

# Elizabeth Ormondroyd

## List of Publications by Year in descending order

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Version: 2024-02-01

28  
papers

1,631  
citations

516710

16  
h-index

526287

27  
g-index

28  
all docs

28  
docs citations

28  
times ranked

3851  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic health data generation in the UK: a 360 view. <i>European Journal of Human Genetics</i> , 2022, 30, 782-789.	2.8	8
2	Provocation Testing and Therapeutic Response in a Newly Described Channelopathy: RyR2 Calcium Release Deficiency Syndrome. <i>Circulation Genomic and Precision Medicine</i> , 2022, 15, CIRCGEN121003589.	3.6	15
3	Genomic sequencing in oncology: Considerations for integration in routine cancer care. <i>European Journal of Cancer Care</i> , 2022, 31, e13584.	1.5	4
4	Common genetic variants and modifiable risk factors underpin hypertrophic cardiomyopathy susceptibility and expressivity. <i>Nature Genetics</i> , 2021, 53, 135-142.	21.4	165
5	Bi-allelic MCM10 variants associated with immune dysfunction and cardiomyopathy cause telomere shortening. <i>Nature Communications</i> , 2021, 12, 1626.	12.8	22
6	Maximal Wall Thickness Measurement in Hypertrophic Cardiomyopathy. <i>JACC: Cardiovascular Imaging</i> , 2021, 14, 2123-2134.	5.3	18
7	Taking it to the bank: the ethical management of individual findings arising in secondary research. <i>Journal of Medical Ethics</i> , 2021, 47, 689-696.	1.8	4
8	Incremental value of left atrial booster and reservoir strain in predicting atrial fibrillation in patients with hypertrophic cardiomyopathy: a cardiovascular magnetic resonance study. <i>Journal of Cardiovascular Magnetic Resonance</i> , 2021, 23, 109.	3.3	14
9	Secondary findings in inherited heart conditions: a genotype-first feasibility study to assess phenotype, behavioural and psychosocial outcomes. <i>European Journal of Human Genetics</i> , 2020, 28, 1486-1496.	2.8	13
10	Whole-genome sequencing of patients with rare diseases in a national health system. <i>Nature</i> , 2020, 583, 96-102.	27.8	338
11	Reevaluation of the South Asian <i>MYBPC3</i> <sup>25bp</sup> Intronic Deletion in Hypertrophic Cardiomyopathy. <i>Circulation Genomic and Precision Medicine</i> , 2020, 13, e002783.	3.6	31
12	Implementation of a genomic medicine multi-disciplinary team approach for rare disease in the clinical setting: a prospective exome sequencing case series. <i>Genome Medicine</i> , 2019, 11, 46.	8.2	25
13	Do health professionals value genomic testing? A discrete choice experiment in inherited cardiovascular disease. <i>European Journal of Human Genetics</i> , 2019, 27, 1639-1648.	2.8	11
14	RV function deteriorates earlier than LV function and predicts adverse cardiovascular outcomes. , 2019, , .		0
15	Analysis of 51 proposed hypertrophic cardiomyopathy genes from genome sequencing data in sarcomere negative cases has negligible diagnostic yield. <i>Genetics in Medicine</i> , 2019, 21, 1576-1584.	2.4	44
16	Views of rare disease participants in a UK whole-genome sequencing study towards secondary findings: a qualitative study. <i>European Journal of Human Genetics</i> , 2018, 26, 652-659.	2.8	30
17	Not pathogenic until proven otherwise: perspectives of UK clinical genomics professionals toward secondary findings in context of a Genomic Medicine Multidisciplinary Team and the 100,000 Genomes Project. <i>Genetics in Medicine</i> , 2018, 20, 320-328.	2.4	56
18	From Genotype to Phenotype. <i>Circulation Genomic and Precision Medicine</i> , 2018, 11, e002316.	3.6	8

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19	Distinct ECG Phenotypes Identified in Hypertrophic Cardiomyopathy Using Machine Learning Associate With Arrhythmic Risk Markers. <i>Frontiers in Physiology</i> , 2018, 9, 213.	2.8	57
20	Insights from early experience of a Rare Disease Genomic Medicine Multidisciplinary Team: a qualitative study. <i>European Journal of Human Genetics</i> , 2017, 25, 680-686.	2.8	24
21	Stakeholder views on secondary findings in whole-genome and whole-exome sequencing: a systematic review of quantitative and qualitative studies. <i>Genetics in Medicine</i> , 2017, 19, 283-293.	2.4	119
22	Exploring the potential duty of care in clinical genomics under UK law. <i>Medical Law International</i> , 2017, 17, 158-182.	1.1	17
23	Factors influencing success of clinical genome sequencing across a broad spectrum of disorders. <i>Nature Genetics</i> , 2015, 47, 717-726.	21.4	310
24	Pre-symptomatic genetic testing for inherited cardiac conditions: a qualitative exploration of psychosocial and ethical implications. <i>European Journal of Human Genetics</i> , 2014, 22, 88-93.	2.8	65
25	Attitudes to reproductive genetic testing in women who had a positive BRCA test before having children: a qualitative analysis. <i>European Journal of Human Genetics</i> , 2012, 20, 4-10.	2.8	68
26	Genome Wide Identification of Recessive Cancer Genes by Combinatorial Mutation Analysis. <i>PLoS ONE</i> , 2008, 3, e3380.	2.5	12
27	Molecular Cloning, cDNA Sequence, and Chromosomal Localization of the Human Phosphatidylinositol 3-Kinase p110 $\alpha$ (PIK3CA) Gene. <i>Genomics</i> , 1994, 24, 472-477.	2.9	107
28	Mapping of cosmid clones in Huntington's disease region of chromosome 4. <i>Somatic Cell and Molecular Genetics</i> , 1991, 17, 83-91.	0.7	46