Tatijana Zemunik

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Ancient human genomes suggest three ancestral populations for present-day Europeans. Nature, 2014, 513, 409-413.	27.8	1,179
2	Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits. Nature Genetics, 2018, 50, 1412-1425.	21.4	924
3	A genome-wide approach accounting for body mass index identifies genetic variants influencing fasting glycemic traits and insulin resistance. Nature Genetics, 2012, 44, 659-669.	21.4	762
4	Large-scale association analyses identify new loci influencing glycemic traits and provide insight into the underlying biological pathways. Nature Genetics, 2012, 44, 991-1005.	21.4	746
5	New loci associated with kidney function and chronic kidney disease. Nature Genetics, 2010, 42, 376-384.	21.4	710
6	Genome-wide association analyses identify 18 new loci associated with serum urate concentrations. Nature Genetics, 2013, 45, 145-154.	21.4	675
7	Genetic associations at 53 loci highlight cell types and biological pathways relevant for kidney function. Nature Communications, 2016, 7, 10023.	12.8	412
8	FTO genotype is associated with phenotypic variability of body mass index. Nature, 2012, 490, 267-272.	27.8	383
9	Genome-wide association and large-scale follow up identifies 16 new loci influencing lung function. Nature Genetics, 2011, 43, 1082-1090.	21.4	367
10	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. Nature Genetics, 2015, 47, 1294-1303.	21.4	357
11	The trans-ancestral genomic architecture of glycemic traits. Nature Genetics, 2021, 53, 840-860.	21.4	341
12	Meta-analyses identify 13 loci associated with age at menopause and highlight DNA repair and immune pathways. Nature Genetics, 2012, 44, 260-268.	21.4	303
13	Genome-Wide Association and Functional Follow-Up Reveals New Loci for Kidney Function. PLoS Genetics, 2012, 8, e1002584.	3.5	166
14	Meta-Analysis of Genome-Wide Association Studies Identifies Six New Loci for Serum Calcium Concentrations. PLoS Genetics, 2013, 9, e1003796.	3.5	142
15	Genome-wide association analysis identifies six new loci associated with forced vital capacity. Nature Genetics, 2014, 46, 669-677.	21.4	131
16	Multiethnic Meta-Analysis of Genome-Wide Association Studies in >100 000 Subjects Identifies 23 Fibrinogen-Associated Loci but No Strong Evidence of a Causal Association Between Circulating Fibrinogen and Cardiovascular Disease. Circulation, 2013, 128, 1310-1324.	1.6	128
17	Novel Blood Pressure Locus and Gene Discovery Using Genome-Wide Association Study and Expression Data Sets From Blood and the Kidney. Hypertension, 2017, 70,	2.7	123
18	Sixteen new lung function signals identified through 1000 Genomes Project reference panel imputation. Nature Communications, 2015, 6, 8658.	12.8	108

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19	"10 001 Dalmatians:―Croatia Launches Its National Biobank. Croatian Medical Journal, 2009, 50, 4-6.	0.7	99
20	Association Between Chromosome 9p21 Variants and the Ankle-Brachial Index Identified by a Meta-Analysis of 21 Genome-Wide Association Studies. Circulation: Cardiovascular Genetics, 2012, 5, 100-112.	5.1	98
21	Ethical aspects of human biobanks: a systematic review. Croatian Medical Journal, 2011, 52, 262-279.	0.7	95
22	Sex-dimorphic genetic effects and novel loci for fasting glucose and insulin variability. Nature Communications, 2021, 12, 24.	12.8	87
23	The peopling of Europe and the cautionary tale of Y chromosome lineage R-M269. Proceedings of the Royal Society B: Biological Sciences, 2012, 279, 884-892.	2.6	84
24	Evidence of Inbreeding Depression on Human Height. PLoS Genetics, 2012, 8, e1002655.	3.5	79
25	Polymorphisms in B3GAT1, SLC9A9 and MGAT5 are associated with variation within the human plasma N-glycome of 3533 European adults. Human Molecular Genetics, 2011, 20, 5000-5011.	2.9	74
26	Environmental Factors Affecting Thyroid-Stimulating Hormone and Thyroid Hormone Levels. International Journal of Molecular Sciences, 2021, 22, 6521.	4.1	74
27	A meta-analysis of 120 246 individuals identifies 18 new loci for fibrinogen concentration. Human Molecular Genetics, 2016, 25, 358-370.	2.9	73
28	Hearing function and thresholds: a genome-wide association study in European isolated populations identifies new loci and pathways. Journal of Medical Genetics, 2011, 48, 369-374.	3.2	71
29	Discovery and Fine Mapping of Serum Protein Loci through Transethnic Meta-analysis. American Journal of Human Genetics, 2012, 91, 744-753.	6.2	69
30	The Role of Recent Admixture in Forming the Contemporary West Eurasian Genomic Landscape. Current Biology, 2015, 25, 2518-2526.	3.9	68
31	Integration of genome-wide association studies with biological knowledge identifies six novel genes related to kidney function. Human Molecular Genetics, 2012, 21, 5329-5343.	2.9	64
32	Vitamin D receptor polymorphism and susceptibility to type 1 diabetes in the Dalmatian population. Diabetes Research and Clinical Practice, 2003, 59, 31-35.	2.8	63
33	Genome-wide Association Study of Biochemical Traits in KorÄula Island, Croatia. Croatian Medical Journal, 2009, 50, 23-33.	0.7	49
34	A meta-analysis of genome-wide data from five European isolates reveals an association of COL22A1, SYT1, and GABRR2with serum creatinine level. BMC Medical Genetics, 2010, 11, 41.	2.1	48
35	Fokl Polymorphism, Vitamin D Receptor, and Interleukin-1 Receptor Haplotypes Are Associated with Type 1 Diabetes in the Dalmatian Population. Journal of Molecular Diagnostics, 2005, 7, 600-604.	2.8	41
36	Prognosis in monoclonal gammopathy of undetermined significance. British Journal of Haematology, 1997, 97, 649-651.	2.5	36

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37	Meta-analysis of 49â€549 individuals imputed with the 1000 Genomes Project reveals an exonic damaging variant in <i>ANGPTL4</i> determining fasting TG levels. Journal of Medical Genetics, 2016, 53, 441-449.	3.2	34
38	The <i>TCF7L2</i> Diabetes Risk Variant is Associated with HbA _{1C} Levels: a Genomeâ€Wide Association Metaâ€Analysis. Annals of Human Genetics, 2010, 74, 471-478.	0.8	33
39	Common Variants in Mendelian Kidney Disease Genes and Their Association with Renal Function. Journal of the American Society of Nephrology: JASN, 2013, 24, 2105-2117.	6.1	33
40	Association of established thyroid peroxidase autoantibody (TPOAb) genetic variants with Hashimoto's thyroiditis. Autoimmunity, 2016, 49, 480-485.	2.6	28
41	The Greeks in the West: genetic signatures of the Hellenic colonisation in southern Italy and Sicily. European Journal of Human Genetics, 2016, 24, 429-436.	2.8	26
42	Environmental Risk Factors for Type 1 Diabetes Mellitus Development. Experimental and Clinical Endocrinology and Diabetes, 2017, 125, 563-570.	1.2	20
43	Family-based analysis of vitamin D receptor gene polymorphisms and type 1 diabetes in the population of South Croatia. Journal of Human Genetics, 2008, 53, 210-214.	2.3	19
44	Thyroglobulin Antibodies are Associated with Symptom Burden in Patients with Hashimoto's Thyroiditis: A Cross-Sectional Study. Immunological Investigations, 2019, 48, 198-209.	2.0	17
45	Genome-wide association analysis suggests novel loci for Hashimoto's thyroiditis. Journal of Endocrinological Investigation, 2019, 42, 567-576.	3.3	17
46	Differences in food consumption between patients with Hashimoto's thyroiditis and healthy individuals. Scientific Reports, 2020, 10, 10670.	3.3	17
47	Oculo-facio-cardio-dental syndrome in three succeeding generations: genotypic data and phenotypic features. Brazilian Journal of Medical and Biological Research, 2012, 45, 1315-1319.	1.5	16
48	Rare and common genetic variations in the Keap1/Nrf2 antioxidant response pathway impact thyroglobulin gene expression and circulating levels, respectively. Biochemical Pharmacology, 2020, 173, 113605.	4.4	16
49	Dietary Factors Associated with Plasma Thyroid Peroxidase and Thyroglobulin Antibodies. Nutrients, 2017, 9, 1186.	4.1	15
50	Genome-wide association analysis suggests novel loci underlying thyroid antibodies in Hashimoto's thyroiditis. Scientific Reports, 2019, 9, 5360.	3.3	15
51	Family-based analysis of tumor necrosis factor and lymphotoxin-α tag polymorphisms with type 1 diabetes in the population of South Croatia. Human Immunology, 2009, 70, 195-199.	2.4	14
52	Predictive Value of 8 Genetic Loci for Serum Uric Acid Concentration. Croatian Medical Journal, 2010, 51, 23-31.	0.7	14
53	Association of TNF promoter polymorphisms with type 1 diabetes in the South Croatian population. Biological Research, 2008, 41, .	3.4	14
54	Association of NOS3 tag polymorphisms with hypoxic-ischemic encephalopathy. Croatian Medical Journal, 2011, 52, 396-402.	0.7	13

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55	Genome-wide meta-analysis identifies novel loci associated with free triiodothyronine and thyroid-stimulating hormone. Journal of Endocrinological Investigation, 2019, 42, 1171-1180.	3.3	13
56	1311-induced changes in rat thyroid gland function. Brazilian Journal of Medical and Biological Research, 2007, 40, 1087-1094.	1.5	12
57	Common Variants in SLC17A3 Gene Affect Intra-personal Variation in Serum Uric Acid Levels in Longitudinal Time Series. Croatian Medical Journal, 2010, 51, 32-39.	0.7	12
58	The OSR1 rs12329305 Polymorphism Contributes to the Development of Congenital Malformations in Cases of Stillborn/Neonatal Death. Medical Science Monitor, 2014, 20, 1531-1538.	1.1	11
59	Association of established hypothyroidism-associated genetic variants with Hashimoto's thyroiditis. Journal of Endocrinological Investigation, 2017, 40, 1061-1067.	3.3	11
60	Genome-wide meta-analysis identifies novel gender specific loci associated with thyroid antibodies level in Croatians. Genomics, 2019, 111, 737-743.	2.9	11
61	AATF and SMARCA2 are associated with thyroid volume in Hashimoto's thyroiditis patients. Scientific Reports, 2020, 10, 1754.	3.3	11
62	Epidemiology of Hypothyroidism, Hyperthyroidism and Positive Thyroid Antibodies in the Croatian Population. Biology, 2022, 11, 394.	2.8	11
63	In Search of a Croatian Model of Nursing Education. Croatian Medical Journal, 2010, 51, 383-395.	0.7	10
64	Historic, Demographic, and Genetic Evidence for Increased Population Frequencies of CCR5Δ32 Mutation in Croatian Island Isolates after Lethal 15th Century Epidemics. Croatian Medical Journal, 2009, 50, 34-42.	0.7	9
65	NeuroD1 gene and interleukin-18 gene polymorphisms in type 1 diabetes in Dalmatian population of Southern Croatia. Croatian Medical Journal, 2006, 47, 571-8.	0.7	9
66	Oxygenation alters ganglioside expression in rat liver following partial hepatectomy. Biochemical and Biophysical Research Communications, 2005, 330, 131-141.	2.1	8
67	Association of NOS3 gene variants and clinical contributors of hypoxic-ischemic encephalopathy. Brazilian Journal of Medical and Biological Research, 2014, 47, 869-875.	1.5	8
68	Genome-wide meta-analysis identifies novel loci associated with parathyroid hormone level. Molecular Medicine, 2018, 24, 15.	4.4	8
69	The effect of food groups and nutrients on thyroid hormone levels in healthy individuals. Nutrition, 2021, 91-92, 111394.	2.4	8
70	Environmental Factors That Affect Parathyroid Hormone and Calcitonin Levels. International Journal of Molecular Sciences, 2022, 23, 44.	4.1	8
71	High prevalence of glaucoma in Veli Brgud, Croatia, is caused by a dominantly inherited T377M mutation in the MYOC gene. British Journal of Ophthalmology, 2008, 92, 1567-1568.	3.9	7
72	Leprosy epidemics during history increased protective allele frequency of PARK2/PACRG genes in the population of the Mljet Island, Croatia. European Journal of Medical Genetics, 2011, 54, e548-52.	1.3	7

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73	Expression of Neutral Glycosphingolipids in Cytokine-Stimulated Human Endothelial Cells. Biochemistry (Moscow), 2004, 69, 513-519.	1.5	6
74	Genome-Wide Analysis Identifies Two Susceptibility Loci for Positive Thyroid Peroxidase and Thyroglobulin Antibodies. Journal of Clinical Endocrinology and Metabolism, 2020, 105, 944-951.	3.6	6
75	IL12RB2 Gene Is Associated with the Age of Type 1 Diabetes Onset in Croatian Family Trios. PLoS ONE, 2012, 7, e49133.	2.5	6
76	Clycosyltransferase B4GALNT1 and type 1 diabetes in Croatian population. Clinical Biochemistry, 2009, 42, 819-822.	1.9	5
77	Association of Established Thyroid-stimulating Hormone and Free Thyroxine Genetic Variants with Hashimoto's Thyroiditis. Immunological Investigations, 2017, 46, 625-638.	2.0	5
78	Genome-wide association meta-analysis for total thyroid hormone levels in Croatian population. Journal of Human Genetics, 2019, 64, 473-480.	2.3	5
79	Genetic Variants in the ST6GAL1 Gene Are Associated with Thyroglobulin Plasma Level in Healthy Individuals. Thyroid, 2019, 29, 886-893.	4.5	5
80	Pregnancy in Adolescent Rats, Growth and Neurodevelopment in their Offspring. Archives of Physiology and Biochemistry, 2001, 109, 450-456.	2.1	4
81	Complete trisomy 10p resulting from an extra stable telocentric chromosome. American Journal of Medical Genetics, Part A, 2012, 158A, 1778-1781.	1.2	4
82	Effects of genetic variants on serum parathyroid hormone in hyperparathyroidism and end-stage renal disease patients. Medicine (United States), 2018, 97, e10834.	1.0	3
83	Genome-Wide Association Study to Identify Common Variants Associated with Brachial Circumference: A Meta-Analysis of 14 Cohorts. PLoS ONE, 2012, 7, e31369.	2.5	3
84	Association of TNF promoter polymorphisms with type 1 diabetes in the South Croatian population. Biological Research, 2008, 41, 157-63.	3.4	3
85	Prognostic Value of B-Symptoms in Low-Grade Non-Hodgkin's Lymphomas. Leukemia and Lymphoma, 1994, 13, 357-358.	1.3	2
86	Immunohistochemical analysis of hepatic ganglioside distribution following a partial hepatectomy and exposure to different hyperbaric oxygen treatments. Acta Histochemica, 2008, 110, 66-75.	1.8	2
87	The effect of multiple nutrients on plasma parathyroid hormone level in healthy individuals. International Journal of Food Sciences and Nutrition, 2019, 70, 638-644.	2.8	2
88	Molecular Characterization of Glucose-6-phosphate Dehydrogenase Deficiency in Families from the Republic of Macedonia and Genotype-phenotype Correlation. Medicinski Arhiv = Medical Archives = Archives De Médecine, 2015, 69, 284.	0.9	2
89	Thyroid hormone levels are associated with metabolic components: a cross-sectional study. Croatian Medical Journal, 2020, 61, 230-238.	0.7	2
90	Nasal dermal sinus cysts with intracranial extension in a child mosaic for a supernumerary ring chromosome 20. International Journal of Pediatric Otorhinolaryngology Extra, 2012, 7, 73-78.	0.1	1

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91	Correlation of serial MRI findings and clinical outcome in the first Croatian patient with acute necrotizing encephalopathy. Croatian Medical Journal, 2014, 55, 431-433.	0.7	1
92	Changes of defense mechanisms and personality profile during group analytic treatment. Collegium Antropologicum, 2005, 29, 551-8.	0.2	1
93	Analysis of the C609T polymorphism of NQO1 gene in South Croatian patients with hematological malignancies. Collegium Antropologicum, 2011, 35, 385-8.	0.2	1
94	Genome-Wide Association Analysis and Genomic Prediction of Thyroglobulin Plasma Levels. International Journal of Molecular Sciences, 2022, 23, 2173.	4.1	1
95	Genetics of Type 1 Diabetes. , 0, , .		0
96	Large-Scale Genomic Analyses Link Reproductive Aging to Hypothalamic Signaling, Breast Cancer Susceptibility, and BRCA1-Mediated DNA Repair. Obstetrical and Gynecological Survey, 2015, 70, 758-762.	0.4	0
97	Limited survivability of unbalanced progeny of carriers of a unique t(4;19)(p15.32;p13.3): a study in multiple generations. Molecular Cytogenetics, 2017, 10, 29.	0.9	0
98	Correction: Environmental Risk Factors for Type 1 Diabetes Mellitus Development. Experimental and Clinical Endocrinology and Diabetes, 2018, , .	1.2	0
99	Distinct Cerebellar Glycosphingolipid Phenotypes in Wistar and Lewis Rats. Neurochemical Journal, 2020, 14, 20-24.	0.5	0
100	Identification of novel genetic loci associated with thyroid function. Endocrine Abstracts, 0, , .	0.0	0
101	Determinants of thyroid volume in healthy young adults of Dalmatia. Periodicum Biologorum, 2020, 121-122, 65-69.	0.1	0
102	Hyperbaric environment up-regulates CD15s expression on leukocytes, down-regulates CD77 expression on renal cells and up-regulates CD34 expression on pulmonary and cardiac cells in rat. Iranian Journal of Basic Medical Sciences, 2016, 19, 821-828.	1.0	0

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