## Dominique Rocha

List of Publications by Year in descending order

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31 2,350 17 31 papers citations h-index g-index

32 32 32 2625
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#	Article	IF	Citations
1	Whole-genome sequencing of 234 bulls facilitates mapping of monogenic and complex traits in cattle. Nature Genetics, 2014, 46, 858-865.	21.4	697
2	Meta-analysis of genome-wide association studies for cattle stature identifies common genes that regulate body size in mammals. Nature Genetics, 2018, 50, 362-367.	21.4	286
3	Molecular cloning, sequencing and expression of the mRNA encoding human $Cdx1$ and $Cdx2$ homeobox. Down-regulation of $Cdx1$ and $Cdx2$ mRNA expression during colorectal carcinogenesis. International Journal of Cancer, 1997, 74, 35-44.	5.1	201
4	Differential Gene Expression in the Murine Thymus Assayed by Quantitative Hybridization of Arrayed cDNA Clones. Genomics, 1995, 29, 207-216.	2.9	185
5	Detection of Haplotypes Associated with Prenatal Death in Dairy Cattle and Identification of Deleterious Mutations in GART, SHBG and SLC37A2. PLoS ONE, 2013, 8, e65550.	2.5	137
6	Identification of large intergenic non-coding RNAs in bovine muscle using next-generation transcriptomic sequencing. BMC Genomics, 2014, 15, 499.	2.8	120
7	Within-breed and multi-breed GWAS on imputed whole-genome sequence variants reveal candidate mutations affecting milk protein composition in dairy cattle. Genetics Selection Evolution, 2017, 49, 68.	3.0	111
8	Uncovering Adaptation from Sequence Data: Lessons from Genome Resequencing of Four Cattle Breeds. Genetics, 2016, 203, 433-450.	2.9	93
9	Novel Insights into the Bovine Polled Phenotype and Horn Ontogenesis in Bovidae. PLoS ONE, 2013, 8, e63512.	2.5	78
10	Genome-Wide Study of Structural Variants in Bovine Holstein, Montbéliarde and Normande Dairy Breeds. PLoS ONE, 2015, 10, e0135931.	2.5	52
11	Multi-breed and multi-trait co-association analysis of meat tenderness and other meat quality traits in three French beef cattle breeds. Genetics Selection Evolution, 2016, 48, 37.	3.0	50
12	Genetic Variability and Linkage Disequilibrium Patterns in the Bovine DNAJA1 Gene. Molecular Biotechnology, 2010, 44, 190-197.	2.4	43
13	From hybridization image to numerical values: a practical, high throughput quantification system for high density filter hybridizations. Genetic Analysis, Techniques and Applications, 1996, 12, 151-162.	1.5	41
14	Identification of copy number variation in French dairy and beef breeds using next-generation sequencing. Genetics Selection Evolution, 2017, 49, 77.	3.0	37
15	Gene-based single nucleotide polymorphism discovery in bovine muscle using next-generation transcriptomic sequencing. BMC Genomics, 2013, 14, 307.	2.8	32
16	Commercial pigs: an untapped resource for human obesity research?. Drug Discovery Today, 2006, 11, 475-477.	6.4	27
17	Genetic characterization of the Blonde d'Aquitaine cattle breed using microsatellite markers and relationship with three other French cattle populations. Journal of Animal Breeding and Genetics, 2011, 128, 201-208.	2.0	16
18	Construction of a large collection of small genome variations in French dairy and beef breeds using whole-genome sequences. Genetics Selection Evolution, 2016, 48, 87.	3.0	15

#	Article	IF	CITATIONS
19	Modulation of mRNA levels in the presence of thymocytes and genome mapping for a set of genes expressed in mouse thymic epithelial cells. Immunogenetics, 1997, 46, 142-151.	2.4	14
20	Usp9y (ubiquitin-specific protease 9 gene on the Y) is associated with a functional promoter and encodes an intact open reading frame homologous to Usp9x that is under selective constraint. Mammalian Genome, 2003, 14, 437-447.	2.2	13
21	Survey of mitochondrial sequences integrated into the bovine nuclear genome. Scientific Reports, 2020, 10, 2077.	3.3	13
22	Concordance analysis for QTL detection in dairy cattle: a case study of leg morphology. Genetics Selection Evolution, 2014, 46, 31.	3.0	12
23	Polymorphism of the Prion Protein in Mammals: A Phylogenetic Approach. Recent Patents on DNA & Gene Sequences, 2009, 3, 63-71.	0.7	9
24	Genetic diversity and selection signatures of the beef â€~Charolais de Cuba' breed. Scientific Reports, 2018, 8, 11005.	3.3	9
25	Specific proteins allow classification of pigs according to sire breed, rearing environment and gender. Livestock Science, 2009, 122, 119-129.	1.6	8
26	Chromosomal assignment of the porcine NALP5 gene, a candidate gene for female reproductive traits. Animal Reproduction Science, 2009, 112, 397-401.	1.5	7
27	Construction and validation of a novel dual reporter vector for studying mammalian bidirectional promoters. Plasmid, 2014, 74, 1-8.	1.4	6
28	Bovine TWINKLE and mitochondrial ribosomal protein L43 genes are regulated by an evolutionary conserved bidirectional promoter. Gene, 2014, 537, 154-163.	2.2	5
29	HSPD1 is not a major susceptibility gene for rheumatoid arthritis in the French Caucasian population. Journal of Human Genetics, 2007, 52, 1036-1039.	2.3	3
30	Genetic variability of the activity of bidirectional promoters: a pilot study in bovine muscle. DNA Research, 2017, 24, 221-233.	3.4	2
31	Allelic variation of the porcine α-1,3-galactosyltransferase 1 (GGTA1) gene. Journal of Applied Genetics, 2008, 49, 75-79.	1.9	o