## Martina Lari

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	The genetic history of Ice Age Europe. Nature, 2016, 534, 200-205.	27.8	729
2	A Revised Timescale for Human Evolution Based on Ancient Mitochondrial Genomes. Current Biology, 2013, 23, 553-559.	3.9	540
3	The genomic history of southeastern Europe. Nature, 2018, 555, 197-203.	27.8	479
4	Evidence for a genetic discontinuity between Neandertals and 24,000-year-old anatomically modern Europeans. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 6593-6597.	7.1	324
5	Pleistocene Mitochondrial Genomes Suggest a Single Major Dispersal of Non-Africans and a Late Glacial Population Turnover in Europe. Current Biology, 2016, 26, 827-833.	3.9	277
6	The origin of European cattle: Evidence from modern and ancient DNA. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 8113-8118.	7.1	271
7	Ancient DNA studies: new perspectives on old samples. Genetics Selection Evolution, 2012, 44, 21.	3.0	150
8	Neandertal Evolutionary Genetics: Mitochondrial DNA Data from the Iberian Peninsula. Molecular Biology and Evolution, 2005, 22, 1077-1081.	8.9	139
9	Understanding 6th-century barbarian social organization and migration through paleogenomics. Nature Communications, 2018, 9, 3547.	12.8	111
10	Tracking down Human Contamination in Ancient Human Teeth. Molecular Biology and Evolution, 2006, 23, 1801-1807.	8.9	105
11	The spread of steppe and Iranian-related ancestry in the islands of the western Mediterranean. Nature Ecology and Evolution, 2020, 4, 334-345.	7.8	95
12	A highly divergent mtDNA sequence in a Neandertal individual from Italy. Current Biology, 2006, 16, R630-R632.	3.9	80
13	The Etruscans: A Population-Genetic Study. American Journal of Human Genetics, 2004, 74, 694-704.	6.2	72
14	The Illumina Sequencing Protocol and the NovaSeq 6000 System. Methods in Molecular Biology, 2021, 2242, 15-42.	0.9	68
15	Specific inactivation of two immunomodulatory <i>SIGLEC</i> genes during human evolution. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 9935-9940.	7.1	64
16	Monitoring DNA Contamination in Handled vs. Directly Excavated Ancient Human Skeletal Remains. PLoS ONE, 2013, 8, e52524.	2.5	58
17	Genealogical Relationships between Early Medieval and Modern Inhabitants of Piedmont. PLoS ONE, 2015, 10, e0116801.	2.5	58
18	Population dynamic of the extinct European aurochs: genetic evidence of a north-south differentiation pattern and no evidence of post-glacial expansion. BMC Evolutionary Biology, 2010, 10, 83	3.2	51

MARTINA LARI

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19	The origin and legacy of the Etruscans through a 2000-year archeogenomic time transect. Science Advances, 2021, 7, eabi7673.	10.3	44
20	Origins and Evolution of the Etruscans' mtDNA. PLoS ONE, 2013, 8, e55519.	2.5	40
21	The Complete Mitochondrial Genome of an 11,450-year-old Aurochsen (Bos primigenius) from Central Italy. BMC Evolutionary Biology, 2011, 11, 32.	3.2	39
22	A 28,000 Years Old Cro-Magnon mtDNA Sequence Differs from All Potentially Contaminating Modern Sequences. PLoS ONE, 2008, 3, e2700.	2.5	37
23	Unexpected presence of Fagus orientalis complex in Italy as inferred from 45,000-year-old DNA pollen samples from Venice lagoon. BMC Evolutionary Biology, 2007, 7, S6.	3.2	36
24	Complete mitochondrial sequences from Mesolithic Sardinia. Scientific Reports, 2017, 7, 42869.	3.3	35
25	Genetic variation in prehistoric Sardinia. Human Genetics, 2007, 122, 327-336.	3.8	34
26	Genetic analysis of the skeletal remains attributed to Francesco Petrarca. Forensic Science International, 2007, 173, 36-40.	2.2	33
27	The Microcephalin Ancestral Allele in a Neanderthal Individual. PLoS ONE, 2010, 5, e10648.	2.5	31
28	Genealogical Discontinuities among Etruscan, Medieval, and Contemporary Tuscans. Molecular Biology and Evolution, 2009, 26, 2157-2166.	8.9	30
29	Archaeogenomic distinctiveness of the Isthmo-Colombian area. Cell, 2021, 184, 1706-1723.e24.	28.9	30
30	Ancient genomes reveal early Andean farmers selected common beans while preserving diversity. Nature Plants, 2021, 7, 123-128.	9.3	29
31	Mitochondrial DNA from El Mirador Cave (Atapuerca, Spain) Reveals the Heterogeneity of Chalcolithic Populations. PLoS ONE, 2014, 9, e105105.	2.5	28
32	Possible Interbreeding in Late Italian Neanderthals? New Data from the Mezzena Jaw (Monti Lessini,) Tj ETQq0 (	) 0 rgBT /0	verlock 10 Tf
33	Genome diversity in the Neolithic Globular Amphorae culture and the spread of Indo-European languages. Proceedings of the Royal Society B: Biological Sciences, 2017, 284, 20171540.	2.6	24
34	The Neanderthal in the karst: First dating, morphometric, and paleogenetic data on the fossil skeleton from Altamura (Italy). Journal of Human Evolution, 2015, 82, 88-94.	2.6	23
35	The first evidence for Late Pleistocene dogs in Italy. Scientific Reports, 2020, 10, 13313.	3.3	21
36	Did Neandertals and anatomically modern humans coexist in northern Italy during the late MIS 3?.	1.5	17

Quaternary International, 2012, 259, 102-112.

Martina Lari

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37	A genetic perspective on Longobard-Era migrations. European Journal of Human Genetics, 2019, 27, 647-656.	2.8	15
38	Combined methodologies for gaining much information from ancient dental calculus: testing experimental strategies for simultaneously analysing DNA and food residues. Archaeological and Anthropological Sciences, 2020, 12, 1.	1.8	13
39	Kinship Determination in Archeological Contexts Through DNA Analysis. Frontiers in Ecology and Evolution, 2020, 8, .	2.2	13
40	The Biarzo case in northern Italy: is the temporal dynamic of swine mitochondrial DNA lineages in Europe related to domestication?. Scientific Reports, 2015, 5, 16514.	3.3	12
41	From unknown to known: Identification of the remains at the mausoleum of fosse Ardeatine. Science and Justice - Journal of the Forensic Science Society, 2018, 58, 469-478.	2.1	12
42	Ancestral mitochondrial N lineage from the Neolithic â€~green' Sahara. Scientific Reports, 2019, 9, 3530.	3.3	10
43	Insights into the Copper-Bronze Age diet in Central Italy: Plant microremains in dental calculus from Grotta dello Scoglietto (Southern Tuscany, Italy). Journal of Archaeological Science: Reports, 2017, 15, 30-39.	0.5	9
44	The mitogenome portrait of Umbria in Central Italy as depicted by contemporary inhabitants and pre-Roman remains. Scientific Reports, 2020, 10, 10700.	3.3	9
45	Maternal DNA lineages at the gate of Europe in the 10th century AD. PLoS ONE, 2018, 13, e0193578.	2.5	8
46	Evaluation of Diammonium hydrogen phosphate and Ca(OH)2 nanoparticles for consolidation of ancient bones. Journal of Cultural Heritage, 2020, 41, 1-12.	3.3	7
47	Ancient human mitochondrial genomes from Bronze Age Bulgaria: new insights into the genetic history of Thracians. Scientific Reports, 2019, 9, 5412.	3.3	6
48	Successful extraction of insect DNA from recent copal inclusions: limits and perspectives. Scientific Reports, 2021, 11, 6851.	3.3	6
49	Combined metagenomic and archaeobotanical analyses on human dental calculus: A cross-section of lifestyle conditions in a Copper Age population of central Italy. Quaternary International, 2023, 653-654, 69-81.	1.5	6
50	Defining criteria for the reintroduction of locally extinct populations based on contemporary and ancient genetic diversity: The case of the Adriatic Beluga sturgeon ( <i>Huso huso</i> ). Diversity and Distributions, 2021, 27, 816-827.	4.1	5
51	The Mountain Meadows Massacre and "poisoned springsâ€ŧ scientific testing of the more recent, anthrax theory. International Journal of Legal Medicine, 2013, 127, 77-83.	2.2	4
52	DNA Sequencing in Cultural Heritage. Topics in Current Chemistry, 2016, 374, 8.	5.8	4
53	First Bronze Age Human Mitogenomes from Calabria (Grotta Della Monaca, Southern Italy). Genes, 2021, 12, 636.	2.4	4
54	Ancient DNA and forensics genetics: The case of Francesco Petrarca. Forensic Science International: Genetics Supplement Series, 2008, 1, 469-470.	0.3	3

Martina Lari

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55	The female ancestor's tale: Longâ€ŧerm matrilineal continuity in a nonisolated region of Tuscany. American Journal of Physical Anthropology, 2018, 167, 497-506.	2.1	3
56	How a Paleogenomic Approach Can Provide Details on Bioarchaeological Reconstruction: A Case Study from the Globular Amphorae Culture. Genes, 2021, 12, 910.	2.4	3
57	Performance of innovative nanomaterials for bone remains consolidation and effect on 14C dating and on palaeogenetic analysis. Scientific Reports, 2022, 12, 6975.	3.3	3
58	Microcomputed tomography and genetic analysis of a rare case of Caffey's disease in a 5–7â€monthâ€old girl. International Journal of Osteoarchaeology, 2019, 29, 854-859.	1.2	1
59	Whole-exome sequencing of the mummified remains of Cangrande della Scala (1291–1329 CE) indicates the first known case of late-onset Pompe disease. Scientific Reports, 2021, 11, 21070.	3.3	1
60	Ancient and Archaic Genomes. Genes, 2021, 12, 1411.	2.4	0