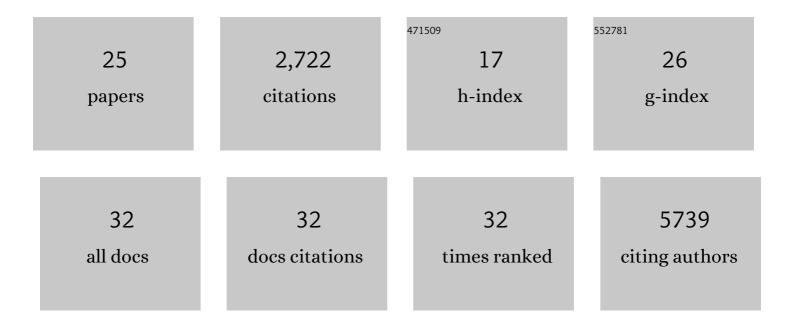
Erin R Riggs

List of Publications by Year in descending order

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FRIN P RICCS

| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 1 | The Human Phenotype Ontology project: linking molecular biology and disease through phenotype data. Nucleic Acids Research, 2014, 42, D966-D974. | 14.5 | 698 |
| 2 | Evaluating the Clinical Validity of Gene-Disease Associations: An Evidence-Based Framework Developed by the Clinical Genome Resource. American Journal of Human Genetics, 2017, 100, 895-906. | 6.2 | 403 |
| 3 | The Monarch Initiative in 2019: an integrative data and analytic platform connecting phenotypes to genotypes across species. Nucleic Acids Research, 2020, 48, D704-D715. | 14.5 | 178 |
| 4 | Mitochondrial Disease Sequence Data Resource (MSeqDR): A global grass-roots consortium to facilitate deposition, curation, annotation, and integrated analysis of genomic data for the mitochondrial disease clinical and research communities. Molecular Genetics and Metabolism, 2015, 114, 388-396. | 1.1 | 76 |
| 5 | GenomeConnect: Matchmaking Between Patients, Clinical Laboratories, and Researchers to Improve Genomic Knowledge. Human Mutation, 2015, 36, 974-978. | 2.5 | 56 |
| 6 | The Gene Curation Coalition: A global effort to harmonize gene–disease evidence resources. Genetics in Medicine, 2022, 24, 1732-1742. | 2.4 | 56 |
| 7 | Phenotypic information in genomic variant databases enhances clinical care and research: The international standards for cytogenomic arrays consortium experience. Human Mutation, 2012, 33, 787-796. | 2.5 | 49 |
| 8 | ClinGen Variant Curation Interface: a variant classification platform for the application of evidence criteria from ACMG/AMP guidelines. Genome Medicine, 2022, 14, 6. | 8.2 | 34 |
| 9 | Incorporating Social Media into your Support Tool Box: Points to Consider from Geneticsâ€Based Communities. Journal of Genetic Counseling, 2018, 27, 470-480. | 1.6 | 33 |
| 10 | The ClinGen Epilepsy Gene Curation Expert Panel—Bridging the divide between clinical domain knowledge and formal gene curation criteria. Human Mutation, 2018, 39, 1476-1484. | 2.5 | 33 |
| 11 | ClinGen's GenomeConnect registry enables patientâ€centered data sharing. Human Mutation, 2018, 39, 1668-1676. | 2.5 | 25 |
| 12 | Points to consider for sharing variant-level information from clinical genetic testing with ClinVar. Journal of Physical Education and Sports Management, 2018, 4, a002345. | 1.2 | 23 |
| 13 | Copy number variant discrepancy resolution using the ClinGen dosage sensitivity map results in updated clinical interpretations in ClinVar. Human Mutation, 2018, 39, 1650-1659. | 2.5 | 23 |
| 14 | Development of a consent resource for genomic data sharing in the clinical setting. Genetics in Medicine, 2019, 21, 81-88. | 2.4 | 20 |
| 15 | Utilizing ClinGen geneâ€disease validity and dosage sensitivity curations to inform variant classification. Human Mutation, 2022, 43, 1031-1040. | 2.5 | 20 |
| 16 | Development of Clinical Domain Working Groups for the Clinical Genome Resource (ClinGen): lessons learned and plans for the future. Genetics in Medicine, 2019, 21, 987-993. | 2.4 | 17 |
| 17 | Ensuring best practice in genomics education and evaluation: reporting item standards for education and its evaluation in genomics (RISE2 Genomics). Genetics in Medicine, 2021, 23, 1356-1365. | 2.4 | 17 |
| 18 | Genomic Variation: Lessons Learned from Whole-Genome CNV Analysis. Current Genetic Medicine Reports, 2014, 2, 146-150. | 1.9 | 16 |

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|----|--|-----|-----------|
| 19 | Variant interpretation is a component of clinical practice among genetic counselors in multiple specialties. Genetics in Medicine, 2020, 22, 785-792. | 2.4 | 14 |
| 20 | The Laboratory linician Team: A Professional Call to Action to Improve Communication and Collaboration for Optimal Patient Care in Chromosomal Microarray Testing. Journal of Genetic Counseling, 2012, 21, 631-637. | 1.6 | 12 |
| 21 | Clinical validity assessment of genes frequently tested on intellectual disability/autism sequencing panels. Genetics in Medicine, 2022, 24, 1899-1908. | 2.4 | 9 |
| 22 | Recontacting registry participants with genetic updates through GenomeConnect, the ClinGen patient registry. Genetics in Medicine, 2021, 23, 1738-1745. | 2.4 | 7 |
| 23 | Response to Maya et al Genetics in Medicine, 2020, 22, 1278-1279. | 2.4 | 4 |
| 24 | Utilizing population-based genomic data to expedite the curation of genes and genomic regions for the ClinGen "dosage sensitivity unlikely―classification. Molecular Genetics and Metabolism, 2021, 132, S228-S229. | 1.1 | 0 |
| 25 | OP029: Genetic updates returned by GenomeConnect, the ClinGen patient registry: A pilot study of participant experience. Genetics in Medicine, 2022, 24, S357-S358. | 2.4 | Ο |