

David A Fasel

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2785076/publications.pdf>

Version: 2024-02-01

19
papers

1,889
citations

567281

15
h-index

713466

21
g-index

22
all docs

22
docs citations

22
times ranked

3407
citing authors

#	ARTICLE	IF	CITATIONS
1	Do research participants share genomic screening results with family members?. <i>Journal of Genetic Counseling</i> , 2022, 31, 447-458.	1.6	12
2	An electronic health record (EHR) log analysis shows limited clinician engagement with unsolicited genetic test results. <i>JAMIA Open</i> , 2021, 4, ooab014.	2.0	5
3	Copy Number Variant Analysis and Genome-wide Association Study Identify Loci with Large Effect for Vesicoureteral Reflux. <i>Journal of the American Society of Nephrology: JASN</i> , 2021, 32, 805-820.	6.1	17
4	Medical records-based chronic kidney disease phenotype for clinical care and "big data" observational and genetic studies. <i>Npj Digital Medicine</i> , 2021, 4, 70.	10.9	39
5	Genomic considerations for FHIR®; eMERGE implementation lessons. <i>Journal of Biomedical Informatics</i> , 2021, 118, 103795.	4.3	15
6	Generalizability of Polygenic Risk Scores for Breast Cancer Among Women With European, African, and Latinx Ancestry. <i>JAMA Network Open</i> , 2021, 4, e2119084.	5.9	31
7	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
8	Type IV Collagen Mutations in Familial IgA Nephropathy. <i>Kidney International Reports</i> , 2020, 5, 1075-1078.	0.8	26
9	Rapid exome sequencing in PICU patients with new-onset metabolic or neurological disorders. <i>Pediatric Research</i> , 2020, 88, 761-768.	2.3	19
10	Harmonizing Clinical Sequencing and Interpretation for the eMERGE III Network. <i>American Journal of Human Genetics</i> , 2019, 105, 588-605.	6.2	99
11	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
12	Diagnostic Utility of Exome Sequencing for Kidney Disease. <i>New England Journal of Medicine</i> , 2019, 380, 142-151.	27.0	456
13	The copy number variation landscape of congenital anomalies of the kidney and urinary tract. <i>Nature Genetics</i> , 2019, 51, 117-127.	21.4	144
14	Whole-Exome Sequencing in Adults With Chronic Kidney Disease. <i>Annals of Internal Medicine</i> , 2018, 168, 100.	3.9	154
15	Genetic Drivers of Kidney Defects in the DiGeorge Syndrome. <i>New England Journal of Medicine</i> , 2017, 376, 742-754.	27.0	120
16	Exome-wide Association Study Identifies GREB1L Mutations in Congenital Kidney Malformations. <i>American Journal of Human Genetics</i> , 2017, 101, 789-802.	6.2	63
17	Genomic imbalances in pediatric patients with chronic kidney disease. <i>Journal of Clinical Investigation</i> , 2015, 125, 2171-2178.	8.2	68
18	Copy number variation analysis identifies novel CAKUT candidate genes in children with a solitary functioning kidney. <i>Kidney International</i> , 2015, 88, 1402-1410.	5.2	65

#	ARTICLE	IF	CITATIONS
19	Discovery of new risk loci for IgA nephropathy implicates genes involved in immunity against intestinal pathogens. <i>Nature Genetics</i> , 2014, 46, 1187-1196.	21.4	505