

# FORBIO Minutes of the 2<sup>nd</sup> follow up committee.

## Present

W. Buysse, G. Geudens, A. Vanderwerf, M. Vanheuckelom, Q. Ponette, K. Verheyen, K. Ceunen (report)

## Excused or absent

Pro silva Wallonie, FEDEMAR, Forêt Wallone, Bosgroepen, BIM, DNF, Natagora, Natuurpunt, Landelijk Vlaanderen

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## See powerpoint for more info on contents of meeting

- Issues that have been discussed:
- Project overview WP1
  - Questionnaire
    - Suggestion to publish an article of the results of the questionnaire (for example in Natuurpunt studie, pro silva, Forêt Wallone,...)
    - From a first analysis of the questionnaire, we can see that the answers for Flanders and Wallonia are similar.
    - We should take into account the differences between the respondents in Flanders and Wallonia (for example Bosgroepen esp. the coordinators and in Wallonia esp. the private forest owners that answered the questionnaire)
    - Another thing that is remarkable is that already 80% of the people that answered the questionnaire is planning to realise mixed forests and 70% is already realising mixed stands.
    - *PDF with summary of results will be distributed*
  - White paper
    - It is more a synthesis of the knowledge that already exists, also for a larger public. We do not aim at a publication for an international, peer-reviewed journal
    - Also for the stakeholders it should be an interesting paper
    - It will be launched at the conferences, scheduled in March '10 and can be placed on the website.
- Task 1.4 Website
  - Etienne will make link to the Belgian Biodiversity Platform and will check whether the site is frequently visited
  - Maybe we should add a remark that persons who want a reprint of the papers listed on the site can contact us
- Task 2.1 Experimental design
  - Management: (selective) cuttings are possible

- A question was raised whether one replication of species combination is enough to be able to compare plots with a higher genetic diversity and lower genetic diversity -> *we think this should be fine, if you consider your species diversity treatments as a block factor*