

MaCuMBA



Fifth Issue of MaCuMBA Project News Now Available

November 2013

It has been an exciting first year for the MaCuMBA (Marine Microorganisms: Cultivation Methods for Improving their Biotechnological Applications) project, which held its first General Assembly in Roscoff, France in September. You can catch up with the latest news from this event and the world of marine microorganisms in the fifth MaCuMBA project newsletter (*attached*), which is now available to download from: www.macumbaproject.eu. MaCuMBA is a four-year EC FP7-funded project that aims to uncover the untold diversity of marine microbes using cultivation-dependent strategies.

In this issue of MaCuMBA Project News: an interview with Prof Francisco Rodriguez-Valera from the Universidad Miguel Hernandez (UMH), Spain, who is the leader of MaCuMBA Work Package 6 (WP6), *Sequencing, genomic and metagenomic libraries and (meta)genome analyses*; a report on the first MaCuMBA General Assembly by MaCuMBA coordinator Lucas Stal; information about the upcoming advanced metagenomics course being organised by the Evolutionary Genomics Group (EGG) of UMH; Prof Brian McNeil from Strathclyde University, Scotland, tells us about the EU-funded SeaBioTech project; an in-brief look at some stories relating to marine microorganisms that have been making the news; and profiles of some of the researchers making MaCuMBA happen.

Note to Editors

MaCuMBA is led by the Royal Netherlands Institute for Sea Research (NIOZ) and is a joint venture of 23 partner institutions from 11 EU countries with the common aim to uncover the untold diversity of marine microbes using cultivation-dependent strategies. MaCuMBA aims to improve the isolation rate and growth efficiency of marine microorganisms from conventional and extreme habitats by applying innovative methods and using automated high-throughput procedures. AquaTT is the communication and dissemination partner for the project.

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Detailed partner profiles are available on request.