Curriculum Vitae for Jessica Louise Ray

PERSONAL INFORMATION

Name: Ray, Jessica Louise

Address: Øyjordsveien 27, 5038 Bergen, Norway

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Date of birth: 05.12.1976

Sex: Female

Nationality: Norway and United States of America Web of Science ResearcherID: D-2210-2018

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EDUCATION

MSc Microbial Biology, University of California at Berkeley, California, USA 2002 2007 PhD, University of Tromsø The Arctic University of Norway, Norway (21.04.2007) Title: Natural transformation with chromosomal DNA in bacteria: occurrence and

determinative factors

CURRENT AND PREVIOUS POSITIONS

2007-2008	Postdoctoral Researcher, Lincoln University, New Zealand
2009-2012	Postdoctoral Researcher, University of Bergen, Norway
2012-2015	Postdoctoral Researcher, Uni Research Environment, Bergen, Norway
2016-2020	Adjunct researcher (20% Researcher II), University of Bergen, Norway
2015-2018	Researcher (Researcher II), Uni Research Environment, Bergen, Norway
2018-	Senior Researcher, NORCE Norwegian Research Centre AS, Norway

FELLOWSHIPS AND AWARDS

2007	AgMARDT Postdoctoral Fellowship, Lincoln University, New Zealand
2009	RCN FRIBIO Mobility Grant ("VIPMAP" nr. 186142), 140 kNOK
2013	Meltzer Foundation travel scholarship, 26.8 kNOK
2014	COST-EU STSM, host University of Glasgow, UK, 1700 EUR
2014	COST-EU STSM, host Friedrich Schiller Univeristy of Jena, Germany, 1350 EUR
2018	RCN KLIMAFORSK Mobility Grant ("aDNAPROX" nr 268062), 124 kNOK

MOBILITY

2003	Plant Research International, The Netherlands. 3-wk research visit (J.D. van Elsas)
2004	AgResearch, Canterbury, New Zealand. 6-wk research visit (M. O'Callaghan)
2004	Plant Research Int'l, The Netherlands. 5-wk research visit (L. van Overbeek)
2009	Rutgers University, 6-month research visit (K.D. Bidle)
2010	Université Paris-Sud, Orsay, France. 3-wk bioinformatics training (M.S. Dubow)
2014	University of Glasgow, UK. 2-wk bioinformatics training (C. Quince, U.Z. Ijaz)
2014	Friedrich Schiller University Jena. 2-wk data analysis (G. Pohnert)
2017	IS-AURORA. 4-day research exchange visit to Ifremer (R. Siano, K. Mertens)
2018	University of Glasgow, Scotland, UK. 6-mo bioinformatics training (U.Z. Ijaz)

SUPERVISION OF GRADUATE STUDENTS AND RESEARCH FELLOWS

Main supervisor: 1 PhD, 2 MSc; Co-supervisor: 2PhD, 3 MSc

MSc supervisor Anine Veronica Grønlund, University of Bergen, Norway (2016-2018); Rachel Lima, University of Bergen, Norway (2019-2023).

PhD Supervisor Danielle Grant (2019-2023)

PhD Co-Supervisor Julia Endresen Storesund, University of Bergen, Norway (2011-2016); Torill Vik Johannessen, University of Bergen, Norway (2010-2015).

MSc Co-Supervisor Arna Kazazic, Uni Research Environment (2015-2017); Heidi Osmundsen, University of Bergen (2011-2013); Helga Andersen, University of Tromsø (2004-2005); Marius Nilsen, University of Bergen (2019-2020)

RESEARCH PROJECT FUNDING AS PROJECT LEADER

2018-2021 VIROVAC, Research Council of Norway (RCN) project 275710, 9.9 MNOK	2018-2021	VIROVAC,	Research	Council of	Norway	(RCN)	project	275710,	9.9 MNOK
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2017-2018 RECNOR, Institute of Marine Research pilot initiative, 500 kNOK

2016-2017 ABRes. research commission from Norwegian Environmental Agency, 500 kNOK

2021-2024 PATHDNA, Research Council of Norway project 236900, 12 MNOK

ORGANISATION OF SCIENTIFIC MEETINGS

2011	Session Co-Chair at ASLO Aquatic Sciences Meeting, San Juan, Puerto Rico
2018	Co-organizer, 2nd Workshop on eDNA for Fisheries Mgmt, Tromsø, Norway
2019	Co-organizer, 3rd Workshop on eDNA for Fisheries Mgmt, Bergen, Norway

INSTITUTIONAL RESPONSIBILITIES

Member of PhD appointment committee, University of Bergen, Norway
External sensor for evaluation of MSc thesis, University of Tromsø, Norway
Member of PhD appointment committee, University of Bergen, Norway
External sensor for evaluation of MSc thesis, University of Tromsø, Norway
Member of committee for laboratory renovation, Uni Research Environment
Member of PhD appointment committee, University of Bergen, Norway
Member of Postdoctoral appointment committee, University of Bergen, Norway
External sensor for evaluation of MSc thesis, University of Tromsø, Norway
External sensor for evaluation of MSc thesis, University of Tromsø, Norway

COMMISSIONS OF TRUST

Scientific Advisory Board, Hjort Centre for Marine Ecosystem Dynamics, Bergen, Norway (2015-2017)

Invited Round Table Participant, eDNA Workshop, Woods Hole Oceanographic Institute, USA (2018-present)

Co-Editor, NOAA and IMR internal white paper on eDNA as a tool for fisheries stock assessments and ecosystem-based fisheries management (2020)

Reviewer for Journals: Environmental Microbiology, Aquatic Microbial Ecology, Journal of Phytoplankton Research, Nature Scientific Reports, FEMS Microbiology Ecology, Viruses, Nature Ecology and Evolution, Marine Ecology Progress Series, Antibiotics, Water, Environmental Pollution

MEMBERSHIPS OF SCIENTIFIC SOCIETIES

2009	Association for the Sciences of Limnology and Oceanography (ASLO), regular

member

2010 American Association for the Advancement of Science (AAAS), regular member

SEA-GOING EXPEDITIONS AND FIELD WORK

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COURSES AND CERTIFICATION

2019	Cruise Leader Course, Institute of Marine Research, Bergen, Norway
2019	Survival Suit Course, RelyOnNutec, Bergen, Norway
2021	Seafarer's Health Certificate, Bergen, Norway

MAJOR COLLABORATIONS

Ruth-Anne Sandaa, Gunnar Bratbak, University of Bergen: marine microbial food web; Stijn De Schepper, NORCE Climate and Bjerknes Centre for Climate Research, molecular palaeoceangraphy - sedimentary ancient DNA as sea ice proxy (aDNAPROX and AGENSI). Kay D. Bidle, Rutgers University, USA: programmed cell death in marine phytoplankton; Janice Lawrence, University of New Brunswick, Canada: role of appendicularians (Chordata:Tunicata)

in marine viral ecology; **Umer Z. Ijaz**, University of Glasgow: bioinformatic and statistical analysis of high-throughput sequencing data; **Olav Kjesbu**, **Martina Stiasny**, Institute of Marine Research, Bergen, Norway: molecular quantification of predation pressure on Norwegian spring-spawning herring (RECNOR); **Anders Lanzén**, AZTI, Spain: bioinformatic and statistical analysis of sediment metabarcoding for environmental monitoring (MetaMon); **Eric Thompson**, University of Bergen: metabarcoding for environmental monitoring and appendicularian trophic ecology; **Kristi Miller-Saunders**, Fisheries and Oceans Canada: quantitative molecular monitoring of fish pathogen eDNA/eRNA in association with sea-based open aquaculture

TRACK RECORD

Scientific Experience and Achievements

Since the completion of my PhD in 2007 I have worked to establish competence in the field of molecular microbial ecology, first in terrestrial ecosystems and now in aquatic environments. In addition to my project leader role in two research projects (VIROVAC and PATHDNA), I am work package leader for two research projects, including the ERC Consolidator Grant *AGENSI* (818449) (see Major Collaborations above) and the Norway-Poland collaborative project *NEEDED*. My role is these projects is to oversee the development and implementation of molecular biological research tools for the investigation of basic and applied ecology in aquatic environments. In the Molecular Ecology Research Group at NORCE Environment I am also participating in national and international initiatives to map, benchmark and utilize eDNA for the purposes of environmental monitoring.

In the track record, please list:

The total <u>number</u> of publications during the career: 25

Publications (source for number of citations: Publons and GoogleScholar)

2004. Koenig RL, **Ray JL**, Maleki SJ, Smeltzer MS, Hurlburt BK. *Staphylococcus aureus* AgrA binding to the RNAIII-*agr* regulatory region. *Journal of Bacteriology* 15, 7549-7555. (123 citations) **2009. Ray JL**, Harms K, Wikmark O-G, Johnsen PJ and Nielsen KM. 2009. Sexual isolation in *Acinetobacter baylyi* is locus-specific and varies 10,000-fold over the genome. *Genetics* 182, 1165-1181. (23 citations)

2009. Clough TJ, **Ray JL**, Buckthought LE, Calder J, Baird D, O'Callaghan M, Sherlock RR and Condron LM. The mitigation potential of hippuric acid on bovine urine N2O emissions: an *in situ* determination of its effect. *Soil Biology and Biochemistry* 41, 2222-2229. (39 citations)

2011. Ogata H, **Ray JL**, Toyoda K, Sandaa R-A, Nagasaki K, Bratbak G and Claverie J-M. Two new subfamilies of DNA mismatch repair proteins (MutS) specifically abundant in the marine environment. *ISME Journal* 5, 1143-1151. (40 citations)

2012. **Ray JL**, Dondrup M, Modha S, Steen IH, Sandaa R-A and Clokie MJ. Finding a needle in the virus metagenome haystack - micro-metagenome analysis captures a snapshot of the diversity of a bacteriophage armoire. *PLoS One* 7, e34238. *doi: 10.1371/journal.pone.0034238*. (17 citations)

2012. **Ray JL**, Töpper B, An S, Silyakova A, Spindelböck J, Thyrhaug R, DuBow MS, Thingstad TF and Sandaa R-A. Effect of increased *p*CO2 on bacterial assemblage shifts in response to glucose addition in Fram Strait seawater mesocosms. *FEMS Microbiology Ecology* 82, 713-723, *doi:* 10.1111/j.1574-6941.2012.01443.x.(13 citations)

2014. Ray JL, Haramaty L, Thyrhaug R, Fredricks H, Van Mooy B, Larsen A, Bidle KD and Sandaa R-A. Virus infection of *Haptolina ericina* and *Phaeocystis pouchetii* implicates evolutionary conservation of programmed cell death induction in marine haptophyte–virus interactions. *Journal of Plankton Research 36*, 943-955. (6 citations)

2016. Ray JL, Skaar KS, Simonelli P, Larsen A, Sazhin A, Jakobsen HH, Nejstgaard JC, Troedsson CT. Molecular gut content analysis demonstrates that *Calanus* grazing on *Phaeocystis pouchetii* and *Skeletonema marinoi* is sensitive to bloom phase but not prey density. *Marine Ecology Progress Series* 542: 63-77. (5 citations)

2016. Ray JL, Althammer J, Skaar KS, Simonelli P, Larsen A, Stoecker D, Sazhin A, Ijaz UZ, Quince C, Nejstgaard JC, Frischer M, Pohnert G, Troedsson CT. Metabarcoding and metabolome analysis of copepod grazing reveals feeding preference and linkage to metabolite classes in dynamic microbial plankton communities. *Molecular Ecology* 216: 5585-5602. (18 citations) **2018**. Lawrence J, Töpper JP, Petelenz-Kurdziel E, Bratbak G, Larsen A, Thompson E, Troedsson C, Ray JL. Viruses on the menu: The appendicularian *Oikopleura dioica* efficiently removes viruses from seawater. *Limnology* & *Oceanography* 63(S1): S244-S253. (4 citations)

- **2019**. De Schepper S, **Ray JL**, Skaar KS, Sadatzki H, Ijaz UZ, Stein R, Larsen A. The potential of sedimentary ancient DNA for reconstructing past sea ice evolution. *The ISME Journal*, doi:10.1038/s41396-019-0457-1. (<u>4 citations</u>)
- **2020**. Brown TA, Rad-Menéndez C, **Ray JL**, Skaar KS, Thomas N, Ruiz-Gonzalez C, Leu E. Influence of nutrient availability on Arctic sea ice diatom HBI lipid synthesis. *Organic Geochemistry* 20: 103977. *doi: https://doi.org/10.1016/j.orggeochem.2020.103977*. (1 citation)
- **2020**. Kutti T, Johnsen IA, Skaar KS, **Ray JL**, Husa V, Dahlgren TG. Quantification of eDNA to map the distribution of cold-water coral reefs. *Frontiers in Marine Science* 7: 446. *doi: 10.3389/fmars.2020.00446*.
- **2020.** Ray JL, Strom M, Johansen T, Goodwin K (*eds.*) UNIG (U.S.–Norway Intergovernmental Group on eDNA Implementation for Fisheries Stock Assessments and Management). Implementation of Environmental DNA (eDNA) as a Tool for Ecosystem-Based Fisheries Management. U.S. Department of Commerce, NOAA White Paper NMFS- NWFSC-WP-2020-01. doi: https://doi.org/10.25923/e736-vn83.
- **2021**. Állan BJM, **Ray JL**, Tiedemann M, Stiasny MH, Skaar KS, Folkvord A, Nash RDM, Stenevik EK, Komyakova V, Vikebø F, Kjesbu OS. Quantitative molecular detection of Atlantic herring (*Clupea harengus*) identifies regions of high predation pressure by Atlantic mackerel (*Scomber scombrus*) along the Northern Norwegian continental shelf. *Scientific Reports* **11**, 5095 (2021). https://doi.org/10.1038/s41598-021-84545-7
- **2021**. Ævarsson A et al. Going to extremes a metagenomic journey into the dark matter of life. *FEMS Microbiology Letters, in press*.
- **2021**. Mayers KMJ, Lawrence J, Skaar KS, Töpper JP, Petelenz E, Saltvedt MR, Sandaa R-A, Larsen A, Bratbak G, **Ray JL**. Impacts of feeding by the appendicularian *Oikopleura dioica* on large virus abundance and dispersal during *Emiliania huxleyi* bloom conditions. *Limnology* & *Oceanography, in press. doi: 10.1002/lno.11935*.
- **2021**. Dunshea G, Martell L, Bakken T, Budaeva N, Ekrem T, Tandberg AHS, Baussant T, De Boer H, Hestetun JT, Hobæk A, Kallioniemi EP, Larsen A, Markussen SS, Mauvisseau Q, **Ray JL**, Yoccoz N and Willassen E. Kunnskapstatus for bruk av molekylære verktøy i kartlegging og overvåkning av biologisk mangfold i marine miljø. Norwegian Environmental Agency, report M-2062.