

Yoichi Suzuki

List of Publications by Year in descending order

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Version: 2024-02-01

18
papers

709
citations

840776

11
h-index

839539

18
g-index

18
all docs

18
docs citations

18
times ranked

1086
citing authors

#	ARTICLE	IF	CITATIONS
1	The return of individual genomic results to research participants: design and pilot study of Tohoku Medical Megabank Project. <i>Journal of Human Genetics</i> , 2022, 67, 9-17.	2.3	9
2	Maternal Baseline Characteristics and Perinatal Outcomes: The Tohoku Medical Megabank Project Birth and Three-Generation Cohort Study. <i>Journal of Epidemiology</i> , 2022, 32, 69-79.	2.4	13
3	Study Profile of the Tohoku Medical Megabank Community-Based Cohort Study. <i>Journal of Epidemiology</i> , 2021, 31, 65-76.	2.4	81
4	Association between Serum Biotin Levels and Cedar Pollinosis in Japanese Schoolchildren. <i>Journal of Nutritional Science and Vitaminology</i> , 2021, 67, 211-216.	0.6	1
5	Cohort Profile: Tohoku Medical Megabank Project Birth and Three-Generation Cohort Study (TMM) Tj ETQq1 1 0.784314 rgBT /Overlook 2020, 49, 18-19m.	1.9	107
6	Breastfeeding promotes egg white sensitization in early infancy. <i>Pediatric Allergy and Immunology</i> , 2020, 31, 315-318.	2.6	6
7	Oral Microbiome Analysis in Prospective Genome Cohort Studies of the Tohoku Medical Megabank Project. <i>Frontiers in Cellular and Infection Microbiology</i> , 2020, 10, 604596.	3.9	12
8	A training and education program for genome medical research coordinators in the genome cohort study of the Tohoku Medical Megabank Organization. <i>BMC Medical Education</i> , 2019, 19, 297.	2.4	9
9	Skin Care and Synbiotics for Prevention of Atopic Dermatitis or Food Allergy in Newborn Infants: A 2 × 2 Factorial, Randomized, Non-Treatment Controlled Trial. <i>International Archives of Allergy and Immunology</i> , 2019, 180, 202-211.	2.1	54
10	Construction of JRG (Japanese reference genome) with single-molecule real-time sequencing. <i>Human Genome Variation</i> , 2019, 6, 27.	0.7	9
11	Susceptibility Loci for Tanning Ability in the Japanese Population Identified by a Genome-Wide Association Study from the Tohoku Medical Megabank Project Cohort Study. <i>Journal of Investigative Dermatology</i> , 2019, 139, 1605-1608.e13.	0.7	14
12	Estimating carrier frequencies of newborn screening disorders using a whole-genome reference panel of 3552 Japanese individuals. <i>Human Genetics</i> , 2019, 138, 389-409.	3.8	7
13	Genome analyses for the Tohoku Medical Megabank Project towards establishment of personalized healthcare. <i>Journal of Biochemistry</i> , 2019, 165, 139-158.	1.7	33
14	Evaluation of reported pathogenic variants and their frequencies in a Japanese population based on a whole-genome reference panel of 2049 individuals. <i>Journal of Human Genetics</i> , 2018, 63, 213-230.	2.3	35
15	Security controls in an integrated Biobank to protect privacy in data sharing: rationale and study design. <i>BMC Medical Informatics and Decision Making</i> , 2017, 17, 100.	3.0	30
16	The Tohoku Medical Megabank Project: Design and Mission. <i>Journal of Epidemiology</i> , 2016, 26, 493-511.	2.4	236
17	The structural origin of metabolic quantitative diversity. <i>Scientific Reports</i> , 2016, 6, 31463.	3.3	18
18	Exonic Mutations in the SLC12A3 Gene Cause Exon Skipping and Premature Termination in Gitelman Syndrome. <i>Journal of the American Society of Nephrology: JASN</i> , 2015, 26, 271-279.	6.1	35