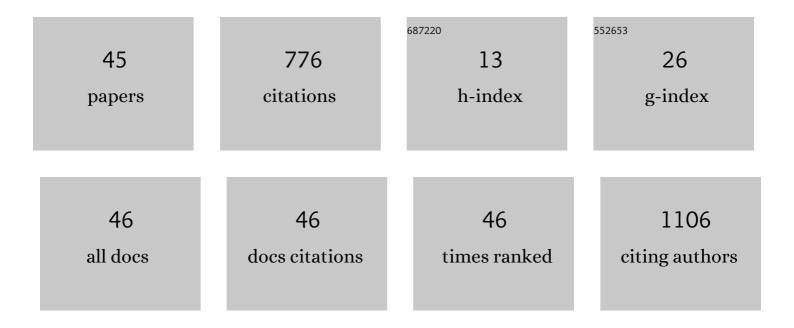
## Maria Teresa Sardina

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2168434/publications.pdf Version: 2024-02-01



| #  | Article   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | Identification of Copy Number Variations and Genetic Diversity in Italian Insular Sheep Breeds. Animals, 2022, 12, 217.   | 1.0 | 12        |
| 2  | Development of "Quadrello di Ovinoâ€, a Novel Fresh Ewe's Cheese. Foods, 2022, 11, 25.  | 1.9 | 2         |
| 3  | Weighted Single-Step Genome-Wide Association Study Uncovers Known and Novel Candidate Genomic<br>Regions for Milk Production Traits and Somatic Cell Score in Valle del Belice Dairy Sheep. Animals,<br>2022, 12, 1155.     | 1.0 | 10        |
| 4  | Genome-Wide Association Study Identifies New Candidate Markers for Somatic Cells Score in a Local<br>Dairy Sheep. Frontiers in Genetics, 2021, 12, 643531.  | 1.1 | 13        |
| 5  | Genome-Wide Patterns of Homozygosity Reveal the Conservation Status in Five Italian Goat<br>Populations. Animals, 2021, 11, 1510.   | 1.0 | 13        |
| 6  | 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         |
| 7  | 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         |
| 8  | The Root Mycobiota of Betula aetnensis Raf., an Endemic Tree Species Colonizing the Lavas of Mt. Etna<br>(Italy). Forests, 2021, 12, 1624.  | 0.9 | 3         |
| 9  | Genomic Structural Diversity in Local Goats: Analysis of Copy-Number Variations. Animals, 2020, 10,<br>1040.  | 1.0 | 7         |
| 10 | Genome-wide detection of copy-number variations in local cattle breeds. Animal Production Science, 2019, 59, 815.   | 0.6 | 9         |
| 11 | 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         |
| 12 | Evaluation of microbiological and physicoâ€chemical parameters of retail readyâ€toâ€eat monoâ€varietal<br>salads. Journal of Food Processing and Preservation, 2019, 43, e13955.  | 0.9 | 6         |
| 13 | Genome-wide association study between CNVs and milk production traits in Valle del Belice sheep.<br>PLoS ONE, 2019, 14, e0215204.   | 1.1 | 31        |
| 14 | A Genome-Wide Detection of Copy Number Variations Using SNP Genotyping Arrays in Braque Français<br>Type Pyrénées Dogs. Animals, 2019, 9, 77.   | 1.0 | 7         |
| 15 | Variation of proteomic profile during lactation in Girgentana goat milk: a preliminary study. Italian<br>Journal of Animal Science, 2019, 18, 88-97.  | 0.8 | 7         |
| 16 | Determination of milk production losses and variations of fat and protein percentages according to<br>different levels of somatic cell count in Valle del Belice dairy sheep. Small Ruminant Research, 2018,<br>162, 39-42. | 0.6 | 12        |
| 17 | Population genetic structure and milk production traits in Girgentana goat breed. Animal Production Science, 2017, 57, 430.   | 0.6 | 4         |
| 18 | Full-length sequencing and identification of novel polymorphisms in the ACACA gene of Valle del<br>Belice sheep breed. Journal of Genetics, 2017, 96, 591-597.  | 0.4 | 5         |

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|----|--|-----|-----------|
| 19 | Genome-wide scan for runs of homozygosity identifies potential candidate genes associated with<br>local adaptation in Valle del Belice sheep. Genetics Selection Evolution, 2017, 49, 84.      | 1.2 | 146       |
| 20 | Whole genome semiconductor based sequencing of farmed European sea bass (Dicentrarchus labrax)<br>Mediterranean genetic stocks using a DNA pooling approach. Marine Genomics, 2016, 28, 63-70. | 0.4 | 11        |
| 21 | 12S rRNA mitochondrial gene as marker to trace Sicilian mono-species dairy products. Livestock<br>Science, 2016, 193, 39-44.   | 0.6 | 8         |
| 22 | Microbial Activation of Wooden Vats Used for Traditional Cheese Production and Evolution of Neoformed Biofilms. Applied and Environmental Microbiology, 2016, 82, 585-595.                     | 1.4 | 41        |
| 23 | Valorization of indigenous dairy cattle breed through salami production. Meat Science, 2016, 114, 58-68.   | 2.7 | 13        |
| 24 | Association study between β-defensin gene polymorphisms and mastitis resistance in Valle del Belice<br>dairy sheep breed. Small Ruminant Research, 2016, 136, 18-21.                           | 0.6 | 11        |
| 25 | Quantitative determination of casein genetic variants in goat milk: Application in Girgentana dairy goat breed. Food Chemistry, 2016, 192, 760-764.  | 4.2 | 16        |
| 26 | Whole mitochondrial genomes unveil the impact of domestication on goat matrilineal variability.<br>BMC Genomics, 2015, 16, 1115.   | 1.2 | 56        |
| 27 | Molecular Characterisation ofl̂ºâ€"CaseinGene inGirgentanaDairy Goat Breed and Identification of Two<br>New Alleles. Italian Journal of Animal Science, 2015, 14, 3464.                        | 0.8 | 9         |
| 28 | 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        |
| 29 | Genetic Variability atαs2-caseinGene inGirgentanaDairy Goat Breed. Italian Journal of Animal Science,<br>2014, 13, 2997.   | 0.8 | 7         |
| 30 | Genome wide linkage disequilibrium and genetic structure in Sicilian dairy sheep breeds. BMC Genetics, 2014, 15, 108.  | 2.7 | 33        |
| 31 | Genetic Characterisation ofCSN2Gene inGirgentanaGoat Breed. Italian Journal of Animal Science, 2014, 13, 3414.   | 0.8 | 12        |
| 32 | Development and validation of RP-HPLC method for the quantitative estimation of αs1-genetic variants<br>in goat milk. Food Chemistry, 2014, 156, 165-169.                                      | 4.2 | 7         |
| 33 | Antibacterial activity of Borago officinalis and Brassica juncea aqueous extracts evaluated inÂvitro<br>and in situ using different food model systems. Food Control, 2014, 40, 157-164.       | 2.8 | 43        |
| 34 | The genome-wide structure of two economically important indigenous Sicilian cattle breeds1. Journal of Animal Science, 2014, 92, 4833-4842.  | 0.2 | 31        |
| 35 | Parentage verification of Valle del Belice dairy sheep using multiplex microsatellite panel. Small<br>Ruminant Research, 2013, 113, 62-65.   | 0.6 | 12        |
| 36 | Effect of hairless gene polymorphism on the breeding values of milk production traits in Valle del<br>Belice dairy sheep. Livestock Science, 2013, 154, 60-63.                                 | 0.6 | 1         |

MARIA TERESA SARDINA

| #  | Article  | IF  | CITATIONS |
|----|--|-----|-----------|
| 37 | Genetic polymorphism at the CSN1S1 gene in Girgentana dairy goat breed. Animal Production Science, 2013, 53, 403.  | 0.6 | 15        |
| 38 | Study of polymorphisms in the promoter region of ovine β-lactoglobulin gene and phylogenetic<br>analysis among the Valle del Belice breed and other sheep breeds considered as ancestors. Molecular<br>Biology Reports, 2012, 39, 745-751. | 1.0 | 14        |
| 39 | Polymorphisms of beta-lactoglobulin promoter region in three Sicilian goat breeds. Molecular<br>Biology Reports, 2012, 39, 3203-3210.  | 1.0 | 10        |
| 40 | Identification of SNPs in the promoter of β-lactoglobulin gene in three Sicilian goat breeds. Italian<br>Journal of Animal Science, 2009, 8, 147-149.  | 0.8 | 1         |
| 41 | Short Communication: Casein Haplotype Variability in Sicilian Dairy Goat Breeds. Journal of Dairy Science, 2008, 91, 3687-3692.  | 1.4 | 23        |
| 42 | Polymorphism and chromosomal localization of the porcine signal transducer and activator of transcription 5B gene (STAT5B). Journal of Animal Breeding and Genetics, 2006, 123, 284-287.   | 0.8 | 5         |
| 43 | Phylogenetic analysis of Sicilian goats reveals a new mtDNA lineage. Animal Genetics, 2006, 37, 376-378.   | 0.6 | 62        |
| 44 | Assignment of Signal Transducer and Activator of Transcription 5A <i>(STAT5A)</i> gene to porcine<br>chromosome 12p13→p11 by radiation hybrid panel mapping. Cytogenetic and Genome Research, 2006, 112,<br>342J-342J.                     | 0.6 | 1         |
| 45 | Phenotypic and genetic analysis of udder health using SCC in Valle del Belice dairy sheep. Italian<br>Journal of Animal Science, 2005, 4, 76-78.   | 0.8 | 1         |