

# Sunday O Peters

## List of Publications by Year in descending order

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Version: 2024-02-01

58  
papers

1,051  
citations

567281

15  
h-index

454955

30  
g-index

59  
all docs

59  
docs citations

59  
times ranked

1408  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic evaluation of semen traits in Friesian bulls raised in Egypt. <i>Theriogenology</i> , 2022, 179, 39-44.	2.1	3
2	Plasma Carboxyl-Metabolome Is Associated with Average Daily Gain Divergence in Beef Steers. <i>Animals</i> , 2021, 11, 67.	2.3	1
3	Comparative accuracies of genetic values predicted for economically important milk traits, genome-wide association, and linkage disequilibrium patterns of Canadian Holstein cows. <i>Journal of Dairy Science</i> , 2021, 104, 1900-1916.	3.4	8
4	Genetic parameters, phenotypic and genetic trends of litter size on different breeds of goats in Egypt. <i>Tropical Animal Health and Production</i> , 2021, 53, 286.	1.4	3
5	Phenotypic and genetic parameters of productive traits in Rahmani and Romanov sheep and crossbreds. <i>Journal of Animal Science and Technology</i> , 2021, 63, 1211-1222.	2.5	2
6	Effects of a blend of mannan and glucan on growth performance, apparent nutrient digestibility, energy status, and whole-blood immune gene expression of beef steers during a 42-d receiving period. <i>Translational Animal Science</i> , 2021, 5, txaa226.	1.1	1
7	Study on the prevalence and genetic diversity of <i>Eimeria</i> species from broilers and free-range chickens in KwaZulu-Natal province, South Africa. <i>Onderstepoort Journal of Veterinary Research</i> , 2020, 87, e1-e10.	1.2	7
8	Evolutionary Pattern of Interferon Alpha Genes in Bovidae and Genetic Diversity of IFNAA in the Bovine Genome. <i>Frontiers in Immunology</i> , 2020, 11, 580412.	4.8	5
9	Genomic Prediction With Different Heritability, QTL, and SNP Panel Scenarios Using Artificial Neural Network. <i>IEEE Access</i> , 2020, 8, 147995-148006.	4.2	4
10	Differential Expression of IGF1, IGFBP5, MSTN, and MYH1 Across Different Age Classes in American Quarter Horses. <i>Journal of Equine Veterinary Science</i> , 2020, 94, 103226.	0.9	0
11	Comparative effects of two multispecies direct-fed microbial products on energy status, nutrient digestibility, and ruminal fermentation, bacterial community, and metabolome of beef steers. <i>Journal of Animal Science</i> , 2020, 98, .	0.5	10
12	Effects of a blend of <i>Saccharomyces cerevisiae</i> -based direct-fed microbial and fermentation products on plasma carbonyl-metabolome and fecal bacterial community of beef steers. <i>Journal of Animal Science and Biotechnology</i> , 2020, 11, 14.	5.3	12
13	RNA-seq profiling of skin in temperate and tropical cattle. <i>Journal of Animal Science and Technology</i> , 2020, 62, 141-158.	2.5	8
14	Leveraging Available Resources and Stakeholder Involvement for Improved Productivity of African Livestock in the Era of Genomic Breeding. <i>Frontiers in Genetics</i> , 2019, 10, 357.	2.3	27
15	Comparison of linear model and artificial neural network using antler beam diameter and length of white-tailed deer ( <i>Odocoileus virginianus</i> ) dataset. <i>PLoS ONE</i> , 2019, 14, e0212545.	2.5	5
16	Use of discriminant analysis for the evaluation of coccidiosis resistance parameters in chickens raised in hot humid tropical environment. <i>Tropical Animal Health and Production</i> , 2018, 50, 1161-1166.	1.4	7
17	Sequence variation of necdin gene in Bovidae. <i>Journal of Animal Science and Technology</i> , 2018, 60, 32.	2.5	3
18	Genetic variation in N- and C-terminal regions of bovine DNAJA1 heat shock protein gene in African, Asian and American cattle. <i>Journal of Genomics</i> , 2018, 6, 1-8.	0.9	5

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19	Genetic Diversity of Bovine Major Histocompatibility Complex Class II DRB3 locus in cattle breeds from Asia compared to those from Africa and America. <i>Journal of Genomics</i> , 2018, 6, 88-97.	0.9	11
20	Computational genome-wide identification of heat shock protein genes in the bovine genome. <i>F1000Research</i> , 2018, 7, 1504.	1.6	10
21	Phylogeny of Pakistani Cattle Breeds using Mitochondrial Cytochrome b Gene. <i>Pakistan Journal of Zoology</i> , 2018, 50, .	0.2	0
22	Association of SNP variants of MHC Class II DRB gene with thermo-physiological traits in tropical goats. <i>Tropical Animal Health and Production</i> , 2017, 49, 323-336.	1.4	5
23	Analysis of culling records and estimation of genetic parameters for longevity and some production traits in Holstein dairy cattle. <i>Journal of Applied Animal Research</i> , 2017, 45, 524-528.	1.2	14
24	Genetic diversity among <i>Babesia rossi</i> detected in naturally infected dogs in Abeokuta, Nigeria, based on 18S rRNA gene sequences. <i>Acta Parasitologica</i> , 2017, 62, 192-198.	1.1	11
25	Molecular evolution of type II MAGE genes from ancestral MAGED2 gene and their phylogenetic resolution of basal mammalian clades. <i>Mammalian Genome</i> , 2017, 28, 443-454.	2.2	11
26	Genetic diversity among <i>Trypanosoma vivax</i> strains detected in naturally infected cattle in Nigeria based on ITS1 of rDNA and diagnostic antigen gene sequences. <i>Journal of Parasitic Diseases</i> , 2017, 41, 433-441.	1.0	3
27	Conservation of Repeats at the Mammalian KCNQ1OT1-CDKN1C Region Suggests a Role in Genomic Imprinting. <i>Evolutionary Bioinformatics</i> , 2017, 13, 117693431771523.	1.2	3
28	Molecular cloning, sequence analysis and tissue expression of bovine imprinted <i>ASCL2</i> gene. <i>South African Journal of Animal Sciences</i> , 2017, 47, 813.	0.5	0
29	Nucleotide sequence variability analysis of Major Histocompatibility Complex Class II DQA1 gene in Nigerian goats. <i>Genetika</i> , 2017, 49, 865-874.	0.4	0
30	Phylogeny of <i>Trypanosoma brucei</i> and <i>Trypanosoma evansi</i> in naturally infected cattle in Nigeria by analysis of repetitive and ribosomal DNA sequences. <i>Tropical Animal Health and Production</i> , 2016, 48, 1235-1240.	1.4	4
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37	A Novel TaqI Polymorphism in the Coding Region of the Ovine TNXB Gene in the MHC Class III Region: Morphostructural and Physiological Influences. <i>Biochemical Genetics</i> , 2014, 52, 1-14.	1.7	5
38	Identification of single nucleotide polymorphisms in the agouti signaling protein (ASIP) gene in some goat breeds in tropical and temperate climates. <i>Molecular Biology Reports</i> , 2013, 40, 4447-4457.	2.3	14
39	Genetic Diversity in Exon 2 of the Major Histocompatibility Complex Class II DQB1 Locus in Nigerian Goats. <i>Biochemical Genetics</i> , 2013, 51, 954-966.	1.7	16
40	Molecular survey of pathogenic trypanosomes in naturally infected Nigerian cattle. <i>Research in Veterinary Science</i> , 2013, 94, 555-561.	1.9	50
41	Application of multivariate heavy-tailed distributions to residuals in the estimation of genetic parameters of growth traits in beef cattle <sup>1</sup> . <i>Journal of Animal Science</i> , 2013, 91, 1552-1561.	0.5	2
42	Heritability and Bayesian genome-wide association study of first service conception and pregnancy in Brangus heifers <sup>1</sup> . <i>Journal of Animal Science</i> , 2013, 91, 605-612.	0.5	58
43	Genotyping-by-Sequencing (GBS): A Novel, Efficient and Cost-Effective Genotyping Method for Cattle Using Next-Generation Sequencing. <i>PLoS ONE</i> , 2013, 8, e62137.	2.5	184
44	Molecular Diagnosis of Subclinical African Trypanosoma vivax Infection and Association with Physiological Indices and Serum Metabolites in Extensively Managed Goats in the Tropics. <i>Open Journal of Veterinary Medicine</i> , 2013, 03, 39-45.	0.4	13
45	Physiological and haematological indices suggest superior heat tolerance of white-coloured West African Dwarf sheep in the hot humid tropics. <i>Tropical Animal Health and Production</i> , 2012, 45, 157-165.	1.4	47
46	Preliminary association of coat colour types and tolerance to Haemonchus contortus infection in West African Dwarf sheep. <i>Journal of Applied Animal Research</i> , 2012, 40, 1-7.	1.2	4
47	Gene network analyses of first service conception in Brangus heifers: Use of genome and trait associations, hypothalamic-transcriptome information, and transcription factors <sup>1</sup> . <i>Journal of Animal Science</i> , 2012, 90, 2894-2906.	0.5	66
48	Morphological and microsatellite DNA diversity of Nigerian indigenous sheep. <i>Journal of Animal Science and Biotechnology</i> , 2012, 3, 38.	5.3	44
49	Multivariate analysis of sexual size dimorphism in local turkeys (Meleagris gallopavo) in Nigeria. <i>Tropical Animal Health and Production</i> , 2012, 44, 1089-1095.	1.4	10
50	Application of principal component and discriminant analyses to morpho-structural indices of indigenous and exotic chickens raised under intensive management system. <i>Tropical Animal Health and Production</i> , 2012, 44, 1247-1254.	1.4	25
51	Effect of crossbreeding on fertility, hatchability and embryonic mortality of Nigerian local chickens. <i>Tropical Animal Health and Production</i> , 2012, 44, 505-510.	1.4	14
52	Application of multivariate principal component analysis to morphological characterization of indigenous goats in Southern Nigeria. <i>Acta Agriculturae Slovenica</i> , 2011, 98, .	0.3	32
53	Haematological studies on frizzled and naked neck genotypes of Nigerian native chickens. <i>Tropical Animal Health and Production</i> , 2011, 43, 631-638.	1.4	36
54	Growth performance of Nigerian local chickens in crosses involving an exotic broiler breeder. <i>Tropical Animal Health and Production</i> , 2011, 43, 643-650.	1.4	22

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55	Comparative Assessment of Growth in Pure and Crossbred Turkeys in a Humid Tropical Environment. International Journal of Poultry Science, 2010, 9, 368-375.	0.1	18
56	Genotype and Sex Effect on Gastrointestinal Nutrient Content, Microflora and Carcass Traits in Nigerian Native Chickens. International Journal of Poultry Science, 2010, 9, 731-737.	0.1	11
57	Semen Quality Traits of Seven Strain of Chickens Raised in the Humid Tropics. International Journal of Poultry Science, 2008, 7, 949-953.	0.1	53
58	Gene Segregation Effects on Fertility and Hatchability of Pure and Crossbred Chicken Genotypes in the Humid Tropics. International Journal of Poultry Science, 2008, 7, 954-958.	0.1	13