List of Publications by Year in descending order

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LUCA FEDDETTI

#	Article	IF	CITATIONS
1	A medium-density genetic linkage map of the bovine genome. Mammalian Genome, 1997, 8, 21-28.	2.2	313
2	Mitochondrial genomes of extinct aurochs survive in domestic cattle. Current Biology, 2008, 18, R157-R158.	3.9	231
3	Total synthesis of a gene for bovine rhodopsin Proceedings of the National Academy of Sciences of the United States of America, 1986, 83, 599-603.	7.1	196
4	The Multifaceted Origin of Taurine Cattle Reflected by the Mitochondrial Genome. PLoS ONE, 2009, 4, e5753.	2.5	157
5	Long range restriction analysis of the bovine casein genes. Nucleic Acids Research, 1990, 18, 6829-6833.	14.5	156
6	A bovine whole-genome radiation hybrid panel and outline map. Mammalian Genome, 2002, 13, 469-474.	2.2	115
7	Origin and Spread of Bos taurus: New Clues from Mitochondrial Genomes Belonging to Haplogroup T1. PLoS ONE, 2012, 7, e38601.	2.5	93
8	The Enigmatic Origin of Bovine mtDNA Haplogroup R: Sporadic Interbreeding or an Independent Event of Bos primigenius Domestication in Italy?. PLoS ONE, 2010, 5, e15760.	2.5	84
9	Chromosomal localization and molecular characterization of 53 cosmid-derived bovine microsatellites. Mammalian Genome, 1995, 6, 629-635.	2.2	71
10	Complete genomic sequence of the bovine prion gene (PRNP) and polymorphism in its promoter region. Animal Genetics, 2001, 32, 231-232.	1.7	57
11	Lactobacillus protoplast transformation. Plasmid, 1987, 17, 73-75.	1.4	54
12	Chromosome evolution and improved cytogenetic maps of the Y chromosome in cattle, zebu, river buffalo, sheep and goat. Chromosome Research, 2005, 13, 349-355.	2.2	54
13	The Paleo-Indian Entry into South America According to Mitogenomes. Molecular Biology and Evolution, 2018, 35, 299-311.	8.9	54
14	Meta-Analysis of Mitochondrial DNA Reveals Several Population Bottlenecks during Worldwide Migrations of Cattle. Diversity, 2014, 6, 178-187.	1.7	51
15	The river buffalo (<i>Bubalus bubalis</i> , 2n = 50) cytogenetic map: assignment of 64 loci by fluorescence in situ hybridization and R-banding. Cytogenetic and Genome Research, 2003, 102, 65-75.	1.1	49
16	Genomic organization, comparative analysis, and genetic polymorphisms of the bovine and ovine prion Doppel genes (PRND). Mammalian Genome, 2001, 12, 729-733.	2.2	43
17	Mitogenomes from Egyptian Cattle Breeds: New Clues on the Origin of Haplogroup Q and the Early Spread of Bos taurus from the Near East. PLoS ONE, 2015, 10, e0141170.	2.5	41
18	Telomeric fusions in cultured human fibroblasts as a source of genomic instability. Cancer Genetics and Cytogenetics, 1997, 95, 130-136.	1.0	40

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19	Mapping human dispersals into the Horn of Africa from Arabian Ice Age refugia using mitogenomes. Scientific Reports, 2016, 6, 25472.	3.3	40
20	A Novel in-Frame 18-bp Microdeletion in <i>MT-CYB</i> Causes a Multisystem Disorder with Prominent Exercise Intolerance. Human Mutation, 2014, 35, 954-958.	2.5	38
21	Combined Q banding and fluorescence in situ hybridization for the identification of bovine chromosomes 1 to 7. Cytogenetic and Genome Research, 1995, 69, 1-6.	1.1	37
22	Analysis of the human Y-chromosome haplogroup Q characterizes ancient population movements in Eurasia and the Americas. BMC Biology, 2019, 17, 3.	3.8	36
23	Cosmid-derived markers anchoring the bovine genetic map to the physical map. Mammalian Genome, 1997, 8, 29-36.	2.2	34
24	Sequence variation in the bovine and ovine PRNP genes. Animal Genetics, 2003, 34, 183-190.	1.7	34
25	Specific and reversible inhibition of the blunt end joining activity of the T4 DNA ligase. Nucleic Acids Research, 1981, 9, 3695-3705.	14.5	31
26	Mitochondrial DNA variants of Podolian cattle breeds testify for a dual maternal origin. PLoS ONE, 2018, 13, e0192567.	2.5	30
27	Temperature dependence of the joining by T4 DNA ligase of termini produced by type II restriction endonucleases. Nucleic Acids Research, 1981, 9, 85-93.	14.5	29
28	A repeated chromosomal DNA sequence is amplified as a circular extrachromosomal molecule in rice (Oryza sativa L.). Molecular Genetics and Genomics, 1990, 222, 58-64.	2.4	24
29	Restriction fragment length polymorphism analysis of the κâ€casein locus in cattle. Animal Genetics, 1990, 21, 107-114.	1.7	23
30	An advanced sheep (<i>Ovis aries</i> , 2 <i>n</i> = 54) cytogenetic map and assignment of 88 new autosomal loci by fluorescence <i>in situ</i> hybridization and Râ€banding. Animal Genetics, 2007, 38, 233-240.	1.7	23
31	Exploring the Y Chromosomal Ancestry of Modern Panamanians. PLoS ONE, 2015, 10, e0144223.	2.5	20
32	The origin of a morphologically unidentifiable human supernumerary minichromosome traced through sorting, molecular cloning, and in situ hybridisation Journal of Medical Genetics, 1991, 28, 92-96.	3.2	19
33	Cloning of the bovine prion-like Shadoo (SPRN) gene by comparative analysis of the predicted genomic locus. Mammalian Genome, 2006, 17, 1130-1139.	2.2	19
34	Reconstructing the genetic history of Italians: new insights from a male (Y-chromosome) perspective. Annals of Human Biology, 2018, 45, 44-56.	1.0	19
35	Interaction between the cellular prion (PrPC) and the 2P domain K+ channel TREK-1 protein. Biochemical and Biophysical Research Communications, 2006, 346, 108-115.	2.1	18
36	Increased SCE levels in Mediterranean Italian buffaloes affected by limb malformation (transversal) Tj ETQq0 0 () rgBT/Ove	rlock 10 Tf 50

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37	On the origin and diversification of Podolian cattle breeds: testing scenarios of European colonization using genome-wide SNP data. Genetics Selection Evolution, 2021, 53, 48.	3.0	18
38	Comparative mapping of the prion gene (PRNP) locus in cattle, sheep and human with PCR-generated probes. Mammalian Genome, 1998, 9, 853-855.	2.2	17
39	Mapping of syntenic groups U7 and U27 to bovine Chromosomes 25 and 12, respectively. Mammalian Genome, 1994, 5, 574-576.	2.2	16
40	Six antimicrobial peptide genes of the cathelicidin family map to bovine chromosome 22q24 by fluorescence in situ hybridization. Cytogenetic and Genome Research, 1996, 75, 240-242.	1.1	16
41	Application of Quantitative Real-Time PCR in the Detection of Prion-Protein Gene Species-Specific DNA Sequences in Animal Meals and Feedstuffs. Journal of Food Protection, 2006, 69, 891-896.	1.7	16
42	Isolation and molecular characterization of rasfadin, a novel gene in the vicinity of the bovine prion gene. Mammalian Genome, 2001, 12, 150-156.	2.2	15
43	Diagnostic value of PRND gene expression profiles in astrocytomas: Relationship to tumor grades of malignancy. Oncology Reports, 2007, 17, 989-96.	2.6	15
44	Differential expression of the prion-like protein doppel gene (PRND) in astrocytomas: a new molecular marker potentially involved in tumor progression. Anticancer Research, 2004, 24, 1507-17.	1.1	15
45	A novel <scp> <i>USP9Y</i> </scp> polymorphism allowing a rapid and unambiguous classification of <scp> <i>Bos taurus</i> Y</scp> chromosomes into haplogroups. Animal Genetics, 2012, 43, 611-613.	1.7	14
46	Comparative mapping of the fragile histidine triad (FHIT) gene in cattle, river buffalo, sheep and goat by FISH and assignment to BTA22 by RH-mapping: a comparison with HSA3. Animal Genetics, 2005, 36, 363-364.	1.7	13
47	Sequence and functional analysis of a divergent promoter from a cryptic plasmid of Lactobacillus acidophilus 168 S. Plasmid, 1987, 17, 69-72.	1.4	12
48	Five polymorphic bovine microsatellite loci: <i>IDVGAâ€62A, IDVGAâ€71, IDVGAâ€82, IDVGAâ€88, IDVGAâ€90<!--<br-->Animal Genetics, 1995, 26, 365-366.</i>	^{i>} 1.7	12
49	Prion-like Doppel gene (PRND) in the goat: genomic structure, cDNA, and polymorphisms. Mammalian Genome, 2005, 16, 963-971.	2.2	11
50	Altered cellular distribution and sub-cellular sorting of doppel (Dpl) protein in human astrocytoma cell lines. Cellular Oncology, 2008, 30, 337-47.	1.9	11
51	Overexpression of the Doppel protein in acute myeloid leukaemias and myelodysplastic syndromes. British Journal of Haematology, 2005, 128, 877-884.	2.5	10
52	FISH Mapping of Bovine U21, U1 and U7 Molecular Markers to River Buffalo Chromosomes 3p, 5q and 5p. Chromosome Research, 1997, 5, 337-340.	2.2	8
53	Survey of uniparental genetic markers in the Maltese cattle breed reveals a significant founder effect but does not indicate local domestication. Animal Genetics, 2016, 47, 267-269.	1.7	8
54	Molecular cloning of DNA from a sorted human minichromosome. Gene, 1991, 99, 229-234.	2.2	7

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55	Functional mapping of the bovine Doppel gene promoter region. Gene, 2005, 356, 101-108.	2.2	7
56	Comparative genomic mapping of the bovine Fragile Histidine Triad (FHIT) tumour suppressor gene: characterization of a 2 Mb BAC contig covering the locus, complete annotation of the gene, analysis of cDNA and of physiological expression profiles. BMC Genomics, 2006, 7, 123.	2.8	6
57	Molecular analysis of the fragile histidine triad <i>(FHIT)</i> tumor suppressor gene in vesical tumors of cattle with chronic enzootic hematuria (CEH). Cytogenetic and Genome Research, 2008, 120, 173-177.	1.1	6
58	Convergent transcription of the Escherichia coll hlsC gene cloned in Bacillus subtllls stops in the vicinity of the attenuator. Gene, 1984, 29, 11-19.	2.2	5
59	lsolation of coding sequences from bovine cosmids by means of exon trapping. Mammalian Genome, 1997, 8, 486-490.	2.2	5
60	Characterization and mapping of three bovine polymorphic microsatellite loci. Animal Genetics, 1996, 27, 121-132.	1.7	5
61	Y-chromosome and Surname Analyses for Reconstructing Past Population Structures: The Sardinian Population as a Test Case. International Journal of Molecular Sciences, 2019, 20, 5763.	4.1	5
62	Overview of the Americas' First Peopling from a Patrilineal Perspective: New Evidence from the Southern Continent. Genes, 2022, 13, 220.	2.4	5
63	The Mitogenome Relationships and Phylogeography of Barn Swallows (<i>Hirundo rustica</i>). Molecular Biology and Evolution, 2022, 39, .	8.9	4
64	Physical mapping of connexin 32 (GJB1) and 43 (GJA1) genes to bovine Chromosomes Xq22 and 9q15/16 by fluorescence in situ hybridization. Mammalian Genome, 1996, 7, 634-635.	2.2	3
65	Six bovine cosmid-derived microsatellites mapping different syntenic groups are fluorescence in situ hybridization mapped to six river buffalo chromosomes. Chromosome Research, 1997, 5, 541-543.	2.2	3
66	Weaving Mitochondrial DNA and Y-Chromosome Variation in the Panamanian Genetic Canvas. Genes, 2021, 12, 1921.	2.4	3
67	Isolation and mapping of a polymorphic DNA sequence (ÂMC.34) on chromosome 2 [D2S63]. Nucleic Acids Research, 1988, 16, 9061-9061.	14.5	2
68	Preparation of high molecular weight plant DNA and its use for artificial chromosome construction. Plant Cell Reports, 1991, 10, 315-20.	5.6	2
69	Construction of a library of bovine genomic fragments enriched in CpG islands. Animal Genetics, 1993, 24, 1-7.	1.7	2
70	Two additional Mspl RFLPs revealed by MC.34 (D2S63). Nucleic Acids Research, 1991, 19, 6345-6345.	14.5	1
71	Eight molecular markers from bovine syntenic groups U2, U5, U24, U14, U12, U28, X and Y were fluorescence in situ mapped to eight river buffalo chromosomes. Chromosome Research, 1998, 6, 656-659.	2.2	1
72	Physical and genetic mapping of two polymorphic bovine dinucleotide repeats: IOBT450 (D6S31) and IDVGA80 (D2S46). Animal Genetics, 1996, 27, 377-378.	1.7	1

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73	Heterologous expression in Bacillus subtilis II. In vitro removal of the attenuator sequence of the Escherichia coli his operon allows expression of the cloned hisG gene in B. subtilis. Gene, 1986, 47, 279-286.	2.2	0
74	T-banding pattern of bovine chromosomes and karyotype reconstitution with physically mapped cosmids. Cytogenetic and Genome Research, 1996, 73, 229-234.	1.1	0
75	Assignment <footref rid="foot01">¹</footref> of interleukin-1 receptor, type I (IL1R1) to bovine chromosome band 11q12 by in situ hybridization. Cytogenetic and Genome Research, 2000, 89, 166-167.	1.1	0