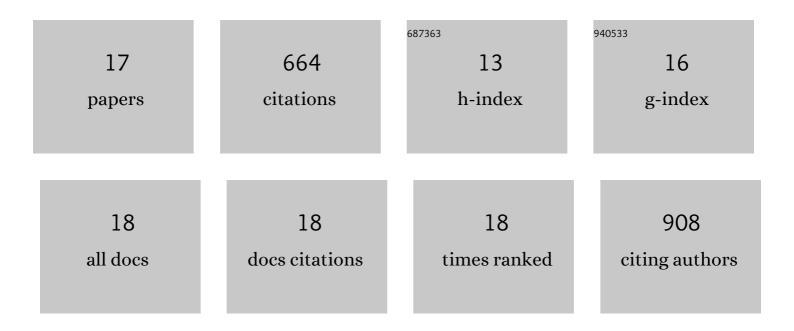
Julie Demars

List of Publications by Year in descending order

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LILLE DEMADS

#	Article	IF	CITATIONS
1	Genome-Wide Association Studies Identify Two Novel BMP15 Mutations Responsible for an Atypical Hyperprolificacy Phenotype in Sheep. PLoS Genetics, 2013, 9, e1003482.	3.5	145
2	Analysis of the IGF2/H19 imprinting control region uncovers new genetic defects, including mutations of OCT-binding sequences, in patients with 11p15 fetal growth disorders. Human Molecular Genetics, 2010, 19, 803-814.	2.9	106
3	Genome-Wide Identification of the Mutation Underlying Fleece Variation and Discriminating Ancestral Hairy Species from Modern Woolly Sheep. Molecular Biology and Evolution, 2017, 34, 1722-1729.	8.9	76
4	Epigenetic and genetic disturbance of the imprinted 11p15 region in Beckwith–Wiedemann and Silver–Russell syndromes. Clinical Genetics, 2012, 81, 350-361.	2.0	69
5	New insights into the pathogenesis of beckwith-wiedemann and silver-russell syndromes: Contribution of small copy number variations to 11p15 imprinting defects. Human Mutation, 2011, 32, 1171-1182.	2.5	53
6	QTL for resistance to <i>Salmonella</i> carrier state confirmed in both experimental and commercial chicken lines. Animal Genetics, 2009, 40, 590-597.	1.7	37
7	Effect of polyethylene glycol-based preservation solutions on graft injury in experimental kidney transplantation. British Journal of Surgery, 2011, 98, 368-378.	0.3	26
8	Epigenetic and Genetic Mechanisms of Abnormal 11p15 Genomic Imprinting in Silver-Russell and Beckwith-Wiedemann Syndromes. Current Medicinal Chemistry, 2011, 18, 1740-1750.	2.4	24
9	New QTL for resistance to Salmonella carrier-state identified on fowl microchromosomes. Molecular Genetics and Genomics, 2011, 285, 237-243.	2.1	23
10	High resolution physical map of porcine chromosome 7 QTL region and comparative mapping of this region among vertebrate genomes. BMC Genomics, 2006, 7, 13.	2.8	21
11	New Insights into the Melanophilin (MLPH) Gene Affecting Coat Color Dilution in Rabbits. Genes, 2018, 9, 430.	2.4	20
12	In-Depth Analysis Reveals Production of Circular RNAs from Non-Coding Sequences. Cells, 2020, 9, 1806.	4.1	16
13	Genetic variants within the second intron of theKCNQ1gene affect CTCF binding and confer a risk of Beckwith–Wiedemann syndrome upon maternal transmission. Journal of Medical Genetics, 2014, 51, 502-511.	3.2	15
14	Metabolic and histochemical characteristics of fat and muscle tissues in homozygous or heterozygous pigs for the body composition QTL located on chromosome 7. Physiological Genomics, 2007, 30, 232-241.	2.3	13
15	Intergenerational Transmission of Characters Through Genetics, Epigenetics, Microbiota, and Learning in Livestock. Frontiers in Genetics, 2019, 10, 1058.	2.3	12
16	Towards candidate genes affecting body fatness at the SSC7 QTL by expression analyses. Journal of Animal Breeding and Genetics, 2012, 129, 316-324.	2.0	8
17	Human Fetal Growth Disorders and Imprinting Anomalies. Research and Perspectives in Endocrine Interactions, 2014, , 101-129.	0.2	0