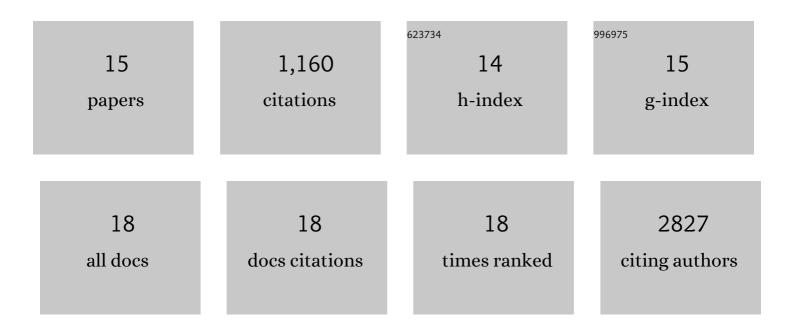
## Erika M Kvikstad

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3702908/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Advancing human genetics research and drug discovery through exome sequencing of the UK Biobank. Nature Genetics, 2021, 53, 942-948.	21.4	234
2	Pan-ancestry exome-wide association analyses of COVID-19 outcomes in 586,157 individuals. American Journal of Human Genetics, 2021, 108, 1350-1355.	6.2	72
3	The complete costs of genome sequencing: a microcosting study in cancer and rare diseases from a single center in the United Kingdom. Genetics in Medicine, 2020, 22, 85-94.	2.4	133
4	Clinically actionable mutation profiles in patients with cancer identified by whole-genome sequencing. Journal of Physical Education and Sports Management, 2018, 4, a002279.	1.2	21
5	A high throughput screen for active human transposable elements. BMC Genomics, 2018, 19, 115.	2.8	14
6	Identification of a new VHL exon and complex splicing alterations in familial erythrocytosis or von Hippel-Lindau disease. Blood, 2018, 132, 469-483.	1.4	70
7	Strong Heterogeneity in Mutation Rate Causes Misleading Hallmarks of Natural Selection on Indel Mutations in the Human Genome. Molecular Biology and Evolution, 2014, 31, 23-36.	8.9	16
8	The origin, evolution, and functional impact of short insertion–deletion variants identified in 179 human genomes. Genome Research, 2013, 23, 749-761.	5.5	206
9	The (r)evolution of SINE versus LINE distributions in primate genomes: Sex chromosomes are important. Genome Research, 2010, 20, 600-613.	5.5	48
10	Ride the wavelet: A multiscale analysis of genomic contexts flanking small insertions and deletions. Genome Research, 2009, 19, 1153-1164.	5.5	27
11	A Macaque's-Eye View of Human Insertions and Deletions: Differences in Mechanisms. PLoS Computational Biology, 2007, 3, e176.	3.2	55
12	Single-Molecule Approach to Bacterial Genomic Comparisons via Optical Mapping. Journal of Bacteriology, 2004, 186, 7773-7782.	2.2	63
13	Shotgun optical mapping of the entire Leishmania major Friedlin genome. Molecular and Biochemical Parasitology, 2004, 138, 97-106.	1.1	41
14	Whole-Genome Shotgun Optical Mapping of Rhodobacter sphaeroides strain 2.4.1 and Its Use for Whole-Genome Shotgun Sequence Assembly. Genome Research, 2003, 13, 2142-2151.	5.5	49
15	A Whole-Genome Shotgun Optical Map of Yersinia pestis Strain KIM. Applied and Environmental Microbiology, 2002, 68, 6321-6331.	3.1	65