RESUME

BAPTESTE, Eric; ORCID: 0000-0003-1966-1215; Date of birth: 18/05/1978; Nationality: French

URL for web site: www.evol-net.fr // https://twitter.com/aire_team

• EDUCATION

2010 : HDR (*Highest academic degree in France, for PhD supervision*), Sorbonne Université/France.
2007: PhD in Philosophy of Biology (2nd)/IHPST/Paris I- Panthéon Sorbonne Université/France.
2003–2007: Post-doctoral fellow in microbial evolution, Doolittle lab/Dalhousie University/Canada
2003: PhD in Evolutionary Biology (1rst)/Université Pierre et Marie Curie/ France.

• CURRENT POSITION(S) & PREVIOUS POSITIONS

2015 – now **CNRS Research Director (DR2)** / UMR 7205 'Institute of Systematics, Evolution and Biodiversity'/ Sorbonne Université/France. **Group leader of the AIRE team** with P. Lopez.

2007 – 2015 CNRS Researcher/ UMR 7138 'Systematics, Adaptation & Evolution'/ Sorbonne Université/ France. Since 2009, group leader of the AIRE team with P. Lopez.

• FELLOWSHIPS AND AWARDS AND SELECTED ESTEEMED INDICATORS

- 2023 **Associate Prof.** at the Dpt of microbiology and infectiology, Faculty of Medecine and Health Sciences, **Sherbrooke University**, Canada
- January 2023 Invited Plenary Speaker (main stage, live on TV) at 'XII Congreso Futuro' (Santiago, Chile) (millions of viewers, and >30 000 attendees, 4th largest event of scientific diffusion in the world)
- **Grant ATM MNHN** "Major phylogenetic extension of evolutionary theories of ageing : first report of cellular ageing and rejuvenation in Archaea" (Bapteste PI : 14 000 euros)
- 2021-2024 Grant 'New tools, methods, and resources for aquatic symbioses', Gordon and Betty Moore Foundation (\$749,000) to B. Llorente (PI) (Bapteste co-PI: \$118 000)
- 2021-2023 Grant Émergence from Alliance Sorbonne Université, 'Evolutionary, systemic, and philosophical dimensions of ageing' (55 000€, Bapteste: PI)
- 2020-2023 **CNRS funding** for the **Thematic Pluridisciplinary Network** 'Epistemic impacts of microbiome studies' (15 000 €, Bapteste: PI)
- January 2019 Invited Plenary Speaker (main stage, live on TV): 'Evolution in the Web of Life', at the 'VIII Futures Congress', (Santiago, Chile), most important open-access scientific event in Latin America, with Nobel Prizes (millions of viewers, and >30 000 attendees)
- 2014 now ERC Consolidator Investigator Grant (PI, 1,2284,240 euros)
- 2009 2019 CNRS Prime for Scientific Excellency (PES, competitive)
- 2009 Paoletti Prize for outstanding research, <u>http://www.cnrs.fr/insb/recherche/prix-2009.htm</u>.
- 2002 Jacques Lebbe Prize, awarded by French Society of Systematics

My work is regularly featured in <u>internationally distributed, renowned media</u> (<u>Press</u>: *Le Monde* (04/2013, 04/2018), *Libération* (11/2015), *Pour la Science* (04/2016, 11/2016, 03/2018, 11/2020), *Mediapart* (2018), *Philosophie Magazine* (04/2016), *New Scientist* (01/2009,03/2011,11/2015,04/2019,06/2021); <u>Radio</u> <u>interviews</u>: *France Inter* (2013/2018), *RFI* (2013/2018), *France culture* (2014), *Radio Anthropocène* (2022), *Radio Cooperativa* (2023) (a major Chilean radio); <u>TV interviews</u>: in 2023, AVANCE 24 HORAS TVN; CNN Chile. Thanks to my <u>well-selling science books</u>, I was plenary speaker in <u>nationally renowned</u> <u>conferences</u>: with *Jean-Claude Ameisen* (2018), *at the Cité des Sciences* (2013, 2023), at the *Festival d'Astronomie de Fleurance*, opened by the French Minister of Research (2017). I also provide public talks and consulting activities for <u>big museums</u>: Centre Pompidou (2022), Palais de la Découverte (2023).

• SUPERVISION OF GRADUATE STUDENTS AND POSTDOCTORAL FELLOWS SINCE 2008 2008-2023: (co)-supervision of *14 master/11 PhD/9 Postdoc* in evolutionary bioinformatics, except *.

<u>Master</u>: 2008-2010: B. Chaeib/U. Pierre & Marie Curie/France; 2011: T. Hujsa/U. Paris 7/France; 2013: M. Grenie/ENS Lyon/France; 2014: M. Rouass/U. Pierre & Marie Curie/France; 2016: A. Danzon/U. Pierre & Marie Curie/France, A. Jaffe/Harvard/USA; 2018: A. Lestage & L. Cavaud/U. Pierre & Marie Curie/France; 2020: P. Martin/U. Nantes/France; 2021: D. Sussfeld & H. Bonnefous/ENS Saclay/France; 2022: A. Omarjee & Y. Dai/U. Paris Cité/France; 2023-present: M. Toucourou/U. Paris-Est Créteil/France. PhD: 2010-2013: C. Bicep/U. Pierre & Marie Curie/France; 2011-2014: PA Jachiet/U. Pierre & Marie Curie/France; 2013-2016: R. Méheust/U. Pierre & Marie Curie/France; 2014-2017: J. Pathmanathan/U. Pierre & Marie Curie/France; C. Vigliotti/MNHN/France; 2016-2019: R. Lannès/U. Pierre & Marie Curie/France; 2019-2021: F. Papale*/U. Montréal/Canada (*Philosophy of sciences); 2019-2022: C. Bernard/U. Pierre & Marie Curie/France; 2021-present: H. Bonnefous/U. Pierre & Marie Curie/France, D. Sussfeld/U. Evry Val d'Essonne /France; 2023-present: Y. Dai/U. Sherbrooke/Canada <u>Postdoc</u>: 2008-2009: J. Leigh/U. Pierre & Marie Curie/France; 2009-2011: K. Schliep/MNHN/France; 2011-2013: S. Karkar/U. Pierre & Marie Curie/France; 2014-2016: E. Corel/U. Pierre & Marie Curie/France; 2015-2017: M. List*/U. Pierre & Marie Curie/France (*linguistics); 2016-2019: A. Watson/U. Pierre & Marie Curie/France; 2017-2018: D. Zea/U. Pierre & Marie Curie/France;2017-2020: G. Bernard/U. Pierre & Marie Curie/France; 2018-2023: J. Teulière/U. Pierre & Marie Curie/France

• **TEACHING ACTIVITIES (only punctual: a CNRS researcher has no teaching activities)** I have taught evolutionary microbiology *classes I created* (master level) at the Ecole Normale Supérieure, Paris, at U. Clermont-Ferrand, and in average in one different university outside France each year: Cortona, Santiago, Kaiserslautern, etc. I also created and taught PhD level classes on networks and evolution through the International Summer School on Networks and Evolution (Roscoff, France, 2016/2018/2019).

• ORGANISATION OF SCIENTIFIC MEETINGS (see also the 2nd part of my CV)

2014: Co-organizer with T. Dagan of a Satellite Meeting SMBE/Germany; 2014-2016: Organizer of the EVOLUNET seminar series/France; 2016: Organizer of the 1rst international summer school on networks and evolution/France; 2017: Organizer of the International Symposium on network and evolution/France; 2018: Organizer of the 2nd international summer school on networks and evolution/France & Co-organizer of the Conference 'Thinking Evolution'/France; 2019: Organizer of the 1rst colloquium on evolutionary networks/France & of the 3rd international summer school on networks and evolution/France; 2020: Organizer of the 2nd Networks and Evolution Conference/France; 2021-2022: Organizer of the New Challenges Induced by Microbiomes Colloquium, 1rst, 2nd and 3rd editions; 2022: Organizer of the 1rst expanding evolutionary theories of ageing Colloquium

• SELECTED RESPONSIBILITIES & REVIEWING ACTIVITIES

Grant reviews for: 2022: the Israel Science Foundation (ISF), the Irish Research Council (IRCLA), 2021: the Canada Foundation for Innovation (CFI)/ John R. Evans Leaders Fund; 2014: the US-Israel Binational Science Foundation, the Programma Nazionale di Ricerche in Antartide (Italy), 2013: the Center for Advanced Study, Univ. of Illinois, USA, 2010: the US-Israel Binational Science Foundation. HDR review: External for J. Filée (CNRS, 2023). PhD reviews: External for L. Pradier (U. Montpellier, Fr., 2022), S. Harzallah Debbabi (U. Jean-Moulin Lyon 3, Fr., 2018), V. Keshri, (U. Aix-Marseille, Fr., 2016 PhD rejected), O. Poirion (Ecole Centrale Lyon, Fr. 2014), E. Skippington (The University of Queensland, 2012), C.A. Michael (Macquarie University, 2008), Evaluator for H. Gardon (U. Clermont Auvergne, Fr., 2022), Committee member for B. Churcheward (Université Nantes, Fr., 2021), M. Hénnart (Pasteur, Fr., 2021), M. Hénnart (Pasteur, Fr., 2020), R. Tignat-Perrier (Ecole Centrale Lyon, Fr., 2017), L. Urbini (Université de Lyon 1, Fr., 2015-2016). Associate Editor: (2021-): Microbiology Research, the German Journal of Microbiology, (2019-): Microorganisms, Biology, (2014-2020): Genome Biology & Evolution, (2008-2020): BMC Research Notes, (2006-2016): Biology Direct. Moreover, I am a regular reviewer for Genome, Trends in Genetics, PloS ONE, BioEssays, Genome Research, Proceedings of the National Academy of Sciences, Molecular Biology and Evolution, Journal of Molecular Evolution, Environmental Microbiology, The International Journal of Systematic and Evolutionary Microbiology, Nucleic Acid Research, Genome Biology and Evolution, BMC Bioinformatics, etc.

• MAJOR COLLABORATIONS

<u>Network and statistics</u>: M. Habib & L. Viennot (University Paris Diderot, Fr.), FJ. Lapointe (U. Montréal, Canada), P. Gambette (U. Marne-la-Vallée, Fr.), S. Chaffron & D. Eveillard (U. Nantes, Fr.); <u>Microbial ecology and evolution</u>: I. Ruiz-Trillo (Barcelona, Spain), D. Bhattacharya (Rutgers, USA), M. Krupovic (Pasteur, France), LP Haraoui (U. Sherbrooke, Canada), L. Hug (U. Waterloo, Canada), C. Larose (Ecole Centrale Lyon, France), F. Not (Roscoff, Fr.), J. Mcinerney (Nottingham, UK); <u>Philosophy of biology</u>: P. Huneman (CNRS, France), J. Dupré (Exeter, UK), F. Bouchard (U. Montréal, Canada); Ageing: A. Baudisch (University of Southern Denmark, Denmark), E. Teeling (University College Dublin, Ireland), W. Ludington (Carnegie Science, USA), C. Franceschi (U. Bologna, Italy)

Ten years track-record

1.Ten significant publications over the last 10 years (out of 52 articles since 2013; supervised students)

In total, I have published **99** papers (25 as first author and 50 as last author, H-index: **43**, > **7273** citations). -Alvarez-Ponce, Lopez, <u>Bapteste</u>, McInerney. (2013) Gene similarity networks provide tools for understanding eukaryote origins and evolution. **PNAS.** 110(17): E1594-603. [*New approach*] -*Corel*, Lopez, *Méheust*, <u>Bapteste</u>. (2016) Network-Thinking: Graphs to Analyze Microbial Complexity and Evolution. **Trends Microbiol.** 24(3):224-37. [*Creative thinking*] -Méheust, Zelzion, Bhattacharya, Lopez, Bapteste. (2016) Protein networks identify novel symbiogenetic genes resulting from plastid endosymbiosis. PNAS.113(13):3579-84. [New phenomena]

-Méheust, Watson, Lapointe, Papke, Lopez, Bapteste. (2018) Hundreds of novel composite genes and chimeric genes with bacterial origins contributed to haloarchaeal evolution. Genome Biology. 19(1):75. [5year Impact Factor: 13.6] [Creative thinking, new findings]

-Watson, Habib, Bapteste. (2020) Phylosystemics: Merging Phylogenomics, Systems Biology, and Ecology to Study Evolution. Trends Microbiol. 28(3):176-190. [Creative framework for evolutionary studies]

-Bernard, Yanyan, Lopez, Bapteste. (2021) Beyond arbitrium: identification of a second communication system in Bacillus phage phi3T that may regulate host defense mechanisms. ISME J. 15(2):545-549. [Impact Factor in 2020: 11.217] [Original analyses of microbial functions and evolution]

-Bernard, Teulière, Lopez, Corel, Lapointe, Bapteste. (2022) Aging at Evolutionary Crossroads: Longitudinal Gene Co-expression Network Analyses of Proximal and Ultimate Causes of Aging in Bats. Mol Biol Evol. 39(1):msab302. [5-year Impact Factor: 14.387] [Original transcriptomics network ageing studies]

-Ocaña-Pallarès, (...), Pathmanathan, Bapteste, Tikhonenkov, Keeling, Szöllősi, Ruiz-Trillo. (2022) Divergent genomic trajectories predate the origin of animals and fungi. Nature 609, 747-753 (2022). [Fruitful training of a visiting scientist, altmetric: top 1%]

-Teulière, Bernard, Corel, Lapointe, Martens, Lopez, Bapteste. (2023) Network analyses unveil ageingassociated pathways evolutionarily conserved from fungi to animals. Geroscience. 45(2):1059-1080. [5-year Impact Factor: 7.608 [Original phylosystemics analyses of ageing]

-Teulière, Bernard, Bonnefous, Martens, Lopez, Bapteste. (2023) Interactomics: Dozens of Viruses, Coevolving With Humans, Including the Influenza A Virus, may Actively Distort Human Aging. Mol Biol Evol.;40(2):msad012. [5-year Impact Factor: 14.387][Original network studies of ageing; altmetric: top 5%] 2. Research monographs

I published 5 single-author synthetic science books: "Les gènes voyageurs", Belin 2013; "Conflits intérieurs", Ed. Matériologiques 2015; "Tous entrelacés!" Belin 2018; "Tout se transforme! Comment marche l'évolution". Ed. Circonflexe 2022; "Le monde surprenant des microbes" Ed. Circonflexe 2022.

selected invited presentations to internationally established 3. 10 conferences -01/2023 : Plenary Speaker (main stage, live on TV) at 'XII Congreso Futuro' (Santiago, Chile) (millions of viewers, and >30 000 attendees, including Nobel Prizes, 4th largest event of scientific diffusion in the world). -01/2019: Plenary Speaker (main stage, live on TV) at the 'VIII Futures Congress' (Santiago, Chile)

-07/2019 : Speaker at 'the SMBE 2019 meeting', Manchester, UK.

-10/2018: Speaker at the International IRIF workshop: 40 Years of Graphs and Algorithms. Paris, France.

-08/2017: Speaker at the 15th International Congress of Protistology (ICOP 2017). Prague, Czech Republic.

-03/2016: Speaker at the International BioPerspectives CNRS School. Cargèse, France.

-10/2015: Speaker at International workshop The Origin of metazoans. Presqu'île de Giens; France.

-07/2015: Speaker at The Phylogenetic Network Workshop. Singapore

-07/2015: Speaker at the International Symbiosis Society. Lisboa, Portugal.

-08/2013: Speaker at the 10th International Phycological Congress, Orlando, Florida, USA.

4. Selected organisations of international conferences

2014 : Co-organizer of the Satellite Meeting SMBE on Reticulated Microbial Evolution/Germany (76 speakers and attendees); 2016-2019: Organizer of the 1rst, 2nd and 3rd editions of the international summer school on networks and evolution/France (27 speakers, 36 selected early carrier researchers, hands-on sessions); 2018 : Co-organizer of the **Conference : 'Penser l'évolution'**/France (20 speakers, 60 attendees); 2019-2020 : Organizer of the 1rst and 2nd colloquium on evolutionary networks/France (11 speakers, 90 attendees); 2021-2022 : Organizer of the New Challenges Induced by Microbiomes Colloquium, 1rst, 2nd and 3rd edition (38 speakers, 270 attendees) – the 4th edition will occur in October 2023. ; 2022 : Organizer of the **1rst** (soon 2nd) expanding evolutionary theories of ageing Colloquium (14 speakers, 70 attendees). 5. Prizes : 2009: Paoletti Price, prestigious award by the CNRS

6. Major contributions to the carriers of 22 excellent researchers

In the past 10 years, I directly supervised/co-supervised: 1 visiting college student: A. Jaffe (2016, 1 coauthored paper in Env. Micro), who next joined the Banfield lab (University of California, Berkeley, USA) for a PhD, 1 visiting PhD student: Dr. E. Ocaña, who developed outlines and methods for a Nature article during his stay in my lab; 8 PhD students: C. Bicep (2010-2013, 5 co-authored papers), who next joined the Embley lab (Newcastle University, UK) as postdoc; PA Jachiet (2011-2014), now data-analyst in a private company; J. Pathmanathan (2014-2017, 6 co-authored papers), who next joined the Landweber lab (Columbia University, USA) as postdoc and is now data-analyst in a private company; R. Méheust (2013-2016, 8 co-authored papers, including a PNAS), who next joined the **Banfield lab** (University of California, Berkeley, USA) as postdoc and is now permanent CEA researcher); C. Vigliotti (2014-2017), who next joined the Institute of Cardiometabolism and Nutrition, Clément lab, as postdoc and is now Assistant Prof

at AgroParisTech; R. Lannes (2016-2019), who next joined the CRCT (Toulouse, ONCOPOLE, France) as postdoc and is now data scientist at the Whitehead Institute; C. Bernard (12/2018-now, 1 book chapter, 2 co-authored papers), who next joined the Dessimoz Lab (UNIL, Switzerland) and is now with E. Rocha at **Pasteur** as postdoc; F. Papale (09/2018-now, 2 co-authored papers, a top co-authored paper in Trends in Micro), who then joined the Doolittle lab (Dalhousie, Canada) as postdoc. I also supervised 6 post-doctoral fellows: S. Karkar (2011-2013, 3 co-authored papers), who next joined the Bhattacharva lab (Rutgers U., USA) as postdoc and is now Assistant Prof at Bordeaux University; E. Corel (2014-2017, 4 co-authored papers), who is now Assistant Professor at UPMC in my team; A. Watson (2016-now, 7 co-authored papers), who next joined the Errington lab (Newcastle University, UK) as postdoc; G. Bernard (2017-04/2019, 2 coauthored papers), who next joined the Bowler lab (ENS, Paris); D. Zea (5 months in 2017/18), who become permanent researcher at CONICET, Argentina and is now Assistant Professor at Paris Saclay); J. Teulière (12/2018-now, 6 co-authored papers, top paper in Ageing Research Reviews); who is applying for Assistant professor position. I also hosted and co-supervised 1 research fellow (Dr. M. List, 01/2015-12/2016, 5 coauthored papers, top paper in the Association of Computational Linguistics Conference, the 2nd best journal in linguistics), who earned a highly competitive ERC starting grant, became Group leader and senior scientist at the Max Planck Institute, Jena, Germany. In addition, Dr. Lopez, member of my former ERC team from 2014-2019 earned the highly competitive permanent position of Full Professor at UPMC.

7. Examples of innovation leadership :

During the last 10 years, my two PhD degrees in Evolutionary Biology and in Philosophy of Sciences allow me i) to lead original interdisciplinary researches at the interface of evolutionary biology, bioinformatics, microbiology, ecology, graph theory, synthetic biology, human sciences and ageing, ii) to gather a powerful network of collaborators and iii) to develop tools and concepts for network-thinking in evolutionary biology. My work unveiled novel kinds of remodelled genes (S-genes, Chic-Genes and OR-genes) in prokaryotic and eukaryotic genomes (PMID: 29880023, 29534719, 34792602); the contribution of ultra-small marine microbes to the cycling of Carbon, Methane, Nitrogen and Sulfur (PMID: 30903144, 33325996); highly diverse quorumsensing systems in CPR and DPANN (PMID: 33051376); the existence of phages and plasmids with multiple quorum-sensing systems (PMID: 33028977, PMID: 36929912); the evolution of robust genetic worlds despite gene transfers (PMID: 29346651); the existence of uncultured oceanic phyla closely related to Holozoa with potential to enlight the origins of animals (PMID: 32533833), etc. In vast majority, the masters, PhD and postdocs I trained during this time pursued a successful academic carrier. Recently, I geared my research to tackle the evolution of ageing from a network and microbial perspective, first using overheads from my ERC Consolidator grant, then by earning an Emergence grant (Sorbonne-University). I investigated the contribution of mitochondria to protist ageing (PMID: 33072737), introduced the original concept of age-distorters (PMID: 34082078), developed novel comparative methods of gene co-expression networks to unveil proximal and ultimate causes of ageing (PMID: 34662394), and predicted that cell division in microbes supports labor division with respect to ageing and rejuvenation (PMID: 32268207), earning an ATM grant to conduct experimental research on archaeal ageing. Based on these works, published in solid scientific journals, attracting some media attention (New Scientist, CurrentScienceDaily, Elementy.ru, Radio Cooperativa, TVN & CNN Chile), I gave international talks, started to organize international conferences on ageing, and new collaborations. I am also leader of a Thematic Pluridisciplinary Network funded by the CNRS, co-PI in an international research project funded by the Moore Foundation and provide my original expertise to a large Prioritary Research Program (PPR Océan et climat).