

L Core Project R03OD032630

Name Methods to maximize the utility of common fund functional genomic

data in multi-ethnic genetic studies

Projects 1R03OD032630-01

Award \$335,407.00

Publications (8)

ID	RCR	Citations	Cit./year	Title	Authors	Journal	Published	Updated
30643251 🗗 DOI 🗹	62	1,040	208	Association studies of up to 1.2 mill	Mengzhen Liu 121 more Scott Vrieze	Nat Genet	2019	Jun 30, 2024
36477530 🗗 DOI 🗗	18.63	102	51	Genetic diversity fuels gene discover	Gretchen R B Saunders 215 more Scott Vrieze	Nature	2022	Jun 30, 2024

36702996 ♂ DOI ♂	6.83	18	18	Multi- ancestry transcripto me-wide ass	Fang Chen 88 more Dajiang J Liu	Nat Genet	2023	Jun 30, 2024
<u>36750564</u>	4.54	12	12	Multi- ancestry and multi- trait genome	Chachrit Khunsrirak sakul 16 more Dajiang J Liu	Nat Commun	2023	Jun 30, 2024
35672318 🗹 DOI 🗹	1.76	14	7	Integrating 3D genomic and epigenomic 	Chachrit Khunsrirak sakul 12 more Dajiang J Liu	Nat Commun	2022	Jun 30, 2024
35927319	1.49	11	5.5	Rare genetic variants explain missing	Seon- Kyeong Jang 88 more Scott Vrieze	Nat Hum Behav	2022	Jun 30, 2024
35833142 🗗 DOI 🗗	0.88	6	3	Constructio n and Application of Polyg	Chachrit Khunsrirak sakul 4 more Dajiang J Liu	Front Immunol	2022	Jun 30, 2024

35178743 2 0.34 2 1	Assessing Roopali reproducibi Singh lity of high1 more Qunhua Li	Stat Med	2022	Jun 30, 2024
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