Paul A Mcgettigan

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

37 papers	1,621	25	37
	citations	h-index	g-index
37 ext. papers	1,933 ext. citations	4.8 avg, IF	4.69 L-index

#	Paper	IF	Citations
37	Expression Quantitative Trait Loci in Equine Skeletal Muscle Reveals Heritable Variation in Metabolism and the Training Responsive Transcriptome. <i>Frontiers in Genetics</i> , 2019 , 10, 1215	4.5	3
36	Intragenic sequences in the trophectoderm harbour the greatest proportion of methylation errors in day 17 bovine conceptuses generated using assisted reproductive technologies. <i>BMC Genomics</i> , 2018 , 19, 438	4.5	10
35	Equine skeletal muscle adaptations to exercise and training: evidence of differential regulation of autophagosomal and mitochondrial components. <i>BMC Genomics</i> , 2017 , 18, 595	4.5	16
34	Atypical benign partial epilepsy of childhood with acquired neurocognitive, lexical semantic, and autistic spectrum disorder. <i>Epilepsy & Behavior Case Reports</i> , 2016 , 6, 42-8	1.2	8
33	GOexpress: an R/Bioconductor package for the identification and visualisation of robust gene ontology signatures through supervised learning of gene expression data. <i>BMC Bioinformatics</i> , 2016 , 17, 126	3.6	16
32	Recessive NEK9 mutation causes a lethal skeletal dysplasia with evidence of cell cycle and ciliary defects. <i>Human Molecular Genetics</i> , 2016 , 25, 1824-35	5.6	28
31	Transcriptomic Study on Ovine Immune Responses to Fasciola hepatica Infection. <i>PLoS Neglected Tropical Diseases</i> , 2016 , 10, e0005015	4.8	33
30	Genome sequencing of the extinct Eurasian wild aurochs, Bos primigenius, illuminates the phylogeography and evolution of cattle. <i>Genome Biology</i> , 2015 , 16, 234	18.3	127
29	Unexpected genetic heterogeneity for primary ciliary dyskinesia in the Irish Traveller population. <i>European Journal of Human Genetics</i> , 2015 , 23, 210-7	5.3	18
28	RNA sequencing provides exquisite insight into the manipulation of the alveolar macrophage by tubercle bacilli. <i>Scientific Reports</i> , 2015 , 5, 13629	4.9	26
27	Epigenetic processes in the male germline. <i>Reproduction, Fertility and Development</i> , 2015 , 27, 725-38	1.8	22
26	Variation in the Ovine Abomasal Lymph Node Transcriptome between Breeds Known to Differ in Resistance to the Gastrointestinal Nematode. <i>PLoS ONE</i> , 2015 , 10, e0124823	3.7	10
25	Towards the identification of a genetic basis for Landau-Kleffner syndrome. <i>Epilepsia</i> , 2014 , 55, 858-65	6.4	30
24	Atypical Alstrom syndrome with novel ALMS1 mutations precluded by current diagnostic criteria. <i>European Journal of Medical Genetics</i> , 2014 , 57, 55-9	2.6	17
23	Proteomic analysis of uterine fluid during the pre-implantation period of pregnancy in cattle. <i>Reproduction</i> , 2014 , 147, 575-87	3.8	83
22	A novel locus for episodic ataxia:UBR4 the likely candidate. <i>European Journal of Human Genetics</i> , 2014 , 22, 505-10	5.3	49
21	Alterations in expression of endometrial genes coding for proteins secreted into the uterine lumen during conceptus elongation in cattle. <i>BMC Genomics</i> , 2013 , 14, 321	4.5	35

(2008-2013)

20	Transcriptomics in the RNA-seq era. Current Opinion in Chemical Biology, 2013, 17, 4-11	9.7	211
19	Polymorphism discovery and allele frequency estimation using high-throughput DNA sequencing of target-enriched pooled DNA samples. <i>BMC Genomics</i> , 2012 , 13, 16	4.5	13
18	Physiological status alters immunological regulation of bovine follicle differentiation in dairy cattle. <i>Journal of Reproductive Immunology</i> , 2012 , 96, 34-44	4.2	11
17	Recessive mutations in MCM4/PRKDC cause a novel syndrome involving a primary immunodeficiency and a disorder of DNA repair. <i>Journal of Medical Genetics</i> , 2012 , 49, 242-5	5.8	48
16	Identification of a mutation in LARS as a novel cause of infantile hepatopathy. <i>Molecular Genetics and Metabolism</i> , 2012 , 106, 351-8	3.7	64
15	Conceptus-endometrium crosstalk during maternal recognition of pregnancy in cattle. <i>Biology of Reproduction</i> , 2012 , 87, 6, 1-9	3.9	45
14	Evidence for an early endometrial response to pregnancy in cattle: both dependent upon and independent of interferon tau. <i>Physiological Genomics</i> , 2012 , 44, 799-810	3.6	66
13	Effect of the metabolic environment at key stages of follicle development in cattle: focus on steroid biosynthesis. <i>Physiological Genomics</i> , 2012 , 44, 504-17	3.6	45
12	Low MAD2 expression levels associate with reduced progression-free survival in patients with high-grade serous epithelial ovarian cancer. <i>Journal of Pathology</i> , 2012 , 226, 746-55	9.4	57
11	Molecular aspects of mucin biosynthesis and mucus formation in the bovine cervix during the periestrous period. <i>Physiological Genomics</i> , 2012 , 44, 1165-78	3.6	26
10	Sequential analysis of global gene expression profiles in immature and in vitro matured bovine oocytes: potential molecular markers of oocyte maturation. <i>BMC Genomics</i> , 2011 , 12, 151	4.5	57
9	First implication of STRA6 mutations in isolated anophthalmia, microphthalmia, and coloboma: a new dimension to the STRA6 phenotype. <i>Human Mutation</i> , 2011 , 32, 1417-26	4.7	62
8	RNA sequencing reveals novel gene clusters in bovine conceptuses associated with maternal recognition of pregnancy and implantation. <i>Biology of Reproduction</i> , 2011 , 85, 1143-51	3.9	68
7	Temporal dysregulation of cortical gene expression in the isolation reared Wistar rat. <i>Journal of Neurochemistry</i> , 2010 , 113, 601-14	6	13
6	Hippocampal region-specific regulation of NF-kappaB may contribute to learning-associated synaptic reorganisation. <i>Brain Research Bulletin</i> , 2010 , 81, 385-90	3.9	27
5	Characterization of the equine skeletal muscle transcriptome identifies novel functional responses to exercise training. <i>BMC Genomics</i> , 2010 , 11, 398	4.5	61
4	A complete mitochondrial genome sequence from a mesolithic wild aurochs (Bos primigenius). <i>PLoS ONE</i> , 2010 , 5, e9255	3.7	64
3	Co-regulation of Gremlin and Notch signalling in diabetic nephropathy. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2008 , 1782, 10-21	6.9	84

Temporal change in gene expression in the rat dentate gyrus following passive avoidance learning.

Journal of Neurochemistry, **2007**, 101, 1085-98

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Integrating transcription factor binding site information with gene expression datasets. *Bioinformatics*, **2007**, 23, 298-305

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