

Julia Wynn

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4894981/publications.pdf>

Version: 2024-02-01

55
papers

1,931
citations

279798

23
h-index

276875

41
g-index

57
all docs

57
docs citations

57
times ranked

3351
citing authors

#	ARTICLE	IF	CITATIONS
1	Questioning the validity of clinically available breast cancer polygenic risk scores: comparison of two labs reveals discrepancies. <i>Familial Cancer</i> , 2022, 21, 125-127.	1.9	0
2	Do research participants share genomic screening results with family members?. <i>Journal of Genetic Counseling</i> , 2022, 31, 447-458.	1.6	12
3	Improving Recruitment for a Newborn Screening Pilot Study with Adaptations in Response to the COVID-19 Pandemic. <i>International Journal of Neonatal Screening</i> , 2022, 8, 23.	3.2	7
4	Being small for gestational age is not an independent risk factor for mortality in neonates with congenital diaphragmatic hernia: a multicenter study. <i>Journal of Perinatology</i> , 2022, , .	2.0	1
5	Information is power: The experiences, attitudes and needs of individuals who chose to have prenatal genomic sequencing for fetal anomalies. <i>Prenatal Diagnosis</i> , 2022, 42, 947-954.	2.3	3
6	Mother and Daughter Perspectives on Genetic Counseling and Testing of Adolescents for Hereditary Breast Cancer Risk. <i>Journal of Pediatrics</i> , 2022, 251, 113-119.e7.	1.8	3
7	Returning negative results from <scp>largeâ€scale</scp> genomic screening: Experiences from the <scp>eMERGE III</scp> network. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 508-516.	1.2	5
8	An electronic health record (EHR) log analysis shows limited clinician engagement with unsolicited genetic test results. <i>JAMIA Open</i> , 2021, 4, oaab014.	2.0	5
9	COVID contingencies: Early epicenter experiences of different genetics clinics at a New York City institution inform emergency adaptation strategies. <i>Journal of Genetic Counseling</i> , 2021, 30, 938-948.	1.6	9
10	Penetrance of Breast Cancer Susceptibility Genes from the eMERGE III Network. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab044.	2.9	14
11	Impact of Genetic Testing for Cardiomyopathy on Emotional Well-Being and Family Dynamics: A Study of Parents and Adolescents. <i>Circulation Genomic and Precision Medicine</i> , 2021, 14, e003189.	3.6	2
12	Rare and de novo variants in 827 congenital diaphragmatic hernia probands implicate LONP1 as candidate risk gene. <i>American Journal of Human Genetics</i> , 2021, 108, 1964-1980.	6.2	22
13	GeneLiFT: A novel test to facilitate rapid screening of genetic literacy in a diverse population undergoing genetic testing. <i>Journal of Genetic Counseling</i> , 2021, 30, 742-754.	1.6	16
14	The influence of genetics in congenital diaphragmatic hernia. <i>Seminars in Perinatology</i> , 2020, 44, 151169.	2.5	39
15	Comparative outcomes of right versus left congenital diaphragmatic hernia: A multicenter analysis. <i>Journal of Pediatric Surgery</i> , 2020, 55, 33-38.	1.6	22
16	Impact of patient education videos on genetic counseling outcomes after exome sequencing. <i>Patient Education and Counseling</i> , 2020, 103, 127-135.	2.2	18
17	Participant choices for return of genomic results in the eMERGE Network. <i>Genetics in Medicine</i> , 2020, 22, 1821-1829.	2.4	25
18	Likely damaging de novo variants in congenital diaphragmatic hernia patients are associated with worse clinical outcomes. <i>Genetics in Medicine</i> , 2020, 22, 2020-2028.	2.4	21

#	ARTICLE	IF	CITATIONS
19	Deep whole-genome sequencing of multiple proband tissues and parental blood reveals the complex genetic etiology of congenital diaphragmatic hernias. <i>Human Genetics and Genomics Advances</i> , 2020, 1, 100008.	1.7	5
20	Advancing the genetic counseling profession through research: Identification of priorities by the National Society of Genetic Counselors research task force. <i>Journal of Genetic Counseling</i> , 2020, 29, 884-887.	1.6	9
21	Understanding the Return of Genomic Sequencing Results Process: Content Review of Participant Summary Letters in the eMERGE Research Network. <i>Journal of Personalized Medicine</i> , 2020, 10, 38.	2.5	15
22	A qualitative study of Latinx parents' experiences of clinical exome sequencing. <i>Journal of Genetic Counseling</i> , 2020, 29, 574-586.	1.6	16
23	Choices, attitudes, and experiences of genetic screening in Latino/a and Ashkenazi Jewish individuals. <i>Journal of Community Genetics</i> , 2020, 11, 391-403.	1.2	4
24	Returning Results in the Genomic Era: Initial Experiences of the eMERGE Network. <i>Journal of Personalized Medicine</i> , 2020, 10, 30.	2.5	39
25	Ethical conflicts in translational genetic research: lessons learned from the eMERGE-III experience. <i>Genetics in Medicine</i> , 2020, 22, 1667-1672.	2.4	10
26	Harmonizing Clinical Sequencing and Interpretation for the eMERGE III Network. <i>American Journal of Human Genetics</i> , 2019, 105, 588-605.	6.2	99
27	Psychological outcomes related to exome and genome sequencing result disclosure: a meta-analysis of seven Clinical Sequencing Exploratory Research (CSER) Consortium studies. <i>Genetics in Medicine</i> , 2019, 21, 2781-2790.	2.4	55
28	Evaluation of the cost and effectiveness of diverse recruitment methods for a genetic screening study. <i>Genetics in Medicine</i> , 2019, 21, 2371-2380.	2.4	10
29	Developing effective and efficient genomic educational tools for our diverse population. <i>Annals of Translational Medicine</i> , 2019, 7, S304-S304.	1.7	1
30	User engagement with web-based genomics education videos and implications for designing scalable patient education materials. <i>AMIA ... Annual Symposium proceedings</i> , 2019, 2019, 923-932.	0.2	0
31	Why Patients Decline Genomic Sequencing Studies: Experiences from the CSER Consortium. <i>Journal of Genetic Counseling</i> , 2018, 27, 1220-1227.	1.6	36
32	Diagnostic exome sequencing in children: A survey of parental understanding, experience and psychological impact. <i>Clinical Genetics</i> , 2018, 93, 1039-1048.	2.0	41
33	Examining the Psychosocial Impact of Genetic Testing for Cardiomyopathies. <i>Journal of Genetic Counseling</i> , 2018, 27, 927-934.	1.6	16
34	Impact of Receiving Secondary Results from Genomic Research: A 12-Month Longitudinal Study. <i>Journal of Genetic Counseling</i> , 2018, 27, 709-722.	1.6	26
35	De novo variants in congenital diaphragmatic hernia identify MYRF as a new syndrome and reveal genetic overlaps with other developmental disorders. <i>PLoS Genetics</i> , 2018, 14, e1007822.	3.5	79
36	Ethical Considerations Related to Return of Results from Genomic Medicine Projects: The eMERGE Network (Phase III) Experience. <i>Journal of Personalized Medicine</i> , 2018, 8, 2.	2.5	44

#	ARTICLE	IF	CITATIONS
37	Parental perceptions of prenatal whole exome sequencing (PPPWES) study. <i>Prenatal Diagnosis</i> , 2018, 38, 801-811.	2.3	31
38	Clinical providers'™ experiences with returning results from genomic sequencing: an interview study. <i>BMC Medical Genomics</i> , 2018, 11, 45.	1.5	55
39	Mutations in <i>BMPR2</i> are not present in patients with pulmonary hypertension associated with congenital diaphragmatic hernia. <i>Journal of Pediatric Surgery</i> , 2017, 52, 1747-1750.	1.6	3
40	Research Participants'™ Preferences for Hypothetical Secondary Results from Genomic Research. <i>Journal of Genetic Counseling</i> , 2017, 26, 841-851.	1.6	39
41	Loss of function in <i>ROBO1</i> is associated with tetralogy of Fallot and septal defects. <i>Journal of Medical Genetics</i> , 2017, 54, 825-829.	3.2	27
42	23andMe Paves the Way for Direct-to-Consumer Genetic Health Risk Tests of Limited Clinical Utility. <i>Annals of Internal Medicine</i> , 2017, 167, 125.	3.9	13
43	Impact of Panel Gene Testing for Hereditary Breast and Ovarian Cancer on Patients. <i>Journal of Genetic Counseling</i> , 2017, 26, 1116-1129.	1.6	90
44	Genome-wide enrichment of damaging de novo variants in patients with isolated and complex congenital diaphragmatic hernia. <i>Human Genetics</i> , 2017, 136, 679-691.	3.8	53
45	Congenital diaphragmatic hernias: from genes to mechanisms to therapies. <i>DMM Disease Models and Mechanisms</i> , 2017, 10, 955-970.	2.4	143
46	Clinical Sequencing Exploratory Research Consortium: Accelerating Evidence-Based Practice of Genomic Medicine. <i>American Journal of Human Genetics</i> , 2016, 98, 1051-1066.	6.2	137
47	Mutations in <i>TKT</i> Are the Cause of a Syndrome Including Short Stature, Developmental Delay, and Congenital Heart Defects. <i>American Journal of Human Genetics</i> , 2016, 98, 1235-1242.	6.2	31
48	Genomic Testing: a Genetic Counselor's Personal Reflection on Three Years of Consenting and Testing. <i>Journal of Genetic Counseling</i> , 2016, 25, 691-697.	1.6	20
49	Brief Report: <i>SETD2</i> Mutation in a Child with Autism, Intellectual Disabilities and Epilepsy. <i>Journal of Autism and Developmental Disorders</i> , 2015, 45, 3764-3770.	2.7	64
50	Association of Researcher Characteristics with Views on Return of Incidental Findings from Genomic Research. <i>Journal of Genetic Counseling</i> , 2015, 24, 833-841.	1.6	17
51	Illustrative case studies in the return of exome and genome sequencing results. <i>Personalized Medicine</i> , 2015, 12, 283-295.	1.5	35
52	Genetic causes of congenital diaphragmatic hernia. <i>Seminars in Fetal and Neonatal Medicine</i> , 2014, 19, 324-330.	2.3	77
53	The usefulness of whole-exome sequencing in routine clinical practice. <i>Genetics in Medicine</i> , 2014, 16, 922-931.	2.4	196
54	Developmental outcomes of children with congenital diaphragmatic hernia: A multicenter prospective study. <i>Journal of Pediatric Surgery</i> , 2013, 48, 1995-2004.	1.6	68

#	ARTICLE	IF	CITATIONS
55	Researchers'™ views on return of incidental genomic research results: qualitative and quantitative findings. <i>Genetics in Medicine</i> , 2013, 15, 888-895.	2.4	103