## <u>Minutes of the 5<sup>th</sup> (and Final) RECIPE progress meeting</u> held on Sunday 28<sup>th</sup> – Wednesday 31<sup>th</sup> of May 2006 at the Macaulay Institute, Aberdeen, Scotland

Present:

Steve Chapman (MLURI) (Project Co-ordinator and Chair)

Harri Vasander (UNHEL); Mika Yli-Petäys (UNHEL); Alexandre Buttler (UFC-CE, AR-WSL); Walter Rosselli (AR-WSL); Andy Siegenthaler (AR-WSL); Edward Mitchell (EPFL) ; Thierry Jean Heger (EPFL); Viviane Froidevaux (LINECO); Fatima Laggoun-Defarge (ISTO); Jean-Robert Disnar (ISTO); Laure Comont (ISTO); Sebastien Gogo (ISTO); André-Jean Francez (ECOBIO); Daniel Gilbert (UFC-LBE); Antonis Chatzinotas (UFZ); Michael Schloter (TUM-BO); Andreas Gattinger (TUM-BO); Brigitte Hai (TUM-BO); Daniel Epron (UHP EEF); Rebekka Artz (MLURI); Gerald Schwarz (MLURI); Clare Trinder (MLURI).

Date	Description	Action
Sunday (28 <sup>th</sup> )	Arrival at Aberdeen 15:10–21:50	
Monday (29 <sup>th</sup> )	Technical reports, giving RECIPE outcomes, specifically relating to deliverables (09:00–13:00):	
	<b>WP 01</b> (Steve) This was completed and the socio-economic publication was being developed following on from the meeting in Frick.	
	<b>WP 03</b> Alexandre presented a spreadsheet of the results showing the samples analysed, accumulated data and highlighting the gaps (see Inventorf work.xls). Some discussion was held on how to deal with the gaps or if some could still be filled.	
	Thierry and Edward reported on progress in characterizing the molecular phylogeny of Arcellinida (see Arcellinida_Aberdeen_Thierry.ppt).	
	Antonis gave a presentation on the genetic diversity of protistan groups using 18S rRNA-based methods. Ther was still a lot of development work to be done before application (see aberdeen 2006_Antonis.ppt).	
	Rebekka presented on the fungal communities work. For Workprogram 1, one paper had been submitted and another paper was in preparation but Mantel tests had yet to be done. There had been problems in Workprogram 2 with samples not extracting due either to humic intereference and/or the low biomass (see MLURI_progress_Aberdeenfinal_WP03.ppt).	
	<b>WP 04</b> Michael gave a review of the deliverables and milestones	
	Brigitte gave a presentation on bacterial diversity determinations using t-RFLP, showing the influence of site, vegetation and regeneration stage. Site grouped very well, vegetation separated grass and moss (Finland), while sampling depth was more important than regeneration stage (Le Russey and Chaux d'Abel). Fragment analysis demonstrated the presence of Firmicutes (mainly Gram+) though nothing can be deduced about their function. The question was raised on how to deal with sites as these were essentially pseudoreplicates. (see RECIPE 290506_Brigette.ppt or RECIPEII 290506_Brigette2.pdf).	
	Lunch 13:00–13:30	
	Continuation of reports 13:30–19:00	

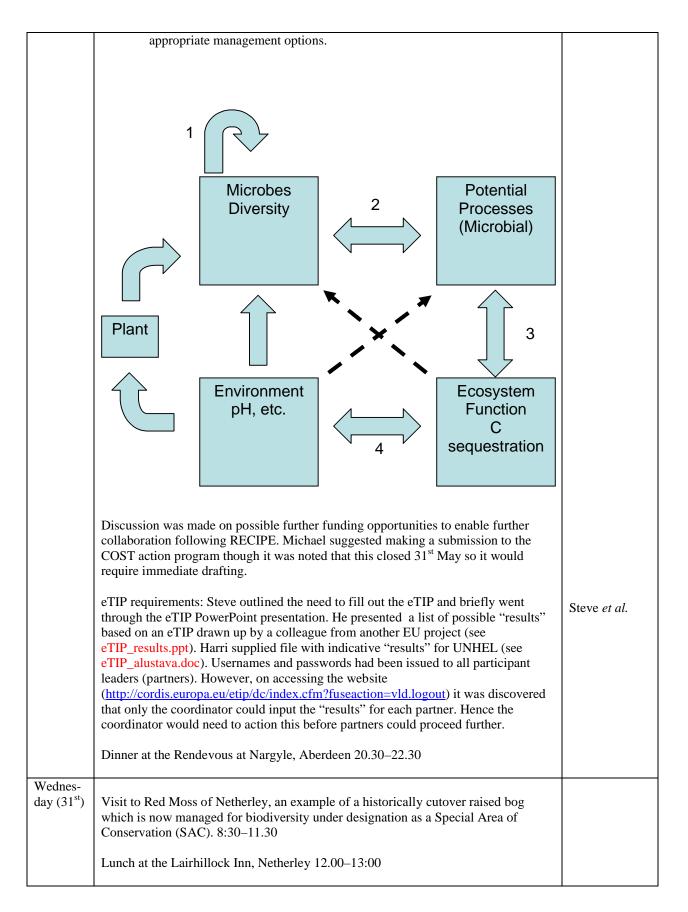
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	Rebekka summarised the CLPP results from both Workprograms 1 and 2. Carbon substrate utilization patterns for the former varied with site (location) and 'humification index' (based on FTIR bands). Together they explained over a third of the variance. Some trends based on regeneration stage were observed but they were not consistent across sites. For Workprogram 2, the dominant variable was site, vegetation had some influence on surface samples while water-table had no effect (see MLURI_progress_finalAberdeen_WP04.ppt).	
	<b>WP 05</b> Fatima and Sebastienne gave an extended report on the results of micromorphology, the C:N ratio of peat fractions and sugar analysis. Significant changes with peat age/depth were seen, as well as between sites, particularly Baupte. Microremains analysis proved to be a useful approach (see ISTO Aberdeen.ppt).	
	Rebekka continued with some data on FTIR analysis. Though country again had a major effect some change with regeneration stage was noted (see MLURI_progress_finalAberdeen_WP05.ppt).	
	<b>WP 06</b> André-Jean reported on carbon turnover studies. For Microbial biomass C or N there were significant responses with plant community and regeneration age. Ratios such as Carbon Turnover also show along the gradient of regeneration stages but CH4/CO2 ratios (potential activity) were not enough sensitive as a regeneration index (see WP6 Aberdeen 28 May-1 June_Andre_Jean.ppt).	
	Andy presented data on gaseous fluxes and estimates of methanotrophs at Le Russey in Workpackages 2 and 3. Peat depth had a major influence, followed by species and water table. There tended to be a reciprocal relationship between carbon dioxide and methane (see Final_Siegenthaler.ppt).	
	Buffer dinner 19:00–22:00	
Tuesday (30 <sup>th</sup> )	Technical reports continued 08:30–12:30	
	Andreas gave a detailed account of the PLEL/PLFA analysis for methanogens/methanotrophs. Methanogenic populations varied across countries and generally increased with depth. Methanotrophs varied even more between countries; they tended to decrease with depth though there was a reverse trend at Le Russey. Using advanced technology, the <sup>13</sup> C signal from litter residues was traced into Gram+ and Gram– bacteria, euryarcheaota, fungi and protozoa (see Aberdeen- May06_gattinger.ppt).	
	<b>WP 02</b> Edward presented on behalf of Emanuela the results of her studies on vegetation and C cycling at Chaux d'Abel. Sphagnum increased along the regeneration gradient while graminoids decreased. Gross photosynthesis and NEE were greatest in the intermediate stage. Methane emissions were greatest for the advanced stage (see Vegetation&Cfluxes_Chaux_d'Abel_Ema.ppt).	
	Mika reported on C flux studies in Finland. In the advanced regeneration stages, C balance was close to zero though GWP (global warming potential) may increase. Good vascular plant growth was achieved in Workprogram 2. For the $CO_2$ models, 90% of the variance could be explained by PAR and LAI. Water table depth was not significant. Methane emission depended upon LAI but not for bare peat and Sphagnum-dominated areas. (need presentation)	
	Rebekka gave an account of the Scottish dataset. This was incomplete and lack of	

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flux in Workprogram 1 and air or soil temperature. In Workprogram 2, CH <sub>4</sub> flux was variable but tended to be greatest where <i>Eriophorum</i> sp. was present (see MLURI_progress_finalAberdeen_WP02.ppt).	
Daniel Epron presented (on behalf of Estelle) on C balance at Le Russey. Dark respiration was satisfactorily modelled by a combination of air temperature, water table depth and dessication index. Photosynthesis under saturating conditions was modelled by air temperature and dessication index. Modelled carbon balance was negative for the bare site but positive for the recent and advanced plots. $CH_4$ emission was related to the LAI of vascular plants (see Prés. 0605 Aberdeen_DanielEpron.ppt).	
<b>WP07</b> Daniel Gilbert reported briefly on the socio-economic work for France carried out by Carine and Amardine. Interviews had been made by phone with producers. The number of exploitations had dropped to 16 (from 23 in 1995).	
Gerald had unfortunately fallen sick and was unable to give his report on WP07.	
A meeting was planned in Franche-Comté for September/October 2007. Philippe Grosvernier would preside over a session on RECIPE (co-organised by Line Rochefort).	
<i>Frontiers in Ecology and Environment</i> was suggested as a possible journal for an article on management options. An alternative was <i>Gaia Ecological Perspectives for Science</i> .	
All PowerPoint presentations should be passed on to Rebekka who will compile these and include them on the <u>RECIPE website</u> .	Presenters/ Rebekka
Lunch 13:15–14:00	
Discussion session on outstanding deliverables/milestones.	
Rebekka reported on the status of the Workprogram 1 database (M10, M16) (See WPI_Environmental_Data.xls). A lot of data had been added but ther were still gaps.	Rebekka/ Everyone
D16 was still in progress; <sup>15</sup> N results were expected by the end of June.	Fatima
D20: some Keeling plots were still to be done but should be completed within two weeks.	André-Jean
D9 Much of the work here was in the development of tools which would then be applied to one site (in CH). Will take at least until the end of July to complete this but relies on Antonis finding time. Looked into possibility of using some frozen samples to redo DNA extractions where data for testate amoebae missing.	Antonis/ Edward
D18: some amino acid analysis still pending.	Sebastien/ Fatima
D11: where indicators are specifically protistan, this has to be worked on. Edward will perform regressions with C turnover.	Edward
D23 This has yet to be fully evaluated; some indicators of change have been shown but this needs a fuller analysis.	Steve <i>et al</i> .
D22 In terms of guidelines for sustainable development, it was felt that to some	Steve

socio-economic paper (see Tabl	onomic issues were tackled by the forthcoming e 1 developed in Frick). Social aspects were still perhaps to review the guidelines from elsewhere and	
website "A Global Peatland Res	ation was available for comment on the IMCG toration Manual" by Martin Schumann and Hans ocess and passed it on to Edward for further review	Edward
management strategies which ha	conomic benefits requires the derivation of specific ave not yet been identified from D22 and D23. This ugh to the socio-economic group for consideration.	Steve
publications. After an intermina Global Change Biology to the e being of general interest to GCE Approaches had been made to J awaited. It was suggested that in be investigated. Possibilities we <i>Ecosystems</i> <i>Oecologia</i> <i>Soil Biology &amp; Biocher</i> <i>Ecography</i>	nistry	Steve
Other possibilities ( <i>Restoration</i> <i>Science</i> ) returned rather low imp	Ecology, Wetlands, European Journal of Soil pact factors.	
itemised for the special publicat draw together the different stram the longer term (6–12 months) of	o consider possible publications beyond those ion. These would seek to be synthesis papers which ds of the RECIPE project. Publication would be in once other papers had reached acceptance. It was levels of synthesis that could lead to four papers (see	
groups specified from of methods (e.g. clone and whether they repre	versity. This would cover bacteria, fungi, protists, lipid analysis. It would look at the complementarity libraries, FISH, DAPI), their relative discrimination, sented genotype or phenotype. There were perhaps filled (e.g. testate amoebae molecular genetics for analysis (D. Gilbert?)).	Michael/ Antonis
results, CLPP data and function". Marker lipid structure and function	ucture and function. This would cover 13C-PLFA soil respiration results, in other words, "organismic ls enabled a direct link possible but relations between for other groupings might be by correlations only , CANOCO, Mantel tests).	Rebekka/ Andreas
<ol> <li>This would relate "orga address the question: D we assess or quantify b How do we relate poten vitro to CH<sub>4</sub> emission i would consider C flux environmental variable</li> </ol>	anismic function" to ecosystem function. It would bo we need biodiversity to restore function? How do iodiversity (Shannon-Wiener index, evenness)? ntial function to actual function, e.g. CH <sub>4</sub> emission in n vivo, or methanotrophs to methane emission? It but would also need to take account of s/vegetation/chemistry of the substrate. Some gaps	Alexandre/ André-Jean
4. This level would cover	nd chemistry that might be useful here were noted. vegetation and modelling with the aim to make certain scenarios. It would then come up with	Steve/ Alexandre



	Visit to Glen Garioch Distillery, Oldmeldrum 14:00–16:00	
	During the course of the day, while traveling, over lunch and before the evening meal, a COST action submission was drafted (largely by Alexandre) and submitted (in Steve's name). The title was "Network for European sustainable peatland management (NETPEAT)" and the full submission is given in PEATNET.pdf.	
	Dinner at Ardoe House Hotel 19:30–22:00	
Thursday (1 <sup>st</sup> )	Participants disperse	