

Tomasz Suchocki

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

561
citations

1040056

9
h-index

642732

23
g-index

33
all docs

33
docs citations

33
times ranked

980
citing authors

#	ARTICLE	IF	CITATIONS
1	History of Heart Failure in Patients Hospitalized Due to COVID-19: Relevant Factor of In-Hospital Complications and All-Cause Mortality up to Six Months. <i>Journal of Clinical Medicine</i> , 2022, 11, 241.	2.4	16
2	Genome-Wide Genomic and Functional Association Study for Workability and Calving Traits in Holstein Cattle. <i>Animals</i> , 2022, 12, 1127.	2.3	1
3	The Thousand Polish Genomes—A Database of Polish Variant Allele Frequencies. <i>International Journal of Molecular Sciences</i> , 2022, 23, 4532.	4.1	15
4	Identification of functional features underlying heat stress response in Sprague–Dawley rats using mixed linear models. <i>Scientific Reports</i> , 2022, 12, 7671.	3.3	0
5	Beyond GWAS—Could Genetic Differentiation within the Allograft Rejection Pathway Shape Natural Immunity to COVID-19?. <i>International Journal of Molecular Sciences</i> , 2022, 23, 6272.	4.1	3
6	High soluble transferrin receptor in patients with heart failure: a measure of iron deficiency and a strong predictor of mortality. <i>European Journal of Heart Failure</i> , 2021, 23, 919-932.	7.1	46
7	Iron deficiency contributes to resistance to endogenous erythropoietin in anaemic heart failure patients. <i>European Journal of Heart Failure</i> , 2021, 23, 1677-1686.	7.1	11
8	SNP prioritization in targeted sequencing data associated with humoral immune responses in chicken. <i>Poultry Science</i> , 2021, 100, 101433.	3.4	2
9	The application of deep learning for the classification of correct and incorrect SNP genotypes from whole-genome DNA sequencing pipelines. <i>Journal of Applied Genetics</i> , 2020, 61, 607-616.	1.9	4
10	Tuning the genomic evaluation system of Holstein-Friesian cattle. <i>Computers and Electronics in Agriculture</i> , 2020, 175, 105594.	7.7	1
11	Exploring the Potential Genetic Heterogeneity in the Incidence of Hoof Disorders in Austrian Fleckvieh and Braunvieh Cattle. <i>Frontiers in Genetics</i> , 2020, 11, 577116.	2.3	3
12	10-year trends in hospitalization rates due to heart failure and related in-hospital mortality in Poland (2010–2019). <i>ESC Heart Failure</i> , 2020, 7, 3365-3373.	3.1	5
13	Two-stage genome-wide association study for the identification of causal variants underlying hoof disorders in cattle. <i>Journal of Dairy Science</i> , 2020, 103, 4483-4494.	3.4	5
14	Assessing the degree of stratification between closely related Holstein-Friesian populations. <i>Journal of Applied Genetics</i> , 2017, 58, 521-526.	1.9	4
15	Effects of intravenous iron therapy in iron-deficient patients with systolic heart failure: a meta-analysis of randomized controlled trials. <i>European Journal of Heart Failure</i> , 2016, 18, 786-795.	7.1	270
16	Utilization of information from gene networks towards a better understanding of functional similarities between complex traits: a dairy cattle model. <i>Journal of Applied Genetics</i> , 2016, 57, 129-133.	1.9	2
17	Identification of candidate genes and mutations in QTL regions for immune responses in chicken. <i>Animal Genetics</i> , 2015, 46, 247-254.	1.7	7
18	Genome-wide association study for semen production traits in Holstein-Friesian bulls. <i>Journal of Dairy Science</i> , 2015, 98, 5774-5780.	3.4	46

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19	Modelling QTL effect on BTA06 using random regression test day models. <i>Journal of Applied Genetics</i> , 2013, 54, 49-60.	1.9	6
20	HIGH SOLUBLE TRANSFERRIN RECEPTOR IN PATIENTS WITH SYSTOLIC HEART FAILURE: A MEASURE OF IRON DEFICIENCY AND A STRONG PREDICTOR OF MORTALITY. <i>Journal of the American College of Cardiology</i> , 2013, 61, E749.	2.8	2
21	Iron Status and Survival in Diabetic Patients With Coronary Artery Disease. <i>Diabetes Care</i> , 2013, 36, 4147-4156.	8.6	61
22	Polymorphism within TATA-box of bovine lactoferrin gene and its association with performance traits in Holstein cattle. <i>Livestock Science</i> , 2012, 149, 267-274.	1.6	3
23	Statistical modelling of growth using a mixed model with orthogonal polynomials. <i>Journal of Applied Genetics</i> , 2011, 52, 95-100.	1.9	4
24	Fitting and validating the genomic evaluation model to Polish Holstein-Friesian cattle. <i>Journal of Applied Genetics</i> , 2011, 52, 363-366.	1.9	15
25	Testing candidate gene effects on milk production traits in dairy cattle under various parameterizations and modes of inheritance. <i>Journal of Dairy Science</i> , 2010, 93, 2703-2717.	3.4	13
26	Additive effects of 19 porcine SNPs on growth rate, meat content and selection index. <i>Journal of Applied Genetics</i> , 2009, 50, 235-243.	1.9	4
27	The impact of single nucleotide polymorphism selection on prediction of genomewide breeding values. <i>BMC Proceedings</i> , 2009, 3, S13.	1.6	8