associated with falseâ€positive outcomes. Journal of Inherited Metabolic Disease, 2020, 43, 934-943.	3.6	23
15	Human Chorionic Gonadotropin modulates CXCL10 Expression through Histone Methylation in human decidua. Scientific Reports, 2020, 10, 5785.	3.3	15
16	Timing of Newborn Blood Collection Alters Metabolic Disease Screening Performance. Frontiers in Pediatrics, 2020, 8, 623184.	1.9	14
17	Cytogenomic Abnormalities in 19 Cases of Salivary Gland Tumors of Parotid Gland Origin. Case Reports in Genetics, 2020, 2020, 1-6.	0.2	1
18	Combining newborn metabolic and DNA analysis for second-tier testing of methylmalonic acidemia. Genetics in Medicine, 2019, 21, 896-903.	2.4	31

#	Article	IF	Citations
19	Elevated methylmalonic acidemia (MMA) screening markers in Hispanic and preterm newborns. Molecular Genetics and Metabolism, 2019, 126, 39-42.	1.1	6
20	ProteomicsBrowser: MS/proteomics data visualization and investigation. Bioinformatics, 2019, 35, 2313-2314.	4.1	2
21	Machine learning selected smoking-associated DNA methylation signatures that predict HIV prognosis and mortality. Clinical Epigenetics, 2018, 10, 155.	4.1	37
22	Efficient computation of the joint probability of multiple inherited risk alleles from pedigree data. Genetic Epidemiology, 2018, 42, 528-538.	1.3	3
23	Estimating <i>TP53</i> Mutation Carrier Probability in Families with Li–Fraumeni Syndrome Using LFSPRO. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 837-844.	2.5	14
24	Data-Independent Acquisition and Parallel Reaction Monitoring Mass Spectrometry Identification of Serum Biomarkers for Ovarian Cancer. Biomarker Insights, 2017, 12, 117727191771094.	2.5	21
25	A Branched-Chain Amino Acid-Related Metabolic Signature Characterizes Obese Adolescents with Non-Alcoholic Fatty Liver Disease. Nutrients, 2017, 9, 642.	4.1	92
26	Next-Generation Molecular Testing of Newborn Dried Blood Spots for Cystic Fibrosis. Journal of Molecular Diagnostics, 2016, 18, 267-282.	2.8	26
27	Prevalence of Corneal Astigmatism in Patients before Cataract Surgery in Northern China. Journal of Ophthalmology, 2014, 2014, 1-7.	1.3	51
28	FamSeq: A Variant Calling Program for Family-Based Sequencing Data Using Graphics Processing Units. PLoS Computational Biology, 2014, 10, e1003880.	3.2	14
29	Rare variant detection using family-based sequencing analysis. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 3985-3990.	7.1	54
30	Gene and pathway-based second-wave analysis of genome-wide association studies. European Journal of Human Genetics, 2010, 18, 111-117.	2.8	228
31	Genome-wide gene and pathway analysis. European Journal of Human Genetics, 2010, 18, 1045-1053.	2.8	99
32	A Novel Statistic for Genome-Wide Interaction Analysis. PLoS Genetics, 2010, 6, e1001131.	3.5	64
33	Genome-Wide Association Studies of Copy Number Variation in Glioblastoma. International Conference on Bioinformatics and Biomedical Engineering: [proceedings] International Conference on Bioinformatics and Biomedical Engineering, 2010, , .	0.0	2
34	The Results on the Stability of Glycolytic Metabolic Networks in Different Cells. Lecture Notes of the Institute for Computer Sciences, Social-Informatics and Telecommunications Engineering, 2009, , 536-540.	0.3	0
35	Multifactor Effects and Evidence of Potential Interaction between Complement Factor H Y402H and LOC387715 A69S in Age-Related Macular Degeneration. PLoS ONE, 2008, 3, e3833.	2.5	64