

Allbio Metagenomics meeting

at the
Netherlands Bioinformatics Conference 2014

Lunteren, April 9th, 2014

<http://www.aanmelder.nl/nbic2014>

The **Netherlands Metagenomics Platform** cordially invites you to join the **Allbio Metagenomics meeting** at the **NBIC 2014**.



The keynote lecture will be presented by **Marcus Claesson** of the University College Cork

Topics include:

All areas of environmental sequencing
(e.g. amplicon sequencing, shotgun metagenomics, metatranscriptomics, metaproteomics)
Prokaryotic, eukaryotic or viral systems
Sequencing strategies and technologies
Quality control, data pre-treatment
Read mapping, taxonomic and functional annotation
Comparative metagenomics
Metagenome assembly
Discovery of novel species, enzymes
Virus discovery by metagenomics
Monitoring, public health surveillance
The tree of life, the rare biosphere
Microbial networks and ecosystem modeling
Host-microbe interactions (e.g. human, plants, corals)
Role of microbes in bioremediation, biogeochemical cycles

Abstracts are solicited in any field related to metagenomics or computational analysis of microbial communities. See the list of example topics above. Abstracts will be peer reviewed. To submit an abstract, please visit the abstract submission page at: <http://www.aanmelder.nl/nbic2014/wiki/metagenomics%20topical%20track>
Please indicate that you are submitting your abstract for the Metagenomics meeting.

