

# Liis Kolberg

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2451282/publications.pdf>

Version: 2024-02-01

11  
papers

5,259  
citations

1162367

8  
h-index

1372195

10  
g-index

14  
all docs

14  
docs citations

14  
times ranked

11606  
citing authors

#	ARTICLE	IF	CITATIONS
1	g:Profiler: a web server for functional enrichment analysis and conversions of gene lists (2019) Tj ETQq1 1 0.784314 rgBT /Overlock 10	6.5	3,302
2	g:Profiler—a web server for functional interpretation of gene lists (2016 update). Nucleic Acids Research, 2016, 44, W83-W89.	6.5	1,179
3	gprofiler2 – an R package for gene list functional enrichment analysis and namespace conversion toolset g:Profiler. F1000Research, 2020, 9, 709.	0.8	331
4	A compendium of uniformly processed human gene expression and splicing quantitative trait loci. Nature Genetics, 2021, 53, 1290-1299.	9.4	193
5	gprofiler2 – an R package for gene list functional enrichment analysis and namespace conversion toolset g:Profiler. F1000Research, 2020, 9, 709.	0.8	128
6	Endometrial transcriptome analysis indicates superiority of natural over artificial cycles in recurrent implantation failure patients undergoing frozen embryo transfer. Reproductive BioMedicine Online, 2016, 32, 597-613.	1.1	38
7	Co-expression analysis reveals interpretable gene modules controlled by trans-acting genetic variants. ELife, 2020, 9, .	2.8	24
8	An exploratory phenome wide association study linking asthma and liver disease genetic variants to electronic health records from the Estonian Biobank. PLoS ONE, 2019, 14, e0215026.	1.1	8
9	funcExplorer: a tool for fast data-driven functional characterisation of high-throughput expression data. BMC Genomics, 2018, 19, 817.	1.2	5
10	The 1st year of the COVID-19 epidemic in Estonia: a population-based nationwide sequential/consecutive cross-sectional study. Public Health, 2022, 205, 150-156.	1.4	4
11	Comparison of variation in frequency for SNPs associated with asthma or liver disease between Estonia, HapMap populations and the 1000 genome project populations. International Journal of Immunogenetics, 2019, 46, 49-58.	0.8	0