

Mohamad Saad

List of Publications by Year in descending order

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Version: 2024-02-01

13
papers

322
citations

1163065

8
h-index

1199563

12
g-index

14
all docs

14
docs citations

14
times ranked

569
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic predisposition to cancer across people of different ancestries in Qatar: a population-based, cohort study. <i>Lancet Oncology</i> , The, 2022, 23, 341-352.	10.7	14
2	Untargeted Metabolomics Profiling Reveals Perturbations in Arginine-NO Metabolism in Middle Eastern Patients with Coronary Heart Disease. <i>Metabolites</i> , 2022, 12, 517.	2.9	5
3	Including diverse and admixed populations in genetic epidemiology research. <i>Genetic Epidemiology</i> , 2022, 46, 347-371.	1.3	11
4	Germline genetic contribution to the immune landscape of cancer. <i>Immunity</i> , 2021, 54, 367-386.e8.	14.3	95
5	Network-based identification of key master regulators associated with an immune-silent cancer phenotype. <i>Briefings in Bioinformatics</i> , 2021, 22, .	6.5	11
6	Distribution of Primary Brain Tumor Subtypes in Lebanon: A Multicenter Eleven-Year Study of 695 Patients. <i>Cureus</i> , 2021, 13, e17918.	0.5	2
7	Oncogenic states dictate the prognostic and predictive connotations of intratumoral immune response. , 2020, 8, e000617.		57
8	Prognostic impact of adenylyl cyclase-associated protein 2 (CAP2) in glioma: A clinicopathological study. <i>Heliyon</i> , 2020, 6, e03236.	3.2	5
9	Quality control and integration of genotypes from two calling pipelines for whole genome sequence data in the Alzheimer's disease sequencing project. <i>Genomics</i> , 2019, 111, 808-818.	2.9	26
10	Post-traumatic stress disorder in adult victims of cluster munitions in Lebanon: a 10-year longitudinal study. <i>BMJ Open</i> , 2017, 7, e017214.	1.9	34
11	Identity-by-descent estimation with population- and pedigree-based imputation in admixed family data. <i>BMC Proceedings</i> , 2016, 10, 295-301.	1.6	4
12	Power of Family-Based Association Designs to Detect Rare Variants in Large Pedigrees Using Imputed Genotypes. <i>Genetic Epidemiology</i> , 2014, 38, 1-9.	1.3	27
13	Combining Family- and Population-Based Imputation Data for Association Analysis of Rare and Common Variants in Large Pedigrees. <i>Genetic Epidemiology</i> , 2014, 38, 579-590.	1.3	28