

# Curriculum Vitae – Chiara Barbieri

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## Current position

Oct/2023 Senior Assistant Professor in Genetics (RTDb). Department of Environment and Life Sciences, University of Cagliari, Italy

Jul/2020 Invited affiliation: Independent Group Leader. Research group “Human genetic diversity across languages and cultures”. University of Zurich, Switzerland.

## Higher education

2010 – 2013 **PhD in Biodiversity and Evolution**, University of Bologna, Italy, and Max Planck Institute for Evolutionary Anthropology, Leipzig, Germany.  
Title of Dissertation: *Comparing genetic and linguistic diversity in African populations with a focus on the Khoisan of southern Africa*. Final score: excellent with mention.

2005 – 2008 **Master Degree in Biodiversity and Evolution**, University of Bologna, Italy. Final score: 110 out of 110 with Honors.

2002 – 2005 **Bachelor Degree in Natural Sciences**, University of Bologna, Italy.

## Academic Career

Feb/18 - Jun/20 **Senior Postdoc (Oberassistent)** at University of Zurich, Switzerland.

Jul/15 – Jan/18 **Postdoctoral position** at Max Planck Institute for the Science of Human History, Jena, Germany.

Dec/14 - Jun/15 **Postdoctoral position** at Max Planck Institute for Evolutionary Anthropology, Leipzig, Germany.

Jul/13 - Oct/14 **Postdoctoral position** at University of Bologna, Italy

Sep/09 - Apr/13 **PhD position** at Max Planck Institute for Evolutionary Anthropology, Leipzig, Germany.

## Scientific experiences and training

Jul/24	<b>Symposium organization.</b> “Human population demography and adaptation signals in the Americas”, Conference of the Society of Molecular Biology and Evolution SMBE, Puerto Vallarta, Mexico
Jun/23	<b>Scientific Committee.</b> International conference “Evolution in Action”. Centro Congressi Stefano Franscini, Monte Verità, Switzerland.
Mar/22	<b>Fieldwork in Chile.</b> Returning the results of a genetic study. Field organization and project coordination.
Aug/21	<b>Workshop organization.</b> “The South American Ethnosphere”. Centro Congressi Stefano Franscini, Monte Verità, Switzerland.
May/19	<b>Fellowship at the DFG Center Words Bones Genes Tools, University of Tübingen, Germany.</b> Collaboration for the study of genetic and linguistic diversity in Peru.
Jan/19	<b>Fieldwork in Chile.</b> Human sample collection with Mapuche communities. Field organization and project coordination.
Dec/18	<b>Symposium organization.</b> “New Methods and Discoveries in Human Archaic Admixture”. University of Zurich.
Apr/16 - May/16	<b>Visiting guest at the Institut für Rechtsmedizin - Charité - Universitätsmedizin Berlin.</b> Y chromosome analysis in the Forensic Genetic department.
Dec/15	<b>Workshop co-organization.</b> “A cross-disciplinary prehistory of Andean Civilisation”. Max Planck Institute for the Science of Human History, Jena.
Nov/15 - Dec/15	<b>Fieldwork in Peru.</b> Human sample collection in the northern coast. Field organization and project coordination.
Jan/15 - Feb/15	<b>Fieldwork in Peru.</b> Human sample collection in the Chachapoyas province. Field organization and project coordination.
Oct/13 and Dec/13	<b>Fieldwork in Italy.</b> Human sample collection among local populations.
Nov/09 – Dec/09	<b>Fieldwork in Botswana.</b> Human sample collection among native populations.
Oct/08 – Jul/09	<b>Internship at Max Planck Institute for Evolutionary Anthropology, Leipzig, Germany,</b> on project “Population relationships and language change in Burkina Faso”.
Jun/08 – Aug/08	<b>Internship on Molecular Anthropology, Parc Recerca Biomèdica de Barcelona (PRBB), Barcelona, Spain.</b> Lab analysis for the <i>Genographic Project</i> (National Geographic, IBM)
Aug/07 – Feb/08	<b>Laboratory experience</b> on Molecular Anthropology, University of Bologna, Italy
Apr/07 – Jun/07	<b>Fieldwork in Peru -</b> Human sample collection and anthropometrical measurements among native populations of the highlands and of the jungle.
Jun/06 and Jul/07	<b>Paleo-archaeological excavations</b> at Isernia-La Pineta, Italy. Collaboration with University of Ferrara, Italy.

## Talks, conference presentations, invited seminars

- October 2024 **Busan, South Korea** – IBS conference on Climate and Human Evolution. Invited keynote: “Contrasting genetic and linguistic histories and population dynamics”
- September 2024 **Caracas, Venezuela** – Venezuelan Institute for Scientific Research. Invited seminar: “Historias genéticas y culturales humanas”
- September 2024 **Rome, Italy** – Congresso Gruppo Italiano Giovani Antropologø, Invited keynote: “Human genetic and cultural histories: blending disciplines and creating narratives”
- May 2024 **Barcelona, Spain** – Invited seminar: “Matches and Mismatches between human genetic and linguistic histories”
- February 2024 **Leiden, the Netherlands** – Invited seminar: “A global analysis of matches and mismatches between human genetic and linguistic histories”
- January 2024 **Vienna, Austria** – HEAS seminar series Ancient Genomics. Invited seminar: “Contrasting linguistic and genetic histories with a global genetic dataset”
- September 2023 **Caracas, Venezuela** – Simposio de Investigación en Arqueología y Paleontología de Venezuela. Invited talk: “Migraciones y diversidad humana en Sudamérica: una perspectiva genética”
- August 2023 **Zurich, Switzerland** – Annual conference of the Latin American Center Zurich. Invited Keynote: “Genética, lenguaje y el poblamiento de América del Sur”
- August 2023 **Leipzig, Germany** – Invited seminar: “Contrasting linguistic and genetic histories with a global genetic dataset”
- April 2023 **Zurich, Switzerland** - Speech Units Workshop. Invited Talk: “Participation and transparency in the field: a conversation on best ethical practice”
- May 2023 **Budapest, Hungary** - 12th Haploid Markers Conference. Invited Keynote: “A global analysis of matches and mismatches between human genetic and linguistic histories”
- March 2023 **Tohoku University, Japan** – Yotta Informatics symposium. Invited speaker (online): “Contrasting linguistic and genetic histories of human populations with a global genetic dataset”
- September 2022 **Heidelberg, Germany** - EMBO | EMBL Symposium Reconstructing the human past: using ancient and modern genomics. Talk: “The genetic history of the southern Andes reconstructed from present day Mapuche ancestry”
- December 2021 **Barcelona, Spain** (online)– Master course of empirical linguistics. Invited lecture: “Genetic and linguistic diversity: reconstructing the past, understanding the present”
- November 2021 **Jena, Germany** (online)– Workshop: Archaeology and language. Invited speaker (with Matthias Urban and Kurt Rademaker): “The languages of the Andes”
- October 2021 **Freiburg, Germany** – Workshop: Critical Life Science. Invited speaker: “Genetic studies and multidisciplinary approaches between history, representation and identity”
- September 2021 **Zurich, Switzerland** – Workshop: The molecular anthropology of language. Invited speaker: “Matches and mismatches between human genetic history and language diversity”

- August 2021 **Zurich, Switzerland** – Workshop: The reconstruction of human history through an interdisciplinary approach. Invited speaker: "Mobility vs continuity: confronting stories and perspectives from the genetic record"
- May 2021 **Zurich, Switzerland** – Conference "Race and racism" (online). Invited speaker: "Racism and representation in genetic and bioanthropological studies".
- January 2021 **Lima, Peru** – Global Microbiome Network-First Latin American Symposium (online). Invited speaker: "Bioanthropological research and community participation in South America".
- October 2020 **Zurich, Switzerland** – NCCR-ISLE colloquium series, UZH (online). Invited speaker: "Integrating human genetics and linguistic diversity"
- May 2020 **Tübingen, Germany** – Weekly kolloquium DFG Center Words, Bones, Genes and Tools, (online). Invited speaker: "Bridging the cultural and biological histories of human populations - from local to global"
- April 2020 **Los Angeles, USA** – AAPA annual meeting (online). Poster presentation: "Genetic relationships and linguistic diversity in present-day populations of South America".
- November 2019 **Lima, Peru** – Simposio Genómica, Salud, Historia y Diversidad Humana. Invited speaker: "Migración, Genética y Lingüística en la Prehistoria de Sudamérica".
- June 2019 **Split, Croatia** – 11th ISABS Conference on Human Genome Project Based Applications in Forensic Science, Anthropology and Individualized Medicine. Poster presentation: "A catalogue of human ancestry components from population studies".
- June 2019 **Genève, Switzerland** – Conference Peopling History of Africa. Poster presentation: "A bioarchaeological study of African individuals from an early Colonial period context from Mexico City".
- April 2019 **Zurich, Switzerland** – Ringvorlesung der Kommission UZH Interdisziplinär. Invited seminar: "Genetic footprints of pre- and post-colonial contact in the Americas".
- March 2019 **Heidelberg, Germany** – EMBO-EMBL symposium Reconstructing the Human Past - Using Ancient and Modern Genomics. Talk and poster: "Gene-Language mismatches in global human history".
- March 2019 **Fribourg, Switzerland** – Invited lecture: "Genes and Languages: bridging our biological and cultural history".
- March 2019 **Zurich, Switzerland** – Inaugural workshop of the Center for the Interdisciplinary Study of Language Evolution. Talk: "Language-Gene mismatches in global human history".
- December 2018 **Jena, Germany** – Historical linguistics workshop. Invited speaker: "Bridging human genetic diversity and language change".
- August 2018 **Zurich, Switzerland** – Mini-symposium of the Latin American Center Zurich. Invited speaker: "Genetic fingerprints of human migration and contact in South America".
- July 2018 **Yokohama, Japan** – workshop "the (co-)evolution of genes, languages, and music from data analyses to theoretical models" Invited speaker: "Matches and mismatches between genes and languages: a case study from South America".

- July 2018 **Yokohama, Japan** – Meeting of the Society of Molecular Biology and Evolution. Poster presentation: “Genome-wide variation in the Andes, Amazonia and Pacific coast of western South America”.
- June 2018 **Nijmegen, Netherlands** – conference Interdisciplinary Approaches in the Language Sciences. Invited Speaker: “Language and genes: bridging our biological and cultural history”.
- November 2017 **Hinxton, UK** – conference Human Evolution: Fossils, Ancient and Modern Genomes. Poster presentation: “Reconstructing human diffusion and collapse in the Americas”.
- October 2017 **Sao Paulo, Brazil** – Invited talk: “Evidencia genética de expansión y colapso poblacionales en las Américas”.
- October 2017 **Belo Horizonte, Brazil** – symposium “Populating the diverse landscapes of South America”. Invited participant.
- September 2017 **Tübingen, Germany** – Invited talk: “Matches and mismatches between genetic and linguistic diversity”.
- June 2017 **Barcelona, Spain** – workshop Co-evolutionary patterns and transition: Analysing processes of cultural change in cross-cultural and diachronic contexts. Invited talk: “Matches and mismatches between genetic and linguistic diversity: A case-study from the Andes”.
- March 2017 **Nijmegen, The Netherlands** – Invited Lecture at Max Planck Institute for Psycholinguistics “Matches and mismatches between genetic and linguistic diversity”.
- December 2016 **Paris, France**– Workshop “DNA polymorphisms in human populations”. Talk: “Reconstructing human diffusion and collapse in the Americas”
- November 2016 **Tübingen, Germany** – Symposium “New perspectives on the peopling of the Americas”. Invited speaker: “Reconstructing demographic change and population movements through the Amazonia to Pacific corridor”
- August 2016 **Jena, Germany** – Mini African Symposium – Linguistics, Genetics and Archaeology. Invited talk: “The role of West Africa within sub-Saharan genetic diversity”
- July 2016 **Gold Coast, Australia** – Meeting of the Society of Molecular Biology and Evolution. Poster presentation: “Reconstructing human diffusion and collapse in the Americas”
- April 2016 **Berlin, Germany** – Workshop Haploid Markers. Talk: “Y chromosome sequences from southern Africa: diversity, coalescence times, and rate variation”
- January 2016 **Huddersfield, UK** – Invited Lecture: “Insights into African population prehistory - From genetic and linguistic data”
- September 2015 **London, UK** – Meeting of the European Society for the study of Human Evolution (ESHE). Talk: “Paleolithic signatures from European glacial refugia revealed by mtDNA genomes”
- May 2015 **Leipzig, Germany** – Workshop “Speaking (of) Khoisan: a symposium reviewing southern African prehistory”. Invited speaker.
- February 2015 **Leipzig, Germany** – The Human Mutation Rate Meeting. Poster presentation: “Estimating the TMRCA of the Y chromosome phylogeny with new sequence data from southern Africa”

June 2014	<b>Halle, Germany</b> – Workshop "The Claims of Descent: Science, Representation, Race and Redress in 21st Century South Africa". <u>Invited speaker</u> . Talk "The use of the "ethnolinguistic" unit in genetic and linguistic research"
April 2014	<b>Leicester, UK</b> – EMBO congress "Human evolution in the genomic era: Origins, populations and phenotypes". Talk and poster: "Language contact and genetic variation in South America: the case of the Peruvian Yanasha"
October 2013	<b>Boston, USA</b> – Meeting of the American Society of Human Genetics. Poster presentation: "Y chromosomal sequences from southern Africa allow direct comparison of paternal and maternal prehistories"
August 2013	<b>Leipzig, Germany</b> – Workshop on Language diversity and prehistory. <u>Invited speaker</u> . Talk: "Prehistorical relationships in the Americas inferred through population genetics"
September 2012	<b>Firenze, Italy</b> – Incontro Nazionale Giovani Antropologi. <u>Invited speaker</u> . Talk: "Variabilità genetica e linguistica a confronto in popolazioni Khoisan africane"
June 2012	<b>Dublin, Ireland</b> – Meeting of the Society of Molecular Biology and Evolution. Poster presentation: "mtDNA genomes of Khoisan populations deciphered under a linguistic and social anthropological perspective"
November 2011	<b>Leipzig, Germany</b> – Symposium on Population Prehistory of the Andes: A Cross-Disciplinary Conspectus. <u>Invited presentation</u> .
October 2011	<b>Berlin, Germany</b> – Workshop on Methodology in Linguistic Prehistory. <u>Invited speaker</u> . Barbieri C. & Pakendorf B. "Integrating molecular data into the linguistic perspective on Khoisan prehistory"
July 2010	<b>Lyon, France</b> – SBE meeting. Poster presentation: "Complete mtDNA genome and Y chromosome data from West Africa: demographic history and population contact in Burkina Faso"
June 2009	<b>Split, Croatia</b> - 6th ISABS Conference on Human Genome Project Based Applications in Forensic Science, Anthropology and Individualized Medicine. Talk and poster: "mtDNA variability in native Andean populations: the case of Lake Titicaca"

## Grants, awards, fellowships

October 2024	Grant "innovation grant" from the NCCR Evolving Language for the project "Upscaling gene-language comparisons through historical paths and present-day diversity" (60,000 CHF)
November 2022	Grant "new research format" to produce a documentary, from the Latin American Zentrum Zurich, Switzerland (2,000 CHF)
August 2021	Fellowship for workshop organization from the NCCR Evolving Language (3,000 CHF) and the Latin American Zentrum Zurich, Switzerland (3,000 CHF)
2017 - 2019	Post-Ph.D. Research Grant from the Wenner-Gren Foundation (20,000 \$)
December 2018	Fellowship for symposium organization from the University Research Priority Program "Evolution in Action" of the University of Zurich, Switzerland (3,200 CHF)
February 2015	Grant for fieldwork in Peru from the Department of Linguistics, Max Planck Institute for Evolutionary Anthropology, Leipzig (1500 €)

September 2013	Award for Best PhD thesis from the Italian Association of Genetics
June 2011	Scholarship for tuition fees of Summer Institute in Statistical Genetics from University of Washington, Seattle, USA. (1200 \$)
June 2009	Split, Croatia - 6th ISABS Conference on Human Genome Project Based Applications in Forensic Science, Anthropology and Individualized Medicine. Recipient of Young Investigator Award (500 €)
April 2007	Grant from University of Bologna for fieldwork in Peru (1000 €)

## Teaching experience

### University of Cagliari

#### Master courses

2024: Semester module “Evolutionary Genomics” 7 CFU.

### University of Zurich

#### Undergraduate courses

2018 - 2023: Block course BIO373, Next Generation Sequencing for Evolutionary and Ecological Genomics.

#### Master courses

2019 - 2023: Semester module “Introduction to Human Genetics and Bioanthropology for linguists”.

#### PhD courses

2019, 2020, 2022: BIO624 - Human Genetic, Demographic and Cultural Diversity.

### University of Leipzig

2010 - 2013: Lectures for the graduate course of Molecular Anthropology.

### University of Bologna

2008 - 2017: Teaching Assistance for undergraduate and graduate courses: Human Genetics, Physical Anthropology, Human Anatomy and Osteology, Human Ecology, Statistical Genetics for Molecular Anthropology. Lectures, laboratory sessions, computer sessions and grading.

### Teaching for graduate programs

2019: Bioinformatic analysis in human genetic diversity. Invited tutor for the workshop on human diversity at University of San Martín de Porres, Lima, Peru. 2 days lectures and exercises.

2016-2017: Tutor for the Spring School of Quantitative Methods/Masterclass, MPI SHH Jena. Introduction to Genetics and phylogenetic reconstruction. 2 days lectures and exercises.

## Professional Memberships

Society for Molecular Biology and Evolution, European Society for the study of Human Evolution, Associazione Antropologica Italiana, Associazione Genetisti Italiani, Società Italiana di Biologia Evoluzionistica, Registro degli Antropologi Forensi Italiani, Lateinamerika-Zentrum Zürich

## Outreach activities

Producer of the documentary “Finding First Peoples: the Mapuche” (CH/IT, 2023, 65 min, director: Renato Giugliano), financed with resources obtained from various institutions in Zurich. The audiovisual project shows how scientists can interact with participants of a study, promote ethical practice in genomic science, and give voice to indigenous minorities of Southern Chile.

Realization of a short movie about Language for the Zurich SciFilm Hackathon.

Curation of press releases for scientific articles. Interviewed for The Atlantic, Smithsonian Magazine, UZH magazine, Pikaia (IT), Betapress (IT), OggiScienza (IT), Radio3 RAI scienza (IT), SRF – Swiss Radio Television. Participation to the documentary “Peru’s Mysterious Cloud Warriors” for Curiosity Stream channel.

Schools: Participation to the program “Skype a Scientist” for scientific outreach in high schools. Zoom seminars to present the job of scientific research and the importance of women in science (in Italy).

General public: invited speaker for events for the 8<sup>th</sup> of March to promote the role of women in science/STEM (in Italy).

## Other academic activity and qualifications

Reviewer for Molecular Biology and Evolution, American Journal of Human Genetics, BMC Genomics, PNAS, Cell, Plos One, American Journal of Physical Anthropology, European Journal of Human Genetics, American Journal of Human Biology, AIMS Genetics, Quaternary International, Human Heredity, Journal of Language Evolution, Forensic Science International: Genetics, Scientific Reports, Nature Communication, Current Anthropology, Annals of Human Genetics.

Ad hoc reviewer for PhD thesis, master thesis, national and international grants. Member of PhD review committees. Member of PhD program of Life Sciences at the University of Zurich and *collegio dei docenti* University of Cagliari. Main supervisor for 3 master students, 2 bachelor thesis, 2 PhD students at University of Zurich.

Associate editor for the Journal of Language Evolution. Academic editor for PeerJ.

Member of the scientific committee of the conference EvoLang – Evolution of Language. Member of the steering committee for the URPP Evolution in Action at the University of Zurich. Senior researcher of the NCCR Evolving Language – National Science Foundation, Switzerland.

Italian habilitation as associated professor in Anthropology and Human Genetics. Abilitazione Scientifica Nazionale: 05/B1, valid until 05/04/28; 05/I1, valid until 26/07/28

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## Publications

Graff, A., Chousou-Polydouri, N., Inman, D., Skirgård, H., Lischka, M., Zakharko, T., **Barbieri, C.**, & Bickel, B. 2025. Curating global datasets of structural linguistic features for independence. *Scientific Data*, 12(1), 106. <https://doi.org/10.1038/s41597-024-04319-4>

Resutik, P., Schneider, J., Aeschbacher, S., Vigeland, M. D., Gysi, M., Moser, C., **Barbieri, C.**, Widmer, P., Currat, M., Kratzer, A., Krützen, M., Haas, C., & Arora, N. 2025. Uncovering genetic signatures of the Walser migration in the Alps: Patterns of diversity and differentiation. *Forensic Science International: Genetics*, 76, 103206. <https://doi.org/10.1016/j.fsigen.2024.103206>

Morris, A. G., Salie, T., Mittnik, A., Rebello, G., **Barbieri, C.**, Parkington, J., Krause, J., & Ramesar, R. 2025. Reconstructing ancient Southern African mitochondrial genomes at Faraoskop. *Archaeological and Anthropological Sciences*, 17(1), 11. <https://doi.org/10.1007/s12520-024-02130-1>



- Passmore S, Wood ALC, **Barbieri C**, Shilton D, Daikoku H, Atkinson QD & Savage PE. 2024. Global musical diversity is largely independent of linguistic and genetic histories. *Nat Commun* 15, 3964. <https://doi.org/10.1038/s41467-024-48113-7>
- Arango-Isaza E, Aninao MJ, Campbell R, Martínez FI, Shimizu KK, **Barbieri C**. 2023. Bridging the Gap: Returning Genetic Results to Indigenous Communities in Latin America. *Frontiers in Genetics* 14. <https://www.frontiersin.org/articles/10.3389/fgene.2023.1304974>.
- Menéndez LP, **Barbieri C**, López Cruz IG, Schmelzle T, Breidenstein A, Barquera R, Borzi G, Schuenemann VJ, Sánchez-Villagra M. 2023. On Roth's "human fossil" from Baradero, Buenos Aires Province, Argentina: morphological and genetic analysis. *Swiss J Palaeontol* 142, 26. <https://doi.org/10.1186/s13358-023-00293-3>
- Arango-Isaza E, Capodiferro MR, Aninao MJ, Babiker H, Aeschbacher S, Achilli A, Posth C, Campbell R, Martínez FI, Heggarty P, Sadowsky S, Shimizu KK, **Barbieri C**. 2023. The genetic history of the southern Andes from present-day Mapuche ancestry. *Current Biology* 33, 1-14. <https://doi.org/10.1016/j.cub.2023.05.013> [LINK](#)
- Capodiferro MR, Chero Osorio AM, Rambaldi Migliore N, Tineo Tineo DH, Raveane A, Xavier C, Bodner M, Simão F, Ongaro L, Montinaro F, Lindo J, Huerta-Sanchez E, Politis G, **Barbieri C**, Parson W, Gusmão L, Achilli A. 2023. The multifaceted genomic history of Ashaninka from Amazonian Peru. *Current Biology*. 33(8):1573-1581.e5. doi: 10.1016/j.cub.2023.02.046. [LINK](#)
- Barbieri, C.**, Blasi, D. E., Arango-Isaza, E., Sotiropoulos, A. G., Hammarström, H., Wichmann, S., Greenhill, S. J., Gray, R. D., Forkel, R., Bickel, B., & Shimizu, K. K. 2022. A global analysis of matches and mismatches between human genetic and linguistic histories. *Proceedings of the National Academy of Sciences*, 119(47), e2122084119 <https://www.pnas.org/doi/10.1073/pnas.2122084119>
- Sotiropoulos, A.G., Arango-Isaza, E., Ban, T., **Barbieri, C.**, Bourras, S., Cowger, C., Czembor, P.C., Ben-David, R., Dinor, A., Ellwood, S.R., Graf, J., Hatta, K., Helguera, M., Sánchez-Martín, J., McDonald, B.A., Morgounov, A.I., Müller, M.C., Shamanin, V., Shimizu, K.K., Yoshihira, T., Zbinden, H., Keller, B., Wicker, T., 2022. Global genomic analyses of wheat powdery mildew reveal association of pathogen spread with historical human migration and trade. *Nat. Commun.* 13, 4315. <https://doi.org/10.1038/s41467-022-31975-0>
- Lankheet I, Vicente M, **Barbieri C**, Schlebusch C. 2021. The performance of common SNP arrays in assigning African mitochondrial haplogroups. *BMC Genomic Data* 221 22, 1–8
- Aguirre-Fernández G\*, **Barbieri C\***, Graff A, Pérez de Arce J, Moreno H, Sánchez-Villagra MR. (\*first authors contributed equally). 2021. "Cultural macroevolution of musical instruments in South America". *Humanit Soc Sci Commun* 8, 208. <https://doi.org/10.1057/s41599-021-00881-z>
- Schlebusch CM, Loog L, Groucutt HS, King T, Rutherford A, **Barbieri C**, Barbujani G, Chikhi L, Stringer C, Jakobsson M, Eriksson A, Manica A, Tishkoff SA, Scerri EM, Scally A, Brierley C, & Thomas MG. 2021. Human origins in Southern African palaeo-wetlands? Strong claims from weak evidence. *Journal of Archaeological Science*, 130, 105374. <https://doi.org/10.1016/j.jas.2021.105374>
- Urban M & **Barbieri C**. 2020. North and South in the ancient Central Andes: contextualizing the archaeological record with evidence from linguistics and molecular anthropology. *Journal of Anthropological Archaeology*, 60, 101233
- Nakatsuka N, Lazaridis I, **Barbieri C**, Skoglund P, Rohland N, Mallick S, Harkins-Kinkaid K, Ferry M, Harney E, Michel M, et al. 2020. A Paleogenomic Reconstruction of the Deep Population History of the Andes. *Cell* 181:1131-1145.e21.
- Barquera R, Lamnidis TC, Lankapalli AK, Kocher A, Hernández-Zaragoza DI, Nelson EA, Zamora-Herrera AC, Ramallo P, Bernal-Felipe N, Immel A, Bos K, Acuña-Alonzo V, **Barbieri C**, Roberts P, Herbig A, Kühnert D, Márquez-Morffín L, Krause J. 2020. Origin and Health Status of First-Generation Africans from Early Colonial Mexico. *Curr. Biol.* 30(11), 2078-2091.e11
- Barbieri C**, Barquera R, Arias L, Sandoval JR, Acosta O, Zurita C, Aguilar-Campos A, Tito-Álvarez AM, Serrano-Osuna R, Gray R, Mafessoni F, Heggarty P, Shimizu KK, Fujita R, Stoneking M, Pugach I, Fehren-Schmitz L. 2019. The current genomic landscape of western South America: Andes, Amazonia and Pacific Coast. *Mol Biol Evol.* 36(12), 2698–2713.

Bajić V, **Barbieri C**, Hübner A, Güldemann T, Naumann C, Gerlach L, Berthold F, Nakagawa H, Mpoloka SW, Roewer L, Purps J, Stoneking M, Pakendorf B. 2018. Genetic structure and sex-biased gene flow in the history of southern African populations. *American Journal of Physical Anthropology*. 167(3), 656-671.

Tucci S, Vohr SH, McCoy RC, Vernet B, Robinson MR, **Barbieri C**, Nelson BJ, Fu W, Purnomo GA, Sudoyo H, Eichler EE, Barbujani G, Visscher PM, Akey JM, Green RE. 2018. Evolutionary history and adaptation of a human pygmy population of Flores Island, Indonesia. *Science* 361:511–516.

Arias L, **Barbieri C**, Barreto G, Stoneking M, Pakendorf B. High-resolution mitochondrial DNA analysis sheds light on human diversity, cultural interactions, and population mobility in Northwestern Amazonia. 2018. *American Journal of Physical Anthropology*; 165(2): 238-255.

**Barbieri C**, Sandoval JR, Valqui J, Shimelman A, Ziemendorff S, Schröder R, Geppert M, Roewer L, Gray R, Stoneking M, Fujita R and Heggarty P. 2017. Enclaves of genetic diversity resisted Inca impacts on population history. *Scientific Reports*, 7: 17411.

Bortolini E, Pagani L, Crema ER, Sarno S, **Barbieri C**, Boattini A, Sazzini M, Da Silva SG, Martini G, Metspalu M, Pettener D. 2017. Inferring patterns of folktale diffusion using genomic data. *Proceedings of the National Academy of Sciences*, p.201614395.

Sarno S, Boattini A, Pagani L, Sazzini M, De Fanti S, Quagliariello A, Gnecci-Ruscione GA, Guichard E, Ciani G, Bortolini E, **Barbieri C**, Cilli E, Petrilli R, Mikerezi I, Sineo L, Vilar M, Wells S, Luiselli D, Pettener D. 2017. Ancient and recent admixture layers in Sicily and Southern Italy trace multiple migration routes along the Mediterranean. *Scientific Reports*, 7(1), p.1984.

Boattini A, Sarno S, Bini C, Pesci V, **Barbieri C**, De Fanti S, Quagliariello A, Pagani L, Ayub Q, Ferri G, Pettener D, Luiselli D, Pelotti S. 2016. Mutation rates and discriminating power for 13 Rapidly-Mutating Y-STRs between related and unrelated individuals. *Plos One*. 11 (11) pp: e0165678

Malaspinas AS, Westaway MC, Muller C, Sousa VC, Lao O, Alves I, Bergström A, Athanasiadis G, Cheng JY, Crawford JE, Heupink TH, Macholdt E, Peischl S, Rasmussen S, Schiffels S, Subramanian S, Wright JL, Albrechtsen A, **Barbieri C**, et al. 2016. A genomic history of Aboriginal Australia. *Nature*. 538: 207–214.

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## Invited contributions

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## Edited volumes

A. Ravignani, **C. Barbieri**, M. Flaherty, Y. Jadoul, E. Lattenkamp, H. Little, M. Martins, K. Mudd, & T. Verhoef (Eds.), *The Evolution of Language: Proceedings of the 13th International Conference (Evolang13)*. 2020. Nijmegen: The Evolution of Language Conferences.

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Guest editor for the special issue "The Genomic Impact of Human Migrations" of the journal *Genes*.

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