## Tessel E Galesloot

List of Publications by Year in descending order

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172457 138484 6,224 54 29 58 citations h-index g-index papers 61 61 61 13307 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Genome-wide Meta-analysis Identifies Novel Genes Associated with Recurrence and Progression in Non–muscle-invasive Bladder Cancer. European Urology Oncology, 2022, 5, 70-83.	5.4	5
2	Transferrin Saturation/Hepcidin Ratio Discriminates TMPRSS6-Related Iron Refractory Iron Deficiency Anemia from Patients with Multi-Causal Iron Deficiency Anemia. International Journal of Molecular Sciences, 2022, 23, 1917.	4.1	4
3	Polygenic prediction of educational attainment within and between families from genome-wide association analyses in 3 million individuals. Nature Genetics, 2022, 54, 437-449.	21.4	215
4	The Effects of Common Genetic Variation in 96 Genes Involved in Thyroid Hormone Regulation on TSH and FT4 Concentrations. Journal of Clinical Endocrinology and Metabolism, 2022, 107, e2276-e2283.	3.6	6
5	Semi-Quantitative Multiplex Profiling of the Complement System Identifies Associations of Complement Proteins with Genetic Variants and Metabolites in Age-Related Macular Degeneration. Journal of Personalized Medicine, 2021, 11, 1256.	2.5	5
6	The power of genetic diversity in genome-wide association studies of lipids. Nature, 2021, 600, 675-679.	27.8	353
7	Identification of ADHD risk genes in extended pedigrees by combining linkage analysis and whole-exome sequencing. Molecular Psychiatry, 2020, 25, 2047-2057.	7.9	17
8	Standardized serum hepcidin values in Dutch children: Set point relative to body iron changes during childhood. Pediatric Blood and Cancer, 2020, 67, e28038.	1.5	16
9	Cigarette Smoking and the Risk of Cutaneous Melanoma: A Case-Control Study. Dermatology, 2020, 236, 228-236.	2.1	21
10	Sex-Specific Regulation of Inflammation and Metabolic Syndrome in Obesity. Arteriosclerosis, Thrombosis, and Vascular Biology, 2020, 40, 1787-1800.	2.4	77
11	Integrating Metabolomics, Genomics, and Disease Pathways in Age-Related Macular Degeneration. Ophthalmology, 2020, 127, 1693-1709.	5.2	43
12	Measurement and genetic architecture of lifetime depression in the Netherlands as assessed by LIDAS (Lifetime Depression Assessment Self-report). Psychological Medicine, 2020, , 1-10.	4.5	4
13	ADHD symptoms in the adult general population are associated with factors linked to ADHD in adult patients. European Neuropsychopharmacology, 2019, 29, 1117-1126.	0.7	23
14	A Potential Role for the STXBP5-AS1 Gene in Adult ADHD Symptoms. Behavior Genetics, 2019, 49, 270-285.	2.1	6
15	Exome chip association study excluded the involvement of rare coding variants with large effect sizes in the etiology of anorectal malformations. PLoS ONE, 2019, 14, e0217477.	2.5	3
16	Genome-wide Association Study for Tumour Stage, Grade, Size, and Age at Diagnosis of Non–muscle-invasive Bladder Cancer. European Urology Oncology, 2019, 2, 381-389.	5 <b>.</b> 4	12
17	Non-Syndromic Cleft Lip with or without Cleft Palate: Genome-Wide Association Study in Europeans Identifies a Suggestive Risk Locus at 16p12.1 and Supports SH3PXD2A as a Clefting Susceptibility Gene. Genes, 2019, 10, 1023.	2.4	26
18	Genome-wide association analyses of risk tolerance and risky behaviors in over 1 million individuals identify hundreds of loci and shared genetic influences. Nature Genetics, 2019, 51, 245-257.	21.4	536

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19	Autism spectrum disorders and autistic traits share genetics and biology. Molecular Psychiatry, 2018, 23, 1205-1212.	7.9	125
20	GWAS and colocalization analyses implicate carotid intima-media thickness and carotid plaque loci in cardiovascular outcomes. Nature Communications, 2018, 9, 5141.	12.8	119
21	Genome-wide analyses identify a role for SLC17A4 and AADAT in thyroid hormone regulation. Nature Communications, 2018, 9, 4455.	12.8	181
22	Association of BRCA2 K3326* With Small Cell Lung Cancer and Squamous Cell Cancer of the Skin. Journal of the National Cancer Institute, 2018, 110, 967-974.	6.3	29
23	Exploring the role of low-frequency and rare exonic variants in alcohol and tobacco use. Drug and Alcohol Dependence, 2018, 188, 94-101.	3.2	10
24	Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. Nature Genetics, 2018, 50, 26-41.	21.4	286
25	Rare and low-frequency coding variants alter human adult height. Nature, 2017, 542, 186-190.	27.8	544
26	Cohort Profile: The Nijmegen Biomedical Study (NBS). International Journal of Epidemiology, 2017, 46, dyw268.	1.9	30
27	Meta-analysis identifies novel risk loci and yields systematic insights into the biology of male-pattern baldness. Nature Communications, 2017, 8, 14694.	12.8	58
28	GLRB allelic variation associated with agoraphobic cognitions, increased startle response and fear network activation: a potential neurogenetic pathway to panic disorder. Molecular Psychiatry, 2017, 22, 1431-1439.	7.9	47
29	Genome-wide meta-analysis associates HLA-DQA1/DRB1 and LPA and lifestyle factors with human longevity. Nature Communications, 2017, 8, 910.	12.8	118
30	Sequence variant at 4q25 near PITX2 associates with appendicitis. Scientific Reports, 2017, 7, 3119.	3.3	14
31	Meta-GWAS and Meta-Analysis of Exome Array Studies Do Not Reveal Genetic Determinants of Serum Hepcidin. PLoS ONE, 2016, 11, e0166628.	2.5	2
32	Variant <i>ASGR1</i> Associated with a Reduced Risk of Coronary Artery Disease. New England Journal of Medicine, 2016, 374, 2131-2141.	27.0	137
33	Genome-wide association study identifies 74 loci associated with educational attainment. Nature, 2016, 533, 539-542.	27.8	1,204
34	Cystatin C and Cardiovascular Disease. Journal of the American College of Cardiology, 2016, 68, 934-945.	2.8	109
35	Meta-analysis of the DRD5 VNTR in persistent ADHD. European Neuropsychopharmacology, 2016, 26, 1527-1532.	0.7	4
36	Exome chip analyses in adult attention deficit hyperactivity disorder. Translational Psychiatry, 2016, 6, e923-e923.	4.8	27

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37	Genetic variants linked to education predict longevity. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 13366-13371.	7.1	110
38	Identification of a novel susceptibility locus at 13q34 and refinement of the 20p12.2 region as a multi-signal locus associated with bladder cancer risk in individuals of European ancestry. Human Molecular Genetics, 2016, 25, 1203-1214.	2.9	38
39	Underestimation of hepcidin concentration by time of flight mass spectrometry and competitive ELISA in hepcidin p.Gly71Asp heterozygotes. Clinical Chemistry and Laboratory Medicine, 2016, 54, e173-6.	2.3	O
40	Implicit stigmatization-related biases in individuals with skin conditions and their significant others Health Psychology, 2016, 35, 861-865.	1.6	26
41	The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. PLoS Genetics, 2015, 11, e1005378.	3.5	331
42	Polygenic risk scores for schizophrenia and bipolar disorder predict creativity. Nature Neuroscience, 2015, 18, 953-955.	14.8	351
43	Known susceptibility SNPs for sporadic prostate cancer show a similar association with "hereditary― prostate cancer. Prostate, 2015, 75, 474-483.	2.3	12
44	Iron and hepcidin as risk factors in atherosclerosis: what do the genes say?. BMC Genetics, 2015, 16, 79.	2.7	23
45	A Comparison of Multivariate Genome-Wide Association Methods. PLoS ONE, 2014, 9, e95923.	2.5	168
46	Identification of Novel Genetic Loci Associated with Thyroid Peroxidase Antibodies and Clinical Thyroid Disease. PLoS Genetics, 2014, 10, e1004123.	3.5	150
47	Serum Hepcidin Is Associated With Presence of Plaque in Postmenopausal Women of a General Population. Arteriosclerosis, Thrombosis, and Vascular Biology, 2014, 34, 446-456.	2.4	40
48	Genome-wide association study yields variants at 20p12.2 that associate with urinary bladder cancer. Human Molecular Genetics, 2014, 23, 5545-5557.	2.9	46
49	The effect of the Taq1A variant in the dopamine D2 receptor gene and common CYP2D6 alleles on prolactin levels in risperidone-treated boys. Pharmacogenetics and Genomics, 2013, 23, 487-493.	1.5	32
50	Associations of common variants in <i>HFE</i> and <i>TMPRSS6</i> with iron parameters are independent of serum hepcidin in a general population: a replication study. Journal of Medical Genetics, 2013, 50, 593-598.	3.2	34
51	Exploration of Gene-Environment Interactions, Maternal Effects and Parent of Origin Effects in the Etiology of Hypospadias. Journal of Urology, 2012, 188, 2354-2360.	0.4	29
52	Candidate Gene Sequencing of SLC11A2 and TMPRSS6 in a Family with Severe Anaemia: Common SNPs, Rare Haplotypes, No Causative Mutation. PLoS ONE, 2012, 7, e35015.	2.5	21
53	Genomeâ€wide association uncovers shared genetic effects among personality traits and mood states. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2012, 159B, 684-695.	1.7	112
54	Serum hepcidin: reference ranges and biochemical correlates in the general population. Blood, 2011, 117, e218-e225.	1.4	246