Salvatore Mastrangelo

List of Publications by Year in descending order

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75 papers 1,642 citations

361045 20 h-index 377514 34 g-index

78 all docs

78 docs citations

78 times ranked 1302 citing authors

#	Article	IF	CITATIONS
1	Genome-wide scan for runs of homozygosity identifies potential candidate genes associated with local adaptation in Valle del Belice sheep. Genetics Selection Evolution, 2017, 49, 84.	1.2	146
2	Genomic inbreeding estimation in small populations: evaluation of runs of homozygosity in three local dairy cattle breeds. Animal, 2016, 10, 746-754.	1.3	129
3	Runs of homozygosity reveal genomeâ€wide autozygosity in Italian sheep breeds. Animal Genetics, 2018, 49, 71-81.	0.6	67
4	Genome-wide identification of runs of homozygosity islands and associated genes in local dairy cattle breeds. Animal, 2018, 12, 2480-2488.	1.3	65
5	On the origin of European sheep as revealed by the diversity of the Balkan breeds and by optimizing population-genetic analysis tools. Genetics Selection Evolution, 2020, 52, 25.	1.2	58
6	Genome-Wide Variation, Candidate Regions and Genes Associated With Fat Deposition and Tail Morphology in Ethiopian Indigenous Sheep. Frontiers in Genetics, 2018, 9, 699.	1.1	56
7	Conservation status and historical relatedness of Italian cattle breeds. Genetics Selection Evolution, 2018, 50, 35.	1.2	50
8	A genomic map of climate adaptation in Mediterranean cattle breeds. Molecular Ecology, 2019, 28, 1009-1029.	2.0	46
9	Genome-wide analysis in endangered populations: a case study in Barbaresca sheep. Animal, 2017, 11, 1107-1116.	1.3	45
10	Novel and known signals of selection for fat deposition in domestic sheep breeds from Africa and Eurasia. PLoS ONE, 2019, 14, e0209632.	1.1	43
11	Genetic diversity and population structure of Sicilian sheep breeds using microsatellite markers. Small Ruminant Research, 2012, 102, 18-25.	0.6	41
12	Genome wide linkage disequilibrium and genetic structure in Sicilian dairy sheep breeds. BMC Genetics, 2014, 15, 108.	2.7	33
13	Application of microsatellite markers as potential tools for traceability of Girgentana goat breed dairy products. Food Research International, 2015, 74, 115-122.	2.9	33
14	The genetic heritage of Alpine local cattle breeds using genomic SNP data. Genetics Selection Evolution, 2020, 52, 40.	1.2	32
15	Insights into Genetic Diversity, Runs of Homozygosity and Heterozygosity-Rich Regions in Maremmana Semi-Feral Cattle Using Pedigree and Genomic Data. Animals, 2020, 10, 2285.	1.0	32
16	The genome-wide structure of two economically important indigenous Sicilian cattle breeds1. Journal of Animal Science, 2014, 92, 4833-4842.	0.2	31
17	Genome-wide association study between CNVs and milk production traits in Valle del Belice sheep. PLoS ONE, 2019, 14, e0215204.	1.1	31
18	Genome-wide scan of fat-tail sheep identifies signals of selection for fat deposition and adaptation. Animal Production Science, 2019, 59, 835.	0.6	29

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19	Genome-Wide SNP Analysis Reveals the Population Structure and the Conservation Status of 23 Italian Chicken Breeds. Animals, 2020, 10, 1441.	1.0	28
20	Preselection statistics and Random Forest classification identify population informative single nucleotide polymorphisms in cosmopolitan and autochthonous cattle breeds. Animal, 2018, 12, 12-19.	1.3	25
21	A Combined Multi-Cohort Approach Reveals Novel and Known Genome-Wide Selection Signatures for Wool Traits in Merino and Merino-Derived Sheep Breeds. Frontiers in Genetics, 2019, 10, 1025.	1.1	24
22	Genome-wide scan for selection signatures reveals novel insights into the adaptive capacity in local North African cattle. Scientific Reports, 2020, 10, 19466.	1.6	24
23	The climatic and genetic heritage of Italian goat breeds with genomic SNP data. Scientific Reports, 2021, 11, 10986.	1.6	23
24	Genomeâ€wide detection of signatures of selection in three Valdostana cattle populations. Journal of Animal Breeding and Genetics, 2020, 137, 609-621.	0.8	22
25	Genomeâ€wide assessment of diversity and differentiation between original and modern Brown cattle populations. Animal Genetics, 2021, 52, 21-31.	0.6	20
26	Refining the genetic structure and relationships of European cattle breeds through meta-analysis of worldwide genomic SNP data, focusing on Italian cattle. Scientific Reports, 2020, 10, 14522.	1.6	19
27	Fifteen Shades of Grey: Combined Analysis of Genome-Wide SNP Data in Steppe and Mediterranean Grey Cattle Sheds New Light on the Molecular Basis of Coat Color. Genes, 2020, 11, 932.	1.0	19
28	Genome-wide analysis reveals the patterns of genetic diversity and population structure of 8 Italian local chicken breeds. Poultry Science, 2021, 100, 441-451.	1.5	18
29	Genome-wide analyses reveal population structure and identify candidate genes associated with tail fatness in local sheep from a semi-arid area. Animal, 2021, 15, 100193.	1.3	18
30	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.	1.2	18
31	The genetics of phenotypic plasticity in livestock in the era of climate change: a review. Italian Journal of Animal Science, 2020, 19, 997-1014.	0.8	17
32	Quantitative determination of casein genetic variants in goat milk: Application in Girgentana dairy goat breed. Food Chemistry, 2016, 192, 760-764.	4.2	16
33	Genetic polymorphism at the CSN1S1 gene in Girgentana dairy goat breed. Animal Production Science, 2013, 53, 403.	0.6	15
34	Genome-wide diversity and runs of homozygosity in the "Braque Français, type Pyrénées―dog breed. BMC Research Notes, 2018, 11, 13.	0.6	15
35	Runs of homozygosity in the Italian goat breeds: impact of management practices in low-input systems. Genetics Selection Evolution, 2021, 53, 92.	1.2	15
36	Study of polymorphisms in the promoter region of ovine \hat{I}^2 -lactoglobulin gene and phylogenetic analysis among the Valle del Belice breed and other sheep breeds considered as ancestors. Molecular Biology Reports, 2012, 39, 745-751.	1.0	14

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37	Genomeâ€wide association studies for milk production traits in Valle del Belice sheep using repeated measures. Animal Genetics, 2019, 50, 311-314.	0.6	14
38	Genomeâ€wide analyses reveal the regions involved in the phenotypic diversity in Sicilian pigs. Animal Genetics, 2020, 51, 101-105.	0.6	14
39	Genome-Wide Analyses Identifies Known and New Markers Responsible of Chicken Plumage Color. Animals, 2020, 10, 493.	1.0	13
40	Genome-Wide Association Study Identifies New Candidate Markers for Somatic Cells Score in a Local Dairy Sheep. Frontiers in Genetics, 2021, 12, 643531.	1.1	13
41	Genome-Wide Patterns of Homozygosity Reveal the Conservation Status in Five Italian Goat Populations. Animals, 2021, 11, 1510.	1.0	13
42	Parentage verification of Valle del Belice dairy sheep using multiplex microsatellite panel. Small Ruminant Research, 2013, 113, 62-65.	0.6	12
43	Genetic Characterisation of CSN2Gene in Girgentana Goat Breed. Italian Journal of Animal Science, 2014, 13, 3414.	0.8	12
44	Determination of milk production losses and variations of fat and protein percentages according to different levels of somatic cell count in Valle del Belice dairy sheep. Small Ruminant Research, 2018, 162, 39-42.	0.6	12
45	Genomic characterization of Algerian Guelmoise cattle and their genetic relationship with other North African populations inferred from SNP genotyping arrays. Livestock Science, 2018, 217, 19-25.	0.6	12
46	Genetic structure of Tunisian sheep breeds as inferred from genome-wide SNP markers. Small Ruminant Research, 2020, 191, 106192.	0.6	12
47	Comparative selection signature analyses identify genomic footprints in Reggiana cattle, the traditional breed of the Parmigiano-Reggiano cheese production system. Animal, 2020, 14, 921-932.	1.3	12
48	Identification of Copy Number Variations and Genetic Diversity in Italian Insular Sheep Breeds. Animals, 2022, 12, 217.	1.0	12
49	Association study between \hat{I}^2 -defensin gene polymorphisms and mastitis resistance in Valle del Belice dairy sheep breed. Small Ruminant Research, 2016, 136, 18-21.	0.6	11
50	Combined approaches to identify genomic regions involved in phenotypic differentiation between low divergent breeds: Application in Sardinian sheep populations. Journal of Animal Breeding and Genetics, 2019, 136, 526-534.	0.8	11
51	Genomeâ€wide association study reveals the locus responsible for microtia in Valle del Belice sheep breed. Animal Genetics, 2018, 49, 636-640.	0.6	10
52	Weighted Single-Step Genome-Wide Association Study Uncovers Known and Novel Candidate Genomic Regions for Milk Production Traits and Somatic Cell Score in Valle del Belice Dairy Sheep. Animals, 2022, 12, 1155.	1.0	10
53	Molecular Characterisation ofl®â€"CaseinGene inGirgentanaDairy Goat Breed and Identification of Two New Alleles. Italian Journal of Animal Science, 2015, 14, 3464.	0.8	9
54	Genomic characterization of the Braque Français type Pyrénées dog and relationship with other breeds. PLoS ONE, 2018, 13, e0208548.	1.1	9

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55	Genome-wide detection of copy-number variations in local cattle breeds. Animal Production Science, 2019, 59, 815.	0.6	9
56	12S rRNA mitochondrial gene as marker to trace Sicilian mono-species dairy products. Livestock Science, 2016, 193, 39-44.	0.6	8
57	Genome-Wide Analysis Reveals Selection Signatures Involved in Meat Traits and Local Adaptation in Semi-Feral Maremmana Cattle. Frontiers in Genetics, 2021, 12, 675569.	1.1	8
58	Genetic Variability atl±s2-caseinGene inGirgentanaDairy Goat Breed. Italian Journal of Animal Science, 2014, 13, 2997.	0.8	7
59	Development and validation of RP-HPLC method for the quantitative estimation of $\hat{l}\pm s1$ -genetic variants in goat milk. Food Chemistry, 2014, 156, 165-169.	4.2	7
60	A combined genome-wide approach identifies a new potential candidate marker associated with the coat color sidedness in cattle. Livestock Science, 2019, 225, 91-95.	0.6	7
61	A Genome-Wide Detection of Copy Number Variations Using SNP Genotyping Arrays in Braque Français Type Pyrénées Dogs. Animals, 2019, 9, 77.	1.0	7
62	Variation of proteomic profile during lactation in Girgentana goat milk: a preliminary study. Italian Journal of Animal Science, 2019, 18, 88-97.	0.8	7
63	Genomic Structural Diversity in Local Goats: Analysis of Copy-Number Variations. Animals, 2020, 10, 1040.	1.0	7
64	Genome-wide diversity of Pagliarola sheep residual population and its conservation implication. Italian Journal of Animal Science, 2021, 20, 1695-1705.	0.8	7
65	Genomeâ€wide survey on three local horse populations with a focus on runs of homozygosity pattern. Journal of Animal Breeding and Genetics, 2022, 139, 540-555.	0.8	7
66	Detection of genomic regions underlying milk production traits in Valle del Belice dairy sheep using regional heritability mapping. Journal of Animal Breeding and Genetics, 2021, 138, 552-561.	0.8	6
67	Full-length sequencing and identification of novel polymorphisms in the ACACA gene of Valle del Belice sheep breed. Journal of Genetics, 2017, 96, 591-597.	0.4	5
68	Genomeâ€wide analysis identifies potentially causative genes explaining the phenotypic variability in Pinzirita sheep. Animal Genetics, 2019, 50, 189-190.	0.6	5
69	High-Density Genomic Characterization of Native Croatian Sheep Breeds. Frontiers in Genetics, 0, 13 , .	1.1	5
70	Population genetic structure and milk production traits in Girgentana goat breed. Animal Production Science, 2017, 57, 430.	0.6	4
71	Penalized classification for optimal statistical selection of markers from high-throughput genotyping: application in sheep breeds. Animal, 2018, 12, 1118-1125.	1.3	3
72	Effect of hairless gene polymorphism on the breeding values of milk production traits in Valle del Belice dairy sheep. Livestock Science, 2013, 154, 60-63.	0.6	1

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73	The Sicilian rock partridge: latest data on genetic integrity from four different relict areas. Turkish Journal of Zoology, 0, , .	0.4	1
74	The Girgentana Goat Breed: A Zootechnical Overview on Genetics, Nutrition and Dairy Production Aspects., 2017,, 191-203.		1
75	Genome-wide association study for milk production traits in an economically important local dairy sheep breed. Italian Journal of Animal Science, 2021, 20, 1500-1505.	0.8	0