

# Valborg Gudmundsdottir

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5965096/publications.pdf>

Version: 2024-02-01

21  
papers

4,570  
citations

623734

14  
h-index

677142

22  
g-index

29  
all docs

29  
docs citations

29  
times ranked

9932  
citing authors

#	ARTICLE	IF	CITATIONS
1	Proteomic Analysis Identifies Circulating Proteins Associated With Plasma Amyloid- $\beta^2$ and Incident Dementia. <i>Biological Psychiatry Global Open Science</i> , 2023, 3, 490-499.	2.2	5
2	A genome-wide association study of serum proteins reveals shared loci with common diseases. <i>Nature Communications</i> , 2022, 13, 480.	12.8	79
3	Four groups of type 2 diabetes contribute to the etiological and clinical heterogeneity in newly diagnosed individuals: An IMI DIRECT study. <i>Cell Reports Medicine</i> , 2022, 3, 100477.	6.5	39
4	Coding and regulatory variants are associated with serum protein levels and disease. <i>Nature Communications</i> , 2022, 13, 481.	12.8	18
5	The Proteomic Profile of Interstitial Lung Abnormalities. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2022, 206, 337-346.	5.6	7
6	A proteogenomic signature of age-related macular degeneration in blood. <i>Nature Communications</i> , 2022, 13, .	12.8	14
7	Itâ€™s in Our Blood: A Glimpse of Personalized Medicine. <i>Trends in Molecular Medicine</i> , 2021, 27, 20-30.	6.7	26
8	Serum levels of ACE2 are higher in patients with obesity and diabetes. <i>Obesity Science and Practice</i> , 2021, 7, 239-243.	1.9	20
9	Multiethnic Genome-Wide Association Study of Subclinical Atherosclerosis in Individuals With Type 2 Diabetes. <i>Circulation Genomic and Precision Medicine</i> , 2021, 14, e003258.	3.6	4
10	Whole blood co-expression modules associate with metabolic traits and type 2 diabetes: an IMI-DIRECT study. <i>Genome Medicine</i> , 2020, 12, 109.	8.2	8
11	Circulating Protein Signatures and Causal Candidates for Type 2 Diabetes. <i>Diabetes</i> , 2020, 69, 1843-1853.	0.6	64
12	Metabolite ratios as potential biomarkers for type 2 diabetes: a DIRECT study. <i>Diabetologia</i> , 2018, 61, 117-129.	6.3	32
13	A computational framework to integrate high-throughput omics datasets for the identification of potential mechanistic links. <i>Nature Protocols</i> , 2018, 13, 2781-2800.	12.0	82
14	Co-regulatory networks of human serum proteins link genetics to disease. <i>Science</i> , 2018, 361, 769-773.	12.6	375
15	Integrative network analysis highlights biological processes underlying GLP-1 stimulated insulin secretion: A DIRECT study. <i>PLoS ONE</i> , 2018, 13, e0189886.	2.5	9
16	Early differences in islets from prediabetic NOD mice: combined microarray and proteomic analysis. <i>Diabetologia</i> , 2017, 60, 475-489.	6.3	31
17	Pancreatic Islet Protein Complexes and Their Dysregulation in Type 2 Diabetes. <i>Frontiers in Genetics</i> , 2017, 8, 43.	2.3	19
18	Human gut microbes impact host serum metabolome and insulin sensitivity. <i>Nature</i> , 2016, 535, 376-381.	27.8	1,506

#	ARTICLE	IF	CITATIONS
19	A proteomic study of the regulatory role for STAT $\alpha$ 1 in cytokine-induced beta cell death. <i>Proteomics - Clinical Applications</i> , 2015, 9, 938-952.	1.6	4
20	Disentangling type 2 diabetes and metformin treatment signatures in the human gut microbiota. <i>Nature</i> , 2015, 528, 262-266.	27.8	1,627
21	The genome of a Late Pleistocene human from a Clovis burial site in western Montana. <i>Nature</i> , 2014, 506, 225-229.	27.8	500