

# Tiina Skoog

## List of Publications by Year in descending order

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Version: 2024-02-01

31  
papers

1,445  
citations

393982

19  
h-index

476904

29  
g-index

32  
all docs

32  
docs citations

32  
times ranked

3016  
citing authors

#	ARTICLE	IF	CITATIONS
1	A Common Functional Polymorphism (C->A Substitution at Position -863) in the Promoter Region of the Tumour Necrosis Factor- $\alpha$ (TNF- $\alpha$ ) Gene Associated With Reduced Circulating Levels of TNF- $\alpha$ . <i>Human Molecular Genetics</i> , 1999, 8, 1443-1449.	1.4	307
2	Plasma tumour necrosis factor- $\alpha$ and early carotid atherosclerosis in healthy middle-aged men. <i>European Heart Journal</i> , 2002, 23, 376-383.	1.0	224
3	Microbe-host interplay in atopic dermatitis and psoriasis. <i>Nature Communications</i> , 2019, 10, 4703.	5.8	217
4	The human GIMAP5 gene has a common polyadenylation polymorphism increasing risk to systemic lupus erythematosus. <i>Journal of Medical Genetics</i> , 2007, 44, 314-321.	1.5	70
5	NOD-like receptor signaling and inflammasome-related pathways are highlighted in psoriatic epidermis. <i>Scientific Reports</i> , 2016, 6, 22745.	1.6	63
6	Cationic gold nanoparticles elicit mitochondrial dysfunction: a multi-omics study. <i>Scientific Reports</i> , 2019, 9, 4366.	1.6	54
7	Matrilysin-2 (Matrix Metalloproteinase-26) Is Upregulated in Keratinocytes During Wound Repair and Early Skin Carcinogenesis. <i>Journal of Investigative Dermatology</i> , 2005, 124, 849-856.	0.3	52
8	Increased expression of matrix metalloproteinases-21 and -26 and TIMP-4 in pancreatic adenocarcinoma. <i>Modern Pathology</i> , 2007, 20, 1128-1140.	2.9	43
9	Aberrant splicing of genes involved in haemoglobin synthesis and impaired terminal erythroid maturation in <i>SF3B1</i> mutated refractory anaemia with ring sideroblasts. <i>British Journal of Haematology</i> , 2015, 171, 478-490.	1.2	37
10	CCHCR1 Is Up-Regulated in Skin Cancer and Associated with EGFR Expression. <i>PLoS ONE</i> , 2009, 4, e6030.	1.1	33
11	MMP-21 is expressed by macrophages and fibroblasts in vivo and in culture. <i>Experimental Dermatology</i> , 2006, 15, 775-783.	1.4	29
12	Fetal HLA-G mediated immune tolerance and interferon response in preeclampsia. <i>EBioMedicine</i> , 2020, 59, 102872.	2.7	25
13	MMP-10 (Stromelysin) and MMP-21 in human and murine squamous cell cancer. <i>Experimental Dermatology</i> , 2009, 18, 1044-1052.	1.4	24
14	Multiparametric Profiling of Engineered Nanomaterials: Unmasking the Surface Coating Effect. <i>Advanced Science</i> , 2020, 7, 2002221.	5.6	24
15	Tumour necrosis factor- $\alpha$ (TNF- $\alpha$ ) polymorphisms -857C/A and -863C/A are associated with TNF- $\alpha$ secretion from human adipose tissue. <i>Diabetologia</i> , 2001, 44, 654-655.	2.9	22
16	Expression of MMP-10, MMP-21, MMP-26, and MMP-28 in Merkel cell carcinoma. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2009, 455, 495-503.	1.4	22
17	The human long non-coding RNA gene RMRP has pleiotropic effects and regulates cell-cycle progression at G2. <i>Scientific Reports</i> , 2019, 9, 13758.	1.6	22
18	Gene expression analysis of skin grafts and cultured keratinocytes using synthetic RNA normalization reveals insights into differentiation and growth control. <i>BMC Genomics</i> , 2015, 16, 476.	1.2	21

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19	Allele-specific chromatin remodeling of the tumor necrosis factor- $\beta$ promoter. <i>Biochemical and Biophysical Research Communications</i> , 2006, 351, 777-783.	1.0	20
20	Expression of Allograft Inflammatory Factor-1 in Inflammatory Skin Disorders. <i>Acta Dermato-Venereologica</i> , 2007, 87, 223-227.	0.6	20
21	Intracellular signalling pathways and cytoskeletal functions converge on the psoriasis candidate gene CCHCR1 expressed at P-bodies and centrosomes. <i>BMC Genomics</i> , 2018, 19, 432.	1.2	17
22	Delineating the Healthy Human Skin UV Response and Early Induction of Interferon Pathway in Cutaneous Lupus Erythematosus. <i>Journal of Investigative Dermatology</i> , 2019, 139, 2058-2061.e4.	0.3	16
23	Biomarkers of nanomaterials hazard from multi-layer data. <i>Nature Communications</i> , 2022, 13, .	5.8	16
24	Toxicogenomic Profiling of 28 Nanomaterials in Mouse Airways. <i>Advanced Science</i> , 2021, 8, 2004588.	5.6	15
25	Identification of MAMDC1 as a Candidate Susceptibility Gene for Systemic Lupus Erythematosus (SLE). <i>PLoS ONE</i> , 2009, 4, e8037.	1.1	14
26	Matrix Metalloproteinase-21 Expression Is Associated with Keratinocyte Differentiation and Upregulated by Retinoic Acid in HaCaT Cells. <i>Journal of Investigative Dermatology</i> , 2009, 129, 119-130.	0.3	13
27	Investigation of rare and low-frequency variants using high-throughput sequencing with pooled DNA samples. <i>Scientific Reports</i> , 2016, 6, 33256.	1.6	13
28	Guide for library design and bias correction for large-scale transcriptome studies using highly multiplexed RNAseq methods. <i>BMC Bioinformatics</i> , 2019, 20, 418.	1.2	9
29	Sequence analysis of pooled bacterial samples enables identification of strain variation in group A streptococcus. <i>Scientific Reports</i> , 2017, 7, 45771.	1.6	3
30	Erythropoiesis In SF3B1 Mutated RARS Is Disrupted During Terminal Erythroid Maturation. <i>Blood</i> , 2013, 122, 2408-2408.	0.6	0
31	INFLUENCE OF FLG LOSS-OF-FUNCTION MUTATIONS IN HOST-MICROBE INTERACTIONS DURING ATOPIC SKIN INFLAMMATION. <i>Journal of Dermatological Science</i> , 2022, , .	1.0	0