Navya S Josyula

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/8513613/publications.pdf

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14	1,470	759233 12	1058476 14
papers	citations	h-index	g-index

19	19	19	3068
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Genetic loci and prioritization of genes for kidney function decline derived from a meta-analysis of 62 longitudinal genome-wide association studies. Kidney International, 2022, 102, 624-639.	5.2	18
2	Differential and shared genetic effects on kidney function between diabetic and non-diabetic individuals. Communications Biology, 2022, 5, .	4.4	17
3	Meta-analysis uncovers genome-wide significant variants for rapid kidney function decline. Kidney International, 2021, 99, 926-939.	5.2	42
4	Exome-wide evaluation of rare coding variants using electronic health records identifies new geneâ€“phenotype associations. Nature Medicine, 2021, 27, 66-72.	30.7	44
5	Genome-wide meta-analysis identifies 127 open-angle glaucoma loci with consistent effect across ancestries. Nature Communications, 2021, 12, 1258.	12.8	196
6	Gene co-regulation and co-expression in the aryl hydrocarbon receptor-mediated transcriptional regulatory network in the mouse liver. Archives of Toxicology, 2020, 94, 113-126.	4.2	11
7	Identifying Ligand Binding Sites of Proteins using Crystallographic Bfactors and Relative Pocket Sizes. Biophysical Journal, 2020, 118, 48a.	0.5	0
8	Genome-wide association meta-analyses and fine-mapping elucidate pathways influencing albuminuria. Nature Communications, 2019, 10, 4130.	12.8	133
9	Harmonizing Clinical Sequencing and Interpretation for the eMERGE III Network. American Journal of Human Genetics, 2019, 105, 588-605.	6.2	99
10	Target genes, variants, tissues and transcriptional pathways influencing human serum urate levels. Nature Genetics, 2019, 51, 1459-1474.	21.4	251
11	A catalog of genetic loci associated with kidney function from analyses of a million individuals. Nature Genetics, 2019, 51, 957-972.	21.4	549
12	PheWAS and Beyond: The Landscape of Associations with Medical Diagnoses and Clinical Measures across 38,662 Individuals from Geisinger. American Journal of Human Genetics, 2018, 102, 592-608.	6.2	66
13	Rare variants in drug target genes contributing to complex diseases, phenome-wide. Scientific Reports, 2018, 8, 4624.	3.3	13
14	Healthcare Utilization and Patientsâ€™ Perspectives After Receiving a Positive Genetic Test for Familial Hypercholesterolemia. Circulation Genomic and Precision Medicine, 2018, 11, e002146.	3.6	23