

Esther van Enckevort

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

616
citations

759055

12
h-index

752573

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21
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21
docs citations

21
times ranked

1890
citing authors

#	ARTICLE	IF	CITATIONS
1	LongITools: Dynamic longitudinal exposome trajectories in cardiovascular and metabolic noncommunicable diseases. <i>Environmental Epidemiology</i> , 2022, 6, e184.	1.4	6
2	The EU Child Cohort Network's core data: establishing a set of findable, accessible, interoperable and re-usable (FAIR) variables. <i>European Journal of Epidemiology</i> , 2021, 36, 565-580.	2.5	24
3	BBMRI-ERIC Negotiator: Implementing Efficient Access to Biobanks. <i>Biopreservation and Biobanking</i> , 2021, 19, 414-421.	0.5	2
4	The de novo FAIRification process of a registry for vascular anomalies. <i>Orphanet Journal of Rare Diseases</i> , 2021, 16, 376.	1.2	8
5	Advancing tools for human early lifecourse exposome research and translation (ATHLETE). <i>Environmental Epidemiology</i> , 2021, 5, e166.	1.4	24
6	The case for open science: rare diseases. <i>JAMIA Open</i> , 2020, 3, 472-486.	1.0	33
7	BBMRI-ERIC's contributions to research and knowledge exchange on COVID-19. <i>European Journal of Human Genetics</i> , 2020, 28, 728-731.	1.4	17
8	Extending the Minimum Information About Biobank Data Sharing Terminology to Describe Samples, Sample Donors, and Events. <i>Biopreservation and Biobanking</i> , 2020, 18, 155-164.	0.5	25
9	Privacy-Preserving Linkage of Genomic and Clinical Data Sets. <i>IEEE/ACM Transactions on Computational Biology and Bioinformatics</i> , 2019, 16, 1342-1348.	1.9	18
10	MOLGENIS research: advanced bioinformatics data software for non-bioinformaticians. <i>Bioinformatics</i> , 2019, 35, 1076-1078.	1.8	58
11	MECP2 variation in Rett syndrome-An overview of current coverage of genetic and phenotype data within existing databases. <i>Human Mutation</i> , 2018, 39, 914-924.	1.1	15
12	Recommendations for Improving the Quality of Rare Disease Registries. <i>International Journal of Environmental Research and Public Health</i> , 2018, 15, 1644.	1.2	116
13	OSSE Goes FAIR - Implementation of the FAIR Data Principles for an Open-Source Registry for Rare Diseases. <i>Studies in Health Technology and Informatics</i> , 2018, 253, 209-213.	0.2	8
14	BiobankUniverse: automatic matchmaking between datasets for biobank data discovery and integration. <i>Bioinformatics</i> , 2017, 33, 3627-3634.	1.8	3
15	Systematically linking tranSMART, Galaxy and EGA for reusing human translational research data. <i>F1000Research</i> , 2017, 6, 1488.	0.8	8
16	BBMRI-ERIC Directory: 515 Biobanks with Over 60 Million Biological Samples. <i>Biopreservation and Biobanking</i> , 2016, 14, 559-562.	0.5	68
17	MOLGENIS/connect: a system for semi-automatic integration of heterogeneous phenotype data with applications in biobanks. <i>Bioinformatics</i> , 2016, 32, 2176-2183.	1.8	12
18	Toward Global Biobank Integration by Implementation of the Minimum Information About Biobank Data Sharing (MIABIS 2.0 Core). <i>Biopreservation and Biobanking</i> , 2016, 14, 298-306.	0.5	66

#	ARTICLE	IF	CITATIONS
19	Integration of EGA secure data access into Galaxy. F1000Research, 2016, 5, 2841.	0.8	7
20	Improved imputation quality of low-frequency and rare variants in European samples using the "Genome of The Netherlands"™. European Journal of Human Genetics, 2014, 22, 1321-1326.	1.4	92