

# Guillem Salazar

ETH Zürich. Switzerland

☑guillems@ethz.ch | ☆guillemsalazar.github.io | ☑0000-0002-9786-1493 | ☑GuillemSalazar

I use computational tools to understand how marine microbial communities are shaped and respond to environmental change.

Valencia, SPAIN

Valencia, SPAIN

12-2006

9-2009

3-2010

5-2010

Z<d9>rich, SWITZERLAND

# **Academic Information**

Degree in Biological Sciences Valencia, SPAIN

UNIVERSITY OF VALENCIA 2007

MSc in Biodiversity: Conservation and Evolution

University of Valencia 2008

PhD. in Oceanography

Barcelona, SPAIN

POLYTECHNIC UNIVERSITY OF CATALONIA & MARINE SCIENCES INSTITUTE (ICM - CSIC)

Sunagawa Lab. ETH ZPrich 2020

### **Grants and Contracts**

Introduction to Scientific Research Grant \_ Education and Science Ministry Valencia, SPAIN

University of Valencia and Cavanilles Intitute for Biodiversity and Evolutionary Biology 7-2007

Research Technician - Geronimo Forteza Programme

University of Valencia and Cavanilles Intitute for Biodiversity and Evolutionary Biology 12-2008

Grant for a stay at Dr. Fuhrman Lab

Valencia, SPAIN

University of Southern California (USC), Los Angeles, CA, USA 12-2011

JAE Predoc (Phd Grant/Contract)

Marine Sciences Institute (ICM), CSIC

8-2013

Research Technician Contract

Barcelona, SPAIN

Marine Sciences Institute (ICM), CSIC

6-2014

Research Technician Contract

Barcelona, SPAIN

Barcelona, SPAIN

Marine Sciences Institute (ICM), CSIC

### Courses\_

**Postdoc** 

II Scientific Workshop of ICBiBE

Valencia, SPAIN

and the second s

International Course on Ecological Modelling Valencia, SPAIN

University of Valenciaand the Spanish Society for Limnology and the Spanish Society for Terrestrial Ecology 11-2007

Socrates Intensive Programme Mathematics and Biology

Paris, FRANCE

Universit\_d<cd>Orsay Paris XI y AgroParisTech 7-2008

Famoso III Scientific Cruise (Barcelona \_ Barcelona, Spain)

CAVANILLES INTITUTE FOR BIODIVERSITY AND EVOLUTIONARY BIOLOGY

Citowork Cruise (Barcelona \_ Barcelona, Spain)

Inter-calibration Scientific Cruise \_ Malspina 2010 Expedition (Las Palmas de Gran

Canaria, Spain \_ Cartagena, Spain)

Methods in Microbial Oceanography Course

Barcelona, SPAIN

Marine Sciences Institute (ICM), CSIC 6-2010

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1-2011

Bioinformatics for metagenomics and environmental sequencing

Oslo, NORWAY

MERG AND UNIVERSITY OF OSLO, DEPARTMENT OF BIOLOGY

4-2011

Tara Oceans Scientific Cruise (Bermudas \_ A<8d>ores, Portugal)

3-2012

# **Teaching**

#### Practical applications to epidemiology

Valencia, SPAIN

VALENCIA'S SCHOOL FOR HEALTH STUDIES (EVES)

5-2013

**Biostatistics for Doctors: practical applications** 

Valencia, SPAIN

VALENCIA'S SCHOOL FOR HEALTH STUDIES (EVES)

11-2014

# **Oral** presentations in conferences

Constataci<d1>n emp<d5>rica de un modelo estoc<e0>stico para predecir la extinci<d1>n de especies en interacci<d1>n

SALAZAR, G., ET AL.

moleculares y gen<d1>micas

IX Congreso de Microbiolog<d5>a

II Congreso Nacional de

del Medio Acu<e0>tico

Biodiversidad

2-2009

CORNEJO-CASTILLO, FM., SALAZAR, G., STAL, L., HINGAMP, P., GASOL, J.M., ACINAS, S.G.

Global beta-diversity patterns in deep-ocean bacterial communities

SALAZAR, G., CORNEJO-CASTILLO, FM., ACINAS, S.G., GASOL, J.M.

ASLO Aquatic Sciences Meeting

IV Congreso de Biodiversidad

Exploraci<d1>n de la diversidad microbiana en el oc\_ano profundo: primeros Resultados de la Expedici<d1>n Malaspina 2010

GASOL, J.M., SALAZAR, G., CORNEJO-CASTILLO, FM., PERNICE, M., GOMES, A., MASSANA, R., MOR<EO>N, X.G., ARRIETA, T., DUARTE, C., ACINAS, S.G.

Biogeograf<d5>a y diversidad de procariotas diazotrofos marinos mediante t\_cnicas

Patrones de diversidad globales de procariotas y picoeucariotas en el oc\_ano f<d1>tico y af<d1>tico

SALAZAR G., LOGARES R., CORNEJO-CASTILLO F.M., SUNAGAWA S., BORK P., GASOL J.M. AND ACINAS S.G.

X Congreso de Microbiolog<d5>a del Medio Acu<e0>tico

A global ocean metagenomic survey reveals events of vertical connectivity in oceanic microbial communities

SALAZAR G., CORNEJO-CASTILLO F.M., LOGARES R., SUNAGAWA S., LUDICONE D., LIONEL G., STEMMANN L., LAXENAIRE R., SPEICH S., GASOL J.M., ACINAS S.G.

SAMF 14

9-2014

8-2015

Living on particles in the ocean surface and the bathypelagic: prokaryotic groups involved and phylogenetic conservation of the particle-association trait

MESTRE M., SALAZAR G., CORNEJO-CASTILLO F.M., SALA M.M., ACINAS S.G., GASOL J.M.

SAME 14

8-2015

# Posters in conferences

SALAZAR, G., ET AL.

El papel del canibalismo en copepodos dentro de una red tr<d1>fica sencilla

XI Conferencia Espa<d0>ola y I Encuentro Iberamericano de Biometr<d5>a

6-2007

SAME 12

2-2013

Un modelo nulo para la conectancia de redes de interacci<d1> planta-animal

XI Conferencia Espa<d0>ola y I Encuentro Iberamericano de Biometr<d5>a

SALAZAR, G., ET AL. 6-2007

Diversity patterns of cultured marine bacteria ussing different organic matter sources as compared to single amplified genomes

SALAZAR, G., SARMENTO, H., D<D5>EZ-VIVES, C., FERRERA, I., GASOL, J.M., AND ACINAS, S.G.

Exploraci<d1>n de la diversidad microbiana en en distintos contextos ecol<d1>gicos IV Congreso de Biodiversidad dentro de la expedici<d1>n TARA Oceans

SALAZAR, G., LOGARES, R., CORNEJO-CASTILLO, FM., FERRERA, I., SARMENTO, H., GASOL, JM., ACINAS, S.G.

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## **Publications**

- 1. Ruiz-González, C, M Mestre, M Estrada, M Sebastián, G Salazar, S Agustí, E Moreno-Ostos, I Reche, XA Álvarez-Salgado, XAG Morán, CM Duarte, MM Sala, and JM Gasol (Apr. 2020). Major imprint of surface plankton on deep ocean prokaryotic structure and activity. *Molecular Ecology*.
- 2. Milanese, A, DR Mende, L Paoli, G Salazar, HJ Ruscheweyh, M Cuenca, P Hingamp, R Alves, PI Costea, LP Coelho, TSB Schmidt, A Almeida, AL Mitchell, RD Finn, J Huerta-Cepas, P Bork, G Zeller, and S Sunagawa (Mar. 2019). Microbial abundance, activity and population genomic profiling with mOTUs2. *Nature Communications* **10**(1).
- 3. Massoni, J, M Bortfeld-Miller, L Jardillier, G Salazar, S Sunagawa, and JA Vorholt (Oct. 2019). Consistent host and organ occupancy of phyllosphere bacteria in a community of wild herbaceous plant species. *The ISME Journal* **14**(1), 245–258.
- 4. Salazar, G, L Paoli, et al. (Nov. 2019). Gene Expression Changes and Community Turnover Differentially Shape the Global Ocean Metatranscriptome. *Cell* **179**(5), 1068–1083.e21.
- 5. Eguíluz, VM, G Salazar, J Fernández-Gracia, JK Pearman, JM Gasol, SG Acinas, S Sunagawa, X Irigoien, and CM Duarte (Dec. 2019). Scaling of species distribution explains the vast potential marine prokaryote diversity. *Scientific Reports* **9**(1).
- 6. Ibarbalz, FM et al. (Nov. 2019). Global Trends in Marine Plankton Diversity across Kingdoms of Life. *Cell* **179**(5), 1084–1097.e21.
- 7. Villarino, E et al. (Jan. 2018). Large-scale ocean connectivity and planktonic body size. *Nature Communications* **9**(1).
- 8. Salazar, G and S Sunagawa (June 2017). Marine microbial diversity. Current Biology 27(11), R489-R494.
- 9. Royo-Llonch, M, I Ferrera, FM Cornejo-Castillo, P Sánchez, G Salazar, R Stepanauskas, JM González, ME Sieracki, S Speich, L Stemmann, C Pedrós-Alió, and SG Acinas (July 2017). Exploring Microdiversity in Novel Kordia sp. (Bacteroidetes) with Proteorhodopsin from the Tropical Indian Ocean via Single Amplified Genomes. *Frontiers in Microbiology* 8.
- 10. Lara, E et al. (Sept. 2017). Unveiling the role and life strategies of viruses from the surface to the dark ocean. *Science Advances* **3**(9), e1602565.
- 11. Cornejo-Castillo, FM, AM Cabello, G Salazar, P Sánchez-Baracaldo, G Lima-Mendez, P Hingamp, A Alberti, S Sunagawa, P Bork, C de Vargas, J Raes, C Bowler, P Wincker, JP Zehr, JM Gasol, R Massana, and SG Acinas (Mar. 2016). Cyanobacterial symbionts diverged in the late Cretaceous towards lineage-specific nitrogen fixation factories in single-celled phytoplankton. *Nature Communications* **7**(1).
- 12. Lima-Mendez, G et al. (May 2015). Determinants of community structure in the global plankton interactome. *Science* **348**(6237), 1262073–1262073.
- 13. Sunagawa, S et al. (May 2015). Structure and function of the global ocean microbiome. *Science* **348**(6237), 1261359–1261359.
- 14. Baltar, F, J Palovaara, M Vila-Costa, G Salazar, E Calvo, C Pelejero, C Marrasé, JM Gasol, and J Pinhassi (June 2015). Response of rare, common and abundant bacterioplankton to anthropogenic perturbations in a Mediterranean coastal site. *FEMS Microbiology Ecology* **91**(6).
- 15. Agusti, S, JI González-Gordillo, D Vaqué, M Estrada, MI Cerezo, G Salazar, JM Gasol, and CM Duarte (July 2015). Ubiquitous healthy diatoms in the deep sea confirm deep carbon injection by the biological pump. *Nature Communications* **6**(1).
- 16. Salazar, G, FM Cornejo-Castillo, V Benítez-Barrios, E Fraile-Nuez, XA Álvarez-Salgado, CM Duarte, JM Gasol, and SG Acinas (Aug. 2015). Global diversity and biogeography of deep-sea pelagic prokaryotes. *The ISME Journal* **10**(3), 596–608.
- 17. Ruiz-González, C, G Salazar, R Logares, L Proia, JM Gasol, and S Sabater (Nov. 2015). Weak Coherence in Abundance Patterns Between Bacterial Classes and Their Constituent OTUs Along a Regulated River. *Frontiers in Microbiology* **6**.
- 18. Acinas, SG, I Ferrera, H Sarmento, C Díez-Vives, I Forn, C Ruiz-González, FM Cornejo-Castillo, G Salazar, and JM Gasol (July 2014). Validation of a new catalysed reporter deposition-fluorescence in situ hybridization probe for the accurate quantification of marine B acteroidetes populations. *Environmental Microbiology* **17**(10), 3557–3569.
- 19. Logares, R, S Sunagawa, G Salazar, FM Cornejo-Castillo, I Ferrera, H Sarmento, P Hingamp, H Ogata, C de Vargas, G Lima-Mendez, J Raes, J Poulain, O Jaillon, P Wincker, S Kandels-Lewis, E Karsenti, P Bork, and SG Acinas (Sept. 2013). Metagenomic 16S rDNA Illumina tags are a powerful alternative to amplicon sequencing to explore diversity and structure of microbial communities. *Environmental Microbiology* **16**(9), 2659–2671.
- 20. Rodríguez-Martínez, R, G Rocap, G Salazar, and R Massana (Apr. 2013). Biogeography of the uncultured marine picoeukary-ote MAST-4: temperature-driven distribution patterns. *The ISME Journal* **7**(8), 1531–1543.
- 21. Ferrera, I, CM Borrego, G Salazar, and JM Gasol (Oct. 2013). Marked seasonality of aerobic anoxygenic phototrophic bacteria in the coastal NW Mediterranean Sea as revealed by cell abundance, pigment concentration and pyrosequencing ofpufMgene. *Environmental Microbiology* **16**(9), 2953–2965.
- 22. Salvadó, Z, FN Arroyo-López, JM Guillamón, G Salazar, A Querol, and E Barrio (Feb. 2011). Temperature Adaptation Markedly Determines Evolution within the GenusSaccharomyces. *Applied and Environmental Microbiology* **77**(7), 2292–2302.
- 23. Rojo, C and G Salazar (June 2010). Why are there so many kinds of planktonic consumers? The answer lies in the allometric diet breadth. *Hydrobiologia* **653**(1), 91–102.
- 24. Rojo, C, M Segura, MA Rodrigo, and G Salazar (Sept. 2008). Factors controlling the colonial structure of Pediastrum tetras (Chlorophyceae). *Hydrobiologia* **617**(1), 143–155.