

Marie Forest

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7605921/publications.pdf>

Version: 2024-02-01

18
papers

161
citations

1683934

5
h-index

1872570

6
g-index

18
all docs

18
docs citations

18
times ranked

440
citing authors

#	ARTICLE	IF	CITATIONS
1	Improved prediction of fracture risk leveraging a genome-wide polygenic risk score. <i>Genome Medicine</i> , 2021, 13, 16.	3.6	35
2	Simultaneous SNP selection and adjustment for population structure in high dimensional prediction models. <i>PLoS Genetics</i> , 2020, 16, e1008766.	1.5	5
3	Development of a polygenic risk score to improve screening for fracture risk: A genetic risk prediction study. <i>PLoS Medicine</i> , 2020, 17, e1003152.	3.9	45
4	Title is missing!. , 2020, 16, e1008766.		0
5	Title is missing!. , 2020, 16, e1008766.		0
6	Title is missing!. , 2020, 16, e1008766.		0
7	Title is missing!. , 2020, 16, e1008766.		0
8	Title is missing!. , 2020, 16, e1008766.		0
9	Title is missing!. , 2020, 16, e1008766.		0
10	Title is missing!. , 2020, 17, e1003152.		0
11	Title is missing!. , 2020, 17, e1003152.		0
12	Title is missing!. , 2020, 17, e1003152.		0
13	Title is missing!. , 2020, 17, e1003152.		0
14	Title is missing!. , 2020, 17, e1003152.		0
15	Title is missing!. , 2020, 17, e1003152.		0
16	Agreement in DNA methylation levels from the Illumina 450K array across batches, tissues, and time. <i>Epigenetics</i> , 2018, 13, 19-32.	1.3	39
17	Gene networks show associations with seed region connectivity. <i>Human Brain Mapping</i> , 2017, 38, 3126-3140.	1.9	32
18	The performance of a new local false discovery rate method on tests of association between coronary artery disease (CAD) and genome-wide genetic variants. <i>PLoS ONE</i> , 2017, 12, e0185174.	1.1	5