Appendix 5

AMBIO/ANTAR-IMPACT meeting 15.12.08

Report
AMBIO / ANTAR-IMPACT meeting

15th of December 2008

Report

University of Liège, Sart Tilman campus, Liège, Belgium
Context

The AMBIO project started two years ago, officially with the Kick-off meeting which took place at the Belgian Royal Library (Brussels) on the 26-27/03/2007. During the first two years, samples were collected from Antarctica during the MERLIN expedition and from international collaborators. Cultivation and molecular work has begun by the three partners and the first results started to be published (Taton et al. 2006, Zakhia et al. 2007). Besides, many activities related to the International Polar Year programs were done by the partners and the AMBIO website was constructed (www.ambio.ulg.ac.be).

The ANTAR-IMPACT project started in the beginning of 2007 with the BELAR expedition. Dr. D. Ertz collected samples from the region around the Belgian Polar Station “Princess Elisabeth” before its construction. The molecular and morphological characterization already started on these samples. In addition to future samples from the same regions, a comparison will assess the impact of the station on the Antarctic ecosystems.

These two projects are closely linked as they are conducted by almost the same partners and they share the same users and follow-up committee. For this reason a joint meeting was organized on the 15th of December 2008 at the University of Liège, Liège, Belgium. The day was divided into two parts: the morning session with the members’ presentations and the afternoon workshop with general talk from the users.

Morning session: members’ presentations

Welcome and presentation of the meeting: Dr. Annick Wilmotte (C, AMBIO & ANTAR-IMPACT)

The opening session started with Dr Annick Wilmotte, who welcomed the audience, reminding during her talk the context of AMBIO and ANTAR-IMPACT, in the frame of the International Polar Year program MERGE (Microbiological and Ecological Responses to Global Environmental Changes in Polar Regions), and introducing the different partners and participants, in relation with the Workpackages. The talk started with a special emphasise, on the microbial character of the Antarctic continent. The molecular tools allowed a revolution in the exploration of the molecular diversity of both cultured and uncultured microorganisms. The project is framed in the current scientific discussion:

- Is ecology of microorganisms driven by the same factors as eukarya?
- Do endemic microbial taxa exist? (or due to their small size, they can be everywhere?)
- To explain the biogeography of microorganisms

The Antarctic continent is the best place to address these questions: it is a remote place, under extreme conditions. However, it presents a gradient of environmental conditions, from harsh (continental biotopes) to milder (Antarctic Peninsula).

Our aim is to ultimately generate molecular data, which we can deposit in a database and to know better about the communities turn over and learn about the biodiversity patterns.

Then Dr. Wilmotte talked about the objectives, work packages and work to be done in the second phase (starting January 2009). Finally, she presented the ANTAR-IMPACT project that
contributes to the evaluation of the environmental impact of the construction and functioning of the future Belgian Polar Base “Princess Elisabeth” on the Antarctic ecosystem, with its aims and work packages.

**Molecular Diversity of Antarctic Cyanobacteria: Mr. Pedro De Carvalho Maalouf (P1, AMBIO)**

P. De Carvalho Maalouf started by explaining the different techniques used in the AMBIO project in order to study the diversity of cyanobacteria in Antarctica (PCR, DGGE, cloning, sequencing). He then presented the work that has been done previously (F. Zakhia) in the cyanobacteria laboratory on the Sample 41 coming from the Antarctic Peninsula. This was the first report of the diversity in a bi-laminated mat. Some results from the work on West Ongul lakes (East Antarctica) of S. Chalfami were also presented. These, in addition to other samples, were used to construct a Phylogenetic tree. Three new – potentially endemic – OTUs were discovered. A clone library was also constructed with the sample WO4, it showed a rather low diversity (2 OTUs). The presentation ended with the work to be done in the second AMBIO phase.

**First Biological Assessment of Utsteinen with Focus on Lichens: Dr. Damien Ertz (National Botanical Garden, ANTAR-IMPACT)**

During the BELARE expedition in February and January 2008 to the region of the future Belgian Polar Base Princess Elisabeth, Dr. Ertz did a mapping of the Lichens and bryophytes. He talked about the importance of a Petrel population and its impact on the lichen communities by providing them a source of nutrition. After identification of the present organisms, they were differentiated in cosmopolitan, bipolar and endemic species. Two potentially new species were discovered: *Trapelia* sp. and *Lecidella* sp. The ITS sequencing of the *Lecidella* specimens showed that they are all genotypically identical. This campaign also provided 52 samples that are now being analyzed by the different partners for the diversity of bacteria, cyanobacteria, diatoms, green algae, rotifers and tardigrades.

This presentation lead to a discussion on the importance of having an undisturbed monitored site on which studies could be held in parallel on different organisms.

**ANTAR-IMPACT. Diversity of the Surroundings of Princess Elisabeth Antarctic Station: Mr. Rafael Fernandez Carazo (P1, ANTAR-IMPACT)**

R. Fernandez Carazo presented the work that was done on cyanobacteria on the Belgian Polar Base “Princess Elisabeth” samples that were brought by D. Ertz. DGGE with semi-nested PCR was used in order to increase the specificity and have more reliable results. Two methods of DNA extraction were also tested and the Smalla et al. (1993) method was chosen. On the twelve analyzed samples, a relatively low diversity was found compared to other coastal samples. A high degree of endemism and three undiscovered OTUs were found. All samples but one shared at least one OTU, which is in concordance with the theory of distribution of species in near-by
habitats. Finally, R. Fernandez Carazo brought up the importance of the preservation of these sites that hold unique biodiversity against the introduction of alien species.

Exploring Heterophic Bacterial Diversity of Antarctic Samples through Cultivation: Miss Karolien Peeters (P3, AMBIO)

The work on the cultivated bacterial diversity of nine samples coming from different Antarctic regions was presented. The plate counts results as well as the molecular characterization by Rep-PCR and partial sequencing of the 16S rRNA gene of the isolates show a large diversity in the samples (especially for PQ1 sample from Pourquoi-Pas Lake) and between the samples. There are several clusters and separate isolates that have sequence similarities with known taxa below 97% and 95% revealing the presence of probable new species and even new genera. The future work will aim to sequence the complete 16S rRNA gene for at least one representative per cluster and to do a detailed characterization of selected new groups with a view to describing them. (Real-Time) PCR tests will be optimized for fast screening a large number of samples for some specific groups.

Hidden Levels of Phylodiversity in Antarctic Green Algae & Uncultivated Diversity of Green Algae and Bacteria: Influence of Regional vs. Environmental Conditions: Dr. Aaike De Weever (P2, AMBIO)

The first part of the presentation underlined the phylodiversity studies done by P2 on green algae. After the isolation of the strains, their microscopic characterization and ARDRA screening, the 18S rRNA gene of 61 strains was sequenced. They were grouped into 14 taxa and a Phylogenetic tree was constructed, it revealed the long Antarctic isolation of the microorganisms. 18S rDNA results showed that there is a distinct Antarctic green algae flora. As *Chlorella* and *Scenedesmus* are two clades with identical 18S rDNA sequences and are detected in most regions sampled, these will be studied in detail using more detailed markers such as ITS.

The second part of the presentation focused on the selection and analysis of 83 samples coming from 70 lakes for studying the importance of regional vs. environmental factors in shaping green algal, diatom and bacterial community composition. Multivariate analysis of DGGE data showed that the distribution of the samples was mainly explained by the environmental variables, yet a small percentage of the variation was explained by regional variables (although not significantly for bacteria). Finally, the importance of obtaining environmental data for each studied region was underlined, as this is needed for evaluating the importance of regional vs. environmental factors in microbial communities.

Lunch
Afternoon workshop: general talks

Antarctica’s Biological History – Insights from Terrestrial Ecosystems: Dr. Peter Convey (BAS)

Dr. Convey talked about the ice evolution on Antarctica, its geographical isolation and its specific biology. The high level of endemism (100% of the nematode species) as well as the separation of the continent into 10 biological regions was underlined. Finally, Dr. Convey put forth the importance of joining biological data to the glaciology and evolution of Antarctica.

Climate Change Effects on Antarctic Terrestrial Ecosystems: Dr. Ad Huiskes (NIOO-KNAW)

The effect of climate change on Antarctic ecosystems is being studied with the TARANTELLA project. Open Top Chambers were used to mimic a global warming scenario and studies are in course on different vegetation and soil communities as well as arthropods. The talk ended with the importance of comparing the results of the studies (13 countries involved) using different kinds of Open Top Chambers.

Antarctica: White and Wild: For How Long? Alexandre de Lichtervelde (CCAMLR and IWC Commissioner)

The meeting ended with a more political talk by Mr. De Lichtervelde who started with the history of Antarctica and the birth of the Antarctic Treaty System followed by the policy-making and the governments involved. The focus is now on the conservation of Antarctica (climate change, tourism, pollution, exotic species...). The talk ended with the importance of the surrounding countries and the international collaboration on the scientific as well as the politic levels.

Internal discussions

At the end of the day, the three partners met in order to have an internal discussion. Three main subjects were put on the table:

1. **Phase 1:**

   PAE will not make clone libraries on algae but rather DGGE-based study. The resolution of the gels is less than expected and bands at the same height might come from different organisms. Moreover, there is a special difficulty with algae and protists: lack of specific PCR primers.

   For bacterial clone libraries, PAE will agree with LMG on the basis of the cultivated diversity, early 2009.

   LMG will do the 16S rRNA of remaining isolates, which will probably give new taxa. They will think of more specific tests to carry out in larger sets of samples.
ULg will do the DGGE of the samples studied also in Gent, and drop the ones of Gibson (too late) and Borghini (too late). Frederic Zakhia subsampled Syowa and Schirmacher Oasis. The Borghini samples will be used in Phase 2.

2. Phase 2:

PAE is thinking about doing T-RFLP for bacteria, and ITS sequencing (probe?) for Chlorella and Scenedesmus.

LMG will stop the cultivation, will do Real Time Quantitative PCR for specific groups, describe the taxonomy of the new taxa and integrate the results.

ULg will use the same samples as Gent for the cultivation of cyanobacteria, and start the Real Time Quantitative PCR. Cultivation will continue on samples agreed on with the partners from Gent.

New samples to come from the following regions: Belgian Basis (BELDIVA 2009), Macquarie Islands (Dana Bergstrom, Australia), Byers (Bart Van de Vije) and South Georgia (Dominic Hodgson). Macquarie and South Georgia are from Subantarctic Islands, where we have no samples yet.

3. Publication strategy:

PAE:

- DGGE MICROMAT (problems with referees and data on DGGE gels of cyanobacteria, Elie should write to Arnaud Taton and Stana Grubisic to see if they can help),

- Green algae cultures,

- Limnology of Syowa Oasis/Schirmacher Oasis,

- Uncultivated diversity by DGGE (all organisms),

- Clones versus cultures (PAE/LMG).

LMG : samples of Belgian Basis, comparison with clone libraries

ULg:

- Transantarctic Mountains

- Belgian Basis, DGGE + microscopy

Choice of co-authorship: The people who sampled and the ones contributing with additional data (see algorithm of BAS that was given by Dominic Hodgson for MICROMAT).

Finally, the partners should think about the IPY meeting in 2010 in Oslo (ideas, presentations...).
Addendum: Participant list

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<tr>
<th>Name</th>
<th>Institute</th>
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<tr>
<td>Boistos, Sophie</td>
<td>University of Ghent, Belgium</td>
<td>PhD student</td>
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<tr>
<td>Convey, Peter</td>
<td>BAS, UK</td>
<td>Dr., scientific visitor</td>
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<tr>
<td>De Carvalho Maalouf, Pedro</td>
<td>University of Liège, Belgium</td>
<td>PhD student</td>
</tr>
<tr>
<td>de Lichtervelde, Alexandre</td>
<td>Federal Department of the Environment, Belgium</td>
<td>Follow-up committee</td>
</tr>
<tr>
<td>De Wever, Aaike</td>
<td>University of Ghent, Belgium</td>
<td>Dr., Postdoc P</td>
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<tr>
<td>Ertz, Damien</td>
<td>National Botanic Garden of Belgium</td>
<td>Dr., P</td>
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<tr>
<td>Fernandez Carazo, Rafael</td>
<td>University of Liège, Belgium</td>
<td>PhD student</td>
</tr>
<tr>
<td>Huiskies, Ad</td>
<td>NIOO - KNAW, The Netherlands</td>
<td>Dr., scientific visitor</td>
</tr>
<tr>
<td>Mano, Marie-José</td>
<td>University of Liège, Belgium</td>
<td>PhD student</td>
</tr>
<tr>
<td>Moermans, Coraline</td>
<td>National Botanic Garden of Belgium</td>
<td>Technician</td>
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<tr>
<td>Namsaraev, Zorigto</td>
<td>University of Liège, Belgium</td>
<td>Dr., Postdoc P</td>
</tr>
<tr>
<td>Peeters, Karolein</td>
<td>University of Ghent, Belgium</td>
<td>PhD student</td>
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<tr>
<td>Savichtcheva, Olga</td>
<td>University of Liège, Belgium</td>
<td>Dr., Postdoc P</td>
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<td>Simon, Patricia</td>
<td>University of Liège, Belgium</td>
<td>Technician</td>
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<tr>
<td>Van Isaker, Nathalie</td>
<td>IPF, Belgium</td>
<td>User</td>
</tr>
<tr>
<td>Vancauwengerghe, Maaike</td>
<td>BELSPO, Belgium</td>
<td>Manager</td>
</tr>
<tr>
<td>Vereecke, Claudine</td>
<td>University of Ghent, Belgium</td>
<td>BCCM manager</td>
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<tr>
<td>Verleyen, Elie</td>
<td>University of Ghent, Belgium</td>
<td>Dr., P</td>
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<tr>
<td>Vyverman, Wim</td>
<td>University of Ghent, Belgium</td>
<td>Prof., P</td>
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<tr>
<td>Wilmotte, Annick</td>
<td>University of Liège, Belgium</td>
<td>Dr., C</td>
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P: partner
C: coordinator
BCCM: Belgian Collection of Microorganisms
IPF: International Polar Foundation
BELSPO: Belgian Federal Science Policy

N.B. the following persons apologized for their absence: Dr. Gibson, Dr. Pearce, Prof. Naganuma, Prof. Marinelli, Prof. Vincent, Prof. Quesada, Dr. Chapelle, Dr. Hodgson, Dr. Danis, Dr. Bosschaerts, Dr. Sergers and Dr. Sabbe.

Annex: program meeting
AMBIO / ANTAR-IMPACT meeting

15th of December 2008, University of Liège, Sart Tilman campus, Liège, Belgium

Morning session: members’ presentations - room 2.71, building B6c

09h50 Welcome and presentation of the meeting
Dr. Annick Wilmotte, Cyanobacteria group, University of Liège, Belgium

10h00 Molecular Diversity of Antarctic Cyanobacteria
Mr. Pedro De Carvalho Maalouf, Cyanobacteria group, University of Liège, Belgium

10h20 First Biological Assessment of Utsteinen with a Focus on the Lichens
Dr. Damien Ertz, National Botanic Garden of Belgium, Brussels, Belgium

10h30 ANTAR-IMPACT. Diversity in the Surroundings of Princess Elizabeth Antarctic Station
Mr. Rafael Fernandez Carazo, Cyanobacteria group, University of Liège, Belgium

10h50 Exploring Heterotrophic Bacterial Diversity of Antarctic Samples Through Cultivation
Miss Karolien Peeters, Laboratory of Microbiology, University of Ghent, Belgium

11h20 Uncultivated Diversity of Green Algae and Bacteria: Influence of Regional vs. Environmental Factors
Dr. Aaike De Weever, Laboratory of Protistology and Aquatic Ecology, University of Ghent, Belgium

12h00 Discussion, questions and suggestions

12h30 Lunch (sandwiches and beverages)

Afternoon workshop - amphitheatre A.4, building B7b

14h00 Antarctica’s Biological History – Insights from Terrestrial Ecosystems
Dr. Peter Convey, British Antarctic Survey, Cambridge, United Kingdom

14h45 Climate Change Effects on Antarctic Terrestrial Ecosystems
Dr. Ad Huiskes, Unit for Polar Ecology, Netherlands Institute of Ecology, Yerseke, The Netherlands

15h30 Antarctica: White and Wild: How long for?
Mr. Alexandre de Lichtervelde, Federal Public Service Health, Food Chain security and Environment - National Contact point for the Committee for Environmental Protection of the Antarctic Treaty - CCAMLR and IWC Commissioner