



Prof. Ludovic Orlando, HDR, PhD, Agrégé, Normalien

Director, Centre for Anthropobiology & Genomics of Toulouse, CAGT

Université Paul Sabatier Toulouse III, France

Born in Marseille (France), January 24th 1977 (46 years old)

Married, three children

Email: Ludovic.orlando@univ-tlse3.fr, orlando.ludovic@gmail.com

Ph: +33 6 18 27 95 10; Skype: lorlando1369, Twitter: LudovicLorlando

<https://cagt.cnrs.fr/orlando-ludovic/>

Researcher ID A-8932-2013 / Orcid ID: 0000-0003-3936-1850

Scientific Areas

Ancient DNA, Animal Domestication, Molecular Anthropology, Evolutionary Biology, Statistical Genomics

Scientific Achievements

- Led the international teams that:
 - Bridged western and indigenous science by revealing the true origins of First Nations horses [Science 2023]
 - Identified the horse [Nature 2021] and donkey [Science 2022] domestication homelands
 - reconstructed the genomic history of horse domestication [Science 2017, Cell 2019]
 - discovered that horses were domesticated twice [Science 2018]
 - unraveled the genetic basis of cold adaptations in Yakutian horses [PNAS 2015]
 - carried out the first conservation genomic study of feral horses [Curr Biol 2015]
 - tracked equine domestication and speciation [PNAS 2014ab]
 - described the first ancient epigenome [Genome Research 2014, Science 2014]
 - sequenced the first Middle Pleistocene genome [Nature 2013]
 - tracked the genetic consequences of the last great Megafauna extinction [Nature 2011]
- Contributed to the international teams that:
 - assessed the environmental consequences of global climate change in the Arctic [Nature 2014, 2016, 2020]
 - sequenced 500+ ancient human genomes [Nature 2014ab, 2015, 2018ab; Science 2014, 2017, 2018ab]
 - sequenced the oldest proteome [Nature 2019]
 - demonstrated the existence of plague epidemics as early as the Bronze Age [Cell 2015]
 - resolved the bird tree of life [Science 2014ab]
 - sequenced the first ancient human genome [Nature 2010]
 - discovered the genetic presence of Neanderthals in Asia [Nature 2007]
- Developed user-friendly, open-source and community-oriented software for analyzing next-generation sequencing data [Bioinformatics 2015, 2016, 2017, 2022ab; Nat Protoc 2014; Mol Ecol 2016; MBE 2016, 2018, Genetics 2019, Gigasciences 2019; Frontiers Ecol 2020]

Track Record of Early Achievements

As of Sept 18th 2023, I have co-authored 208 articles in international peer-reviewed journals (75, ~36 %, of which during the last five years). These appeared in *Nature* (23), *Science* (15), *Cell* (4), *PNAS* (12), as well as top-specialist journals in the field of genomics (*Genome Res*), evolutionary biology (*Mol Biol Evol*, *Mol Ecol*), computational biology (*Bioinformatics*) and archaeological science (*J Arc Sci*). I have also published 9 Book Chapters, 2 books and multiple reviews in high-impact journals, including in *Nat Rev* (2015, 2020, 2021) and the *Annual Reviews* series (2014, 2017, 2020, 2021). I have co-edited a full issue dedicated to ancient DNA for *J Hum Evol* (2015) and been invited to write opinion articles for *Nature* (2023), *Cell* (2020), *PNAS* (2016, 2018, 2023), *Curr Biol* (2015), *Genome Biology* (2015) and *Bioessays* (2014, 2020). Collectively, my papers have attracted 38,172 citations (Google Scholar), 18,666 (~49%) of which in the last three years, and my H-index is 95. I have co-organized 11 international conferences, chaired/co-organized 7 sessions (including at Cold Spring Harbor *Biology of Genomes* and *UISPP*) and been the opponent for 23 international PhD committees in 7 countries, including at high-rank universities such as Oxford. I have lectured 1,200+ hours at all university levels and tutored 11 international workshops. I have won 1 ERC CoG and 1 ERC SyG grant. I am an Academic Editor for *Scientific Reports* and *PeerJ*, and Editor-in-Chief of *Science and Technology for Archaeological Research*. During the last five years, my projects have collectively been granted ~16 M€, from diverse funding bodies including *EU*, *ANR*, *FNU*, *Simone & Cino Del Duca* and *Vellux Fonden*.

Education

Dec 14th 2009

Professorial thesis (HDR), ENS Lyon, France

July 7th 2003

PhD with highest honors. University of Lyon, France

July 2000

MSc in Development, Genetics and Immunology, University of Lyon, France (ranking: 1st)

July 1999

Aggrégation of Life and Earth Sciences (ranking: 4th / >2,000 applicants)

Sept 1996-July 1998

Magistère of Cellular and Molecular Biology, ENS Lyon, France

Sept 15th 1996

Admitted after national competitive examination of >2,000 applicants as **Normalien** at ENS Lyon

Academic Positions

Summary	I established my research group at the Centre for GeoGenetics (CGG), Denmark, in April 2010, first as an Associate Prof, then as a Prof of Molecular Archaeology from March 2016 and until June 2020. While my main appointment remained at CGG until September 2016, I have been appointed as an Invited Professor at the University Paul Sabatier, Toulouse France, since January 2015. From October 2016, I have relocated my research group there, following my appointment as a CNRS Research Director. I have acted as the CNRS 5288 Deputy Director since October 2016, and I have acted as the CAGT founding director from Jan 2020.
Jan 2021-now	Director , CAGT Laboratory, Univ. Paul Sabatier, Toulouse, France
Jan 2020-Dec 2020	Director , AMIS Laboratory, Univ. Paul Sabatier, Toulouse, France
Sept 2017-now	Group Leader , Archaeology, Genomes, Evolution & Societies (AGES)
Oct 2016-Jun 2020	Deputy Director , CNRS AMIS 5288, Univ. Paul Sabatier, Toulouse, France Research Director , CNRS, Univ. Paul Sabatier, Toulouse, France Professor of Molecular Archaeology, CGG, Copenhagen, Denmark (20% full-time equivalent) Invited Professor , Chaires d'Excellence IDEX, Univ. Toulouse (20% full-time equivalent) Professor of Molecular Archaeology, CGG, Copenhagen, Denmark Curator of the National Cryobank, Natural History Museum of Denmark
Jan 2015-Sept 2016	Group Leader , CGG, Copenhagen, Denmark
Mar 2016-Sept 2016	Associate Professor of Molecular Archaeology, CGG, Copenhagen, Denmark
Apr 2010-Sept 2016	Curator of the Paleontology collection, Natural History Museum of Denmark
Apr 2010-Feb 2016	Curator of the Quaternary Zoology collections, Natural History Museum of Denmark
Jul 2014-Jul 2015	Assistant Professor , ENS Lyon, Institute for Functional Genomics of Lyon (IGFL), France
Jul 2013-Jul 2015	Post-Doctoral Researcher , University Aix-Marseille 1, Phylogenomics lab, France
Sep 2005-Mar 2010	
Sep 2004-Aug 2005	

Prizes, Awards, Miscellaneous

Nov 2023	CNRS Silver Medal
Jan 2018, 2020	Highly Cited Researcher List 2018, 2020
Dec 2015	Article celebrated as AIA's top 10 discoveries of 2015
Jan 2014	Article ranked 8th on the 'Discover Magazine's Top 100 science stories'; celebrated in 'Genomes of the year' in Science (2013) and in '365 days in review for 2013' (Nature)
Nov 2013	videnskab.dk top-10 research project in Denmark for year 2013
2006-2007	Marie-Curie IEF (FP6-010631, Ancient Biomolecules Centre, Oxford) [Declined]
2004	Annual Academic Award Le Monde de la Recherche Scientifique

Professional Service

Dec 2017	HCERES Evaluation committee, ISYEB, Committee President
Apr 2017 – now	Editorial Board , Associate Editor, <i>PeerJ</i>
Sep 2016 – now	Editorial Board , Associate Editor, <i>Scientific Reports</i>
2015, 2016	Expert Member of the Institut Universitaire de France, Young Scientist panel
2014 - now	Editorial Board , Editor-in-Chief, <i>Science and Technology of Archaeological Research</i>
2012 - 2020	Editorial Board , Academic Editor, <i>PLoS ONE</i>
2012-2013	Expert Member of the SVSE7 Evaluation Committee of the French National Agency for Research
2009-2011	Expert Member of the French RTP CNRS "Paleogenetics of ancient humans and their environment"
2007-2009	Scientific Council of the Institute for Functional Genomics, ENS Lyon (CNRS UMR 5242)
Sept 2005-Mar 2010	960h of lectures for BSc, MSc and PhD students in French Universities

Research Supervision and Research Funding

- **Group Leader** of a research group currently consisting of 22 staff members, including 1 CNRS Research Director, 1 Prof, 1 Associate Prof, 3 CNRS Researchers, 2 Research Engineers (CNRS, INRAE), 3 Assistant Engineer, 1 Lab Technician, 7 Post-Doctoral researchers, and 3 PhD students
- **Main Supervisor** of 20 post-doctoral researchers (15 in the last five years), 11 PhD students (8 in the last five years), 19 MSc students (11 in the last five years)
- **Attracted** 20 international post-doctoral researchers since 2011, representing ~2.46 M€ and including: 1 Postdoctoral Exchange Fellowship Program of the China Postdoctoral Council 38,500€; 3 Marie-Curie FP7-PEOPLE-IEF fellowships (2012-2014: 228,082.6 €, 218,560.2 €, 2014-2016 221,154.6€), 8 Marie-Curie MSCA-IF fellowships (2015-2017: 200,194.8 €, 2017-2019: 200,194.8 €, 2018-2020: 173,076.0€, 2020-2023: 186,876.96 €; 2021-2023: 196,707.84 €; 2021-2023: 184,707.84 €; 2022-2024: 184,707.84 €; 2022-2024: 195,914.88 €), 1 VillumFonden Blokstipendier (2015-2016: 1,090,916 DKK), 1 AXA Research Fund fellowship (2012-2014: 120,000€); 1 EMBO long-term fellowship (2016-2017: 791,076.6 DKK+6,018.62 €) and 1 Fondation Simone & Cino

Del Duca (**50,000€**).

- **Main Investigator** for 23 grants representing **~16.26 M€** (2022-2028: ERC SyG Horse Power, **10,478,093 €**; 2021-2025: CNRS IRP AnimalFarm, **50,000 €**; 2022-2023: MARENGO, France Génomique Appel à Grand Projet, 30 sequencing S4 lanes for the NovaSeq 6000 instrument; 2020-2021 : Fondation Simone et Cino Del Duca, **125,000€**; 2020-2021: CNRS MITI **40,000 €**; 2019-2020 : ANR MRSEI AnimalFarm, **29,970€**; 2017-2020 : GENCI A0050310593, 1M CPU hours on HPC servers; ANR LifeChange, **492,032 €**; 2016-2020: ERC-*Consolidator* 2015, **1,999,555 €**; 2016-2019 : *Villum Fonden Research Project*, **5,995,928 DKK** ; 2017-2018: BUCEPHALE, France Génomique Appel à Grand Projet, 32 sequencing flow-cells for the HiSeq 4000 instrument; 2015-2017: Chaire d'attractivité, IDEX, Univ. de Toulouse, France **750,000 €**; 2015, EU Synthesis-3 Programme, **2,000 €**; 2014-2017 : *DFF Danish Research Foundation, Major Research Project*, **5,218,917 DKK**; 2014: *DASI International Network Programme*, **222,507 DKK**; 2014-2016: *Villum Fondens Blokstipendier*, **1,090,916 DKK**; 2014-2015: *International Research Group Program* (IRG14-08), *Deanship of Scientific Research*, King Saud Univ, **75,000 €**; 2012-2015: *Marie-Curie FP7-PEOPLE-CIG*, **100,000 €**; 2012-2015: *DFF FNU Minor Research Project*, **1,958,400 DKK**; 2012: *GEUS/Geogenetics/Geocenter*, **975,000 DKK**; 2008-2009: *French-Australian Science and Technology linkage*, **12,000 €**; 2011, 2012: *French-Danish Co-operation from the French Embassy*, **45,000 DKK**)
- **Co-Investigator for 13 Grants** since 2010 (including 3 ANR, 1 ANRS & 2 Marie Curie ITN, *EUROAST & ArchSci2020*)

Visiting Scientists

Jan-Feb 2007 **MPI for Evolutionary Anthropology**, Leipzig, Germany (Director: Pr. Svante Paabo)

July-Aug 2008 **Australian Centre for Ancient DNA**, Adelaide, Australia, (Director: Pr. Alan Cooper)

Research Production, Summary

- H-index: **95** [Google Scholar]
- Citations: **38,172** [Google Scholar]
- **208 publications** in peer-reviewed journals (23 *Nature*, 15 *Science*, 4 *Cell*, 12 *PNAS*, 10 *Curr Biol*, 3 *Genome Res*)
- **2 Books, 9 Book chapters**
- Referee for **250+ Articles** (~10-15 per year), including *Nature*, *Science*, *Cell*, *PNAS* and *Curr Biol*
- **177 Invited Seminars and Conferences across 28 countries**
- Invited lecturer in **11 international workshops** (Paleogenomics summer school, Cargese, October 2011; Gulbenkian Training Program in Bioinformatics, Lisbon, June 2012; Adelaide Bioinformatics Workshop, Adelaide, November 2012, 2013; Computing for NGS analyses, University of Toulouse, December 2013; Mathematical and Computational Evolutionary Biology, Porquerolles, June 2015; Workshop on Population and Speciation Genomics, Český Krumlov, Czech Republic, February 2016, 2018, 2020; Max Planck Institute for the Science of Human History, Jena, March 2016; Spring School in Bioinformatics and Population Genomics. Leukerbad, Switzerland, May 2016; Italian Society for Evolutionary Biology, Ferrara, Italy, Dec 2016)
- Organization of **10 international conferences** (SBME 2010; ISBA 2010; INQUA 2011; SMBE 2012; SPAAM 2016; UISPP 2018; ISBA 2018; AnimalFarm 2019; AIEM 2019; ISBA 2020)
- Participation to **23 international PhD** (AUS 2; DK 4; FR 10; SP 2; CH 1; IE: 1; SW 1; PT 1; UK 1) and **5 HDR** (French Professorial Thesis, 2016, 2017, 2018, 2020, 2022) committees

Selection of 10 Peer-Reviewed Publications

- 1) Taylor WT, [87 co-authors], **Orlando L.** 2023. Early dispersal of domestic horses into the Great Plains and Northern Rockies. **Science**, in press.
- 2) Todd ET, [47 co-authors], **Orlando L.** 2022. The genomic history and global expansion of domestic donkeys. **Science** 377:1172-80.
- 3) Librado P, [161 co-authors], **Orlando L.** 2021. The origins and spread of domestic horses from the Western Eurasian steppes. **Nature** 598:634-640.
- 4) Frantz L, Bradley D, Larson G, **Orlando L.** 2020. Animal domestication in the era of ancient genomics. **Nat Rev Genet** 21:449-60.
- 5) Fages A, [119 co-authors], **Orlando L.** 2019. Tracking Five Millennia of Horse Management with Extensive Ancient Genome Time Series. **Cell** 2019 Apr 26. pii: S0092-8674(19)30384-8.
- 6) Gaunitz C, [45 co-authors], **Orlando L.** 2018. Ancient genomes revisit the ancestry of domestic and Przewalski's horses. **Science** 360:111-114
- 7) Librado P, [31 co-authors], **Orlando L.** 2017. Ancient genomic changes associated with domestication of the horse. **Science** 356:442-5.
- 8) Der Sarkissian C, [24 co-authors], **Orlando L.** 2015. Evolutionary genomics and conservation of the endangered Przewalski's horse. **Curr Biol** 25:2577-83.
- 9) Pedersen JS, [19 co-authors], **Orlando L.** 2014. Genome-wide nucleosome map and cytosine methylation levels of an ancient human genome. **Genome Res** 24:454-66.
- 10) **Orlando L.**, [54 co-authors], Willerslev E. 2013. Recalibrating *Equus* evolution using the genome sequence of an early

Selection of 10 Invited Seminars

12/07/2023	Hilgendorf Lecture. Tuebingen, Germany.
13/12/2022	Harrison Lecture. The genomic history of the domestic horse. Cornell University, Ithaca, USA
25/07/2022	Plenary. In: Havemeyer 13 th International Horse Genome Workshop. Ithaca, USA
11/01/2022	Plenary. In: Plant and Animal Genomics XXIX. San Diego, USA
20/02/2020	Keynote. In: Inauguration day of the Centre for Palaeogenetics, Swedish Museum of Natural history, Stockholm
17/06/2019	Keynote. International Society for Applied Biological Sciences. Split, Croatia
30/03/2019	Keynote. Reconstructing the Human Past - Using Ancient and Modern Genomics. EMBL Heidelberg, Germany.
04/11/2018	Keynote. In: Australian Genomic Technologies Association (AGTA) Annual Meeting, Adelaide, Australia
02/05/2018	Keynote. In: Ancient DNA from the sub-seafloor. Exploratory Seminar, Harvard University, USA
18/04/2018	Keynote. In: 1 st AsiaEvo Conference, Shenzhen, China

Communication & Outreach [Selection]

Radio & TV/Videos

01/03/2023	New York Times, Carl Zimmer, Ancient DNA reveals history of hunter-gatherers in Europe.
01/05/2022	Marilyn: her final secret. https://www.sbs.com.au/ondemand/video/2045259843619/marilyn-her-final-secret
01/03/2022	Le Monde. La science sur les traces de la peste de 1720 à Marseille
13/01/2021	Arte 28 minutes, Elisabeth Kein. L'ADN remonte le temps
23/02/2021	France 5, C à dire, Mélanie Taravant. ADN : un voyage de plus de 700,000 ans
25/10/2021	France Bleu, Sidonie Bonnec, Minute Papillon. L'ADN fossile, comment peut-il redonner vie aux dinosaures
22/10/2021	CBC Radio, Bob McDonald, Quirks & Quarks. Whoa Nelly! Modern horses come from ancestors we improved 4,000 years ago
21/10/2021	France Info. Boris Hallier, Le cheval a été domestiqué pour la première fois en Russie, 22 siècles avant notre ère
18/02/2022	France Culture, La Série Documentaire. Perrine Kervin, La Grande peste, l'empreinte d'une tueuse, épisode 8/8 : l'ADN du mal
01/02/2021	RFI, Caroline Lachowsky. Comment remonter le temps grâce à l'ADN fossile ? Et jusqu'où ?
12/01/2021	France Inter, La Terre Au Carré. Mathieu Vidard. Voyage dans le passé grâce à la paléogénétique.
23/01/2021	France Culture, Carbone 14, Vincent Charpentier. « Inné et acquis sont dans un bateau », quand la paléogénétique raconte notre histoire
19/05/2019	5000 ans d'histoire entre l'Homme et les chevaux. Les Années Lumières, Radio Canada
17/05/2019	La domestication du cheval a modifié son ADN. Journal des Sciences, France Culture
14/05/2019	The first horse warriors. NOVA, documentary by Dr Niobe Thompson
09/05/2019	5000 ans d'histoire du cheval. La tête au carré, France Inter
22/02/2018	Why the last wild horses really aren't. Nell Greenfieldboyce. National Public Radio
30/05/2017	La révolution paléo-génomique. France Culture. Carbone 14
28/04/2017	France Inter. La tête au carré

Press

05/04/2022	Knowable Magazine. Amber Dance, The tale of the domesticated horse
14/01/2022	The Scientist. Chris Baranuk, Ancient Mesopotamians bred horse-like hybrids
26/01/2022	Horizon: The EU Research & Innovation Magazine. Gareth Willmer, Quest to uncover the origins of horse taming is rewriting our picture of the past
18/11/2021	Pour la Science. François Savatier, La steppe pontique, berceau du cheval domestique
31/10/2021	Le Point. Génétique : la course folle du cheval domestique
27/10/2021	Cheval Magazine. Bettina Hubert, Les origines du cheval domestique dévoilées
26/10/2021	Le Monde. Hervé Morin, Débats éthiques sur l'étude de l'ADN humain ancien
22/10/2021	Nature. Tosin Thompson, Ancient DNA points to origins of modern domestic horses
20/10/2021	The New York Times. Sabrina Imbler, The horse you rode in on may have been made in Southern Russia
20/10/2021	Science. Ann Gibbons, Ancient DNA reveals the long-sought homeland of modern horses
19/09/2021	Pour la Science. Jean-Luc Voisin, L'ADN fossile, une machine à remonter le temps
08/08/2021	Sciences et Avenir. Bernadette Arnaud, L'origine des Scythes révélée par la génétique
17/07/2021	Science & Vie. Anne Debroise, Nous sommes tous des nomades des steppes
09/03/2021	National Geographic. Mehdi Benmakhoul, Quand la paléogénétique bouleverse l'Histoire
01/03/2021	La Nouvelle République. Ambre Philouze-Rousseau, Ludovic Orlando: « La paléogénétique a permis de défantasser le passé »
26/02/2021	Les Échos. Jessica Berthereau, Quand l'ADN fossile redessine le passé...
17/02/2021	Nature. Ewen Callaway, Million-year-old mammoth genomes shatter record for oldest ancient DNA

17/02/2021 Science. Michael Price, Mammoth molars yield the oldest DNA ever sequenced
 24/01/2021 La Dépêche du Midi. Sophie Vigroux, Ludovic Orlando redonne un visage aux hommes du passé
 20/01/2021 L'Écho. Simon Brunfaut, La mémoire du monde tient dans une molécule d'ADN
 06/07/2020 Cheval Magazine. Marie Sénechal, Une préférence pour les chevaux mâles durant l'Âge du Bronze ?
 03/05/2019 5000 ans d'histoire du cheval domestique. Jean-Luc Nothias. Le Figaro
 02/05/2019 Speedy horses evolved only recently. Jason Bittel. National Geographic
 02/05/2019 Ancient DNA reveals two lost lineages of horses—but not their elusive origins. Elisabeth Pennisi. Science
 25/02/2018 La domestication du cheval revisitée. Nathaniel Herzberg. Le Monde
 24/02/2018 How DNA proved wild horses no longer exist. Sarah Gibbens. National Geographic
 23/02/2018 Genetic tests rein in myth of wild horses. Oliver Moody. The Times
 29/04/2017 Les leçons des chevaux des Scythes. Le Figaro. Jean-Luc Nothias
 27/04/2017 Long-frozen DNA shows how humans made horses faster. Washington Post. Ben Guarino
 27/04/2017 Ancient horse DNA shows Scythian warriors were adept domesticators. New York Times. Kenneth Chang
 28/12/2015 The horse that can endure Siberian winters. BBC Earth. Jane Palmer
 11/11/2015 The world's last truly wild horse is making a comeback. BBC Earth. Jane Palmer
 31/08/2015 The taming of the pig took some wild turns. Science. Elizabeth Pennisi
 16/02/2015 La domestication, une affaire de gènes. Le Monde. Catherine Mary
 19/12/2014 The thoroughly bred horse. Science. Ann Gibbons
 17/04/2014 How to build a Neanderthal. Nature. Ewen Callaway

Open Conferences & Exhibition [Non-scientific audience]

04/05/2022 Les Mystérieuses Origines du Cheval Domestique. Académie des Sciences de Toulouse
 31/01/2022 L'ADN fossile : une véritable machine à remonter le temps. Association Club Enjeux de l'Alt, Tournefeuille
 25/12/2021 World of Antiquity Youtube Channel. Who domesticated the horse?
<https://www.youtube.com/watch?v=HwQQM6NvKSg&t=2463s>
 15/11/2021 L'ADN fossile, une machine à remonter le temps Association CVA Lyon
 26/10/2021 L'ADN des fossiles, une machine à remonter dans le temps. AFIS.
<https://www.youtube.com/watch?v=6x1-ch8chMw>
 15/09/2021 XXVIèmes Estivales de Malepère, Arzens. Voyager dans le temps avec l'ADN fossile
<https://www.youtube.com/watch?v=nIVm9Q214pE>
 22/09/2021 Voyager dans le temps avec l'ADN fossile. Lycée Berthelot, Toulouse
 01/08/2021 Le Salon du Livre Préhistorique de la Chapelle-aux-Saints. Table Ronde, ADN : les clefs pour comprendre
 20/05/2021 Festival L'Histoire à Venir. L'ADN dans tous ses états
<https://www.youtube.com/watch?v=xu6HjB33HLE&feature=youtu.be>
 13/12/2019 Voyager dans le temps avec l'ADN fossile. Maison de la Philosophie, Toulouse, France
 19-31/10/2019 Festival Je veux Savoir. Quai des Savoirs, Toulouse, France
 25/05/2019 Histoire à venir. Muséum d'Histoire Naturelle de Toulouse. Discussion with J-M Geneste & N Teyssandier
 14/05/2019 Les grands séminaires de l'Observatoire Midi-Pyrénées. L'origine et l'impact du cheval dans l'histoire de l'humanité.
 12/10/2018 MétamorphOse, 3ème Rendez-vous de l'Innovation.
 13/11/2017 A DNA ride on the horse of the Scythian nomads. Turku, Finland
 30/11/2016 How early nomad riders shaped the modern horse. Videnskabsgalla 2016. Copenhagen, Denmark.
 04/10/2016 Reconstructing the (epi)genomes and microbiomes from ancient DNA fragments. Rejkjavik, Iceland
 12/02/2016 Darwin Day at the Centre for Ecological and Evolution Synthesis. Oslo, Norway

Cover pictures of 17 Scientific Journals [including 13 in the last five years: PNAS (2), Nat Rev Genet (1), Genetics (1), Genome Res, Science (3), Cell (3), Mol Ecol (1), Mol Ecol Res (2)]