Alessio Boattini

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2757918/publications.pdf

Version: 2024-02-01

430874 501196 40 885 18 28 citations h-index g-index papers 42 42 42 1477 all docs docs citations times ranked citing authors

| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Genetic history of Calabrian Greeks reveals ancient events and long term isolation in the Aspromonte area of Southern Italy. Scientific Reports, 2021, 11, 3045. | 3.3 | 9 |
| 2 | The surname structure of Trentino (Italy) and its relationship with dialects and genes. Annals of Human Biology, 2021, 48, 260-269. | 1.0 | 3 |
| 3 | Yâ€chromosome variability and genetic history of <scp>Commons</scp> from <scp>Northern Italy</scp> . American Journal of Physical Anthropology, 2021, 175, 665-679. | 2.1 | 4 |
| 4 | Educational briefings in touristic facilities promote tourist sustainable behavior and customer loyalty. Biological Conservation, 2021, 259, 109122. | 4.1 | 5 |
| 5 | Twenty-Seven Y-Chromosome Short Tandem Repeats Analysis of Italian Mummies of the 16th and 18th Centuries: An Interdisciplinary Research. Frontiers in Genetics, 2021, 12, 720640. | 2.3 | 2 |
| 6 | Genomic history of the Italian population recapitulates key evolutionary dynamics of both Continental and Southern Europeans. BMC Biology, 2020, 18, 51. | 3.8 | 26 |
| 7 | Inter-individual genomic heterogeneity within European population isolates. PLoS ONE, 2019, 14, e0214564. | 2.5 | 3 |
| 8 | Estimating Y-Str Mutation Rates and Tmrca Through Deep-Rooting Italian Pedigrees. Scientific Reports, 2019, 9, 9032. | 3.3 | 16 |
| 9 | Dissecting the Pre-Columbian Genomic Ancestry of Native Americans along the Andes–Amazonia Divide. Molecular Biology and Evolution, 2019, 36, 1254-1269. | 8.9 | 47 |
| 10 | The Genetic Variability of APOE in Different Human Populations and Its Implications for Longevity. Genes, 2019, 10, 222. | 2.4 | 96 |
| 11 | A Historical-Genetic Reconstruction of Human Extra-Pair Paternity. Current Biology, 2019, 29, 4102-4107.e7. | 3.9 | 23 |
| 12 | Ripples on the surface. Surnames and genes in Sicily and Southern Italy. Annals of Human Biology, 2018, 45, 57-65. | 1.0 | 5 |
| 13 | Impact of non-LTR retrotransposons in the differentiation and evolution of anatomically modern humans. Mobile DNA, 2018, 9, 28. | 3.6 | 18 |
| 14 | Transposable Elements Activity is Positively Related to Rate of Speciation in Mammals. Journal of Molecular Evolution, 2018, 86, 303-310. | 1.8 | 50 |
| 15 | Overcoming the dichotomy between open and isolated populations using genomic data from a large European dataset. Scientific Reports, 2017, 7, 41614. | 3.3 | 15 |
| 16 | Ancient and recent admixture layers in Sicily and Southern Italy trace multiple migration routes along the Mediterranean. Scientific Reports, 2017, 7, 1984. | 3.3 | 52 |
| 17 | The genetic variation in the R1a clade among the Ashkenazi Levites' Y chromosome. Scientific Reports, 2017, 7, 14969. | 3.3 | 13 |
| 18 | Reply to d'Huy et al.: Navigating biases and charting new ground in the cultural diffusion of folktales. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E8556. | 7.1 | 4 |

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|----|---|-----|-----------|
| 19 | Inferring patterns of folktale diffusion using genomic data. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 9140-9145. | 7.1 | 37 |
| 20 | Mutation Rates and Discriminating Power for 13 Rapidly-Mutating Y-STRs between Related and Unrelated Individuals. PLoS ONE, 2016, 11, e0165678. | 2.5 | 22 |
| 21 | Positive selection of lactase persistence among people of Southern Arabia. American Journal of Physical Anthropology, 2016, 161, 676-684. | 2.1 | 9 |
| 22 | Complex interplay between neutral and adaptive evolution shaped differential genomic background and disease susceptibility along the Italian peninsula. Scientific Reports, 2016, 6, 32513. | 3.3 | 41 |
| 23 | Shared language, diverging genetic histories: high-resolution analysis of Y-chromosome variability in Calabrian and Sicilian Arbereshe. European Journal of Human Genetics, 2016, 24, 600-606. | 2.8 | 16 |
| 24 | Inferring the genetic history of lactase persistence along the Italian peninsula from a large genomic interval surrounding the <scp><i>LCT</i></scp> gene. American Journal of Physical Anthropology, 2015, 158, 708-718. | 2.1 | 7 |
| 25 | Tracing Behçet's disease origins along the Silk Road: an anthropological evolutionary genetics perspective. Clinical and Experimental Rheumatology, 2015, 33, S60-6. | 0.8 | 1 |
| 26 | An Ancient Mediterranean Melting Pot: Investigating the Uniparental Genetic Structure and Population History of Sicily and Southern Italy. PLoS ONE, 2014, 9, e96074. | 2.5 | 55 |
| 27 | Linguistic, geographic and genetic isolation: a collaborative study of Italian populations. Journal of Anthropological Sciences, 2014, 92, 201-31. | 0.4 | 43 |
| 28 | mtDNA variation in East Africa unravels the history of afroâ€asiatic groups. American Journal of Physical Anthropology, 2013, 150, 375-385. | 2.1 | 22 |
| 29 | What can we do with surnames today? An anthropologist point of view. Physics of Life Reviews, 2013, 10, 418-419. | 2.8 | 3 |
| 30 | Uniparental Markers in Italy Reveal a Sex-Biased Genetic Structure and Different Historical Strata. PLoS ONE, 2013, 8, e65441. | 2.5 | 61 |
| 31 | General Method to Unravel Ancient Population Structures through Surnames, Final Validation on Italian Data. Human Biology, 2012, 84, 235-270. | 0.2 | 24 |
| 32 | The Family Name as Socio-Cultural Feature and Genetic Metaphor: From Concepts to Methods. Human Biology, 2012, 84, 169-214. | 0.2 | 36 |
| 33 | Linking Italy and the Balkans. A Y-chromosome perspective from the Arbereshe of Calabria. Annals of Human Biology, 2011, 38, 59-68. | 1.0 | 19 |
| 34 | Are ethnic minorities synonymous for genetic isolates? Comparing Walser and Romance populations in the Upper Lys Valley (Western Alps). Journal of Anthropological Sciences, 2011, 89, 161-73. | 0.4 | 11 |
| 35 | Dissecting a human isolate: Novel sampling criteria for analysis of the genetic structure of the Val di Scalve (Italian Pre-Alps). Annals of Human Biology, 2010, 37, 604-609. | 1.0 | 11 |
| 36 | Italian isolates today: geographic and linguistic factors shaping human biodiversity. Journal of Anthropological Sciences, 2008, 86, 179-88. | 0.4 | 19 |

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|----|--|-----------------|--------------|
| 37 | Genetic Structure of La Cabrera, Spain, from Surnames and Migration Matrices. Human Biology, 2007, 79, 649-666. | 0.2 | 11 |
| 38 | Ethnicity and Evolution of the Biodemographic Structure of Arbëreshe and Italian Populations of the Pollino Area, southern Italy (1820–1984). American Anthropologist, 2007, 109, 735-746. | 1.4 | 15 |
| 39 | Migration matrices and surnames in populations with different isolation patterns: Val di Lima (Italian) Tj ETQq1 1 18, 676-690. | 0.784314 1.6 | rgBT /Overlo |
| 40 | Inbreeding Patterns in La Cabrera, Spain: Dispensations, Multiple Consanguinity Analysis, and Isonymy. Human Biology, 2004, 76, 191-210. | 0.2 | 19 |