



EVOLTREE

EVOLution of TREEs as drivers of terrestrial biodiversity

Linking genomics and ecology to understand the evolution of diversity in terrestrial ecosystems

<http://www.evoltree.eu/>





EVOLTREE

A Network of Excellence under the
EC 6th framework programme for research

Focus: Assessing and forecasting changes in
biodiversity, structure, function and dynamics
of ecosystems and their services



EVOLTREE



A Network of Excellence under the
EC 6th framework programme for research

Coordinator: Antoine Kremer (INRA)

Consortium of 25 institutions in 15 countries

EC contribution: 14.3 million euros

Starting date: 1 April 2006 (end Sept 2010)

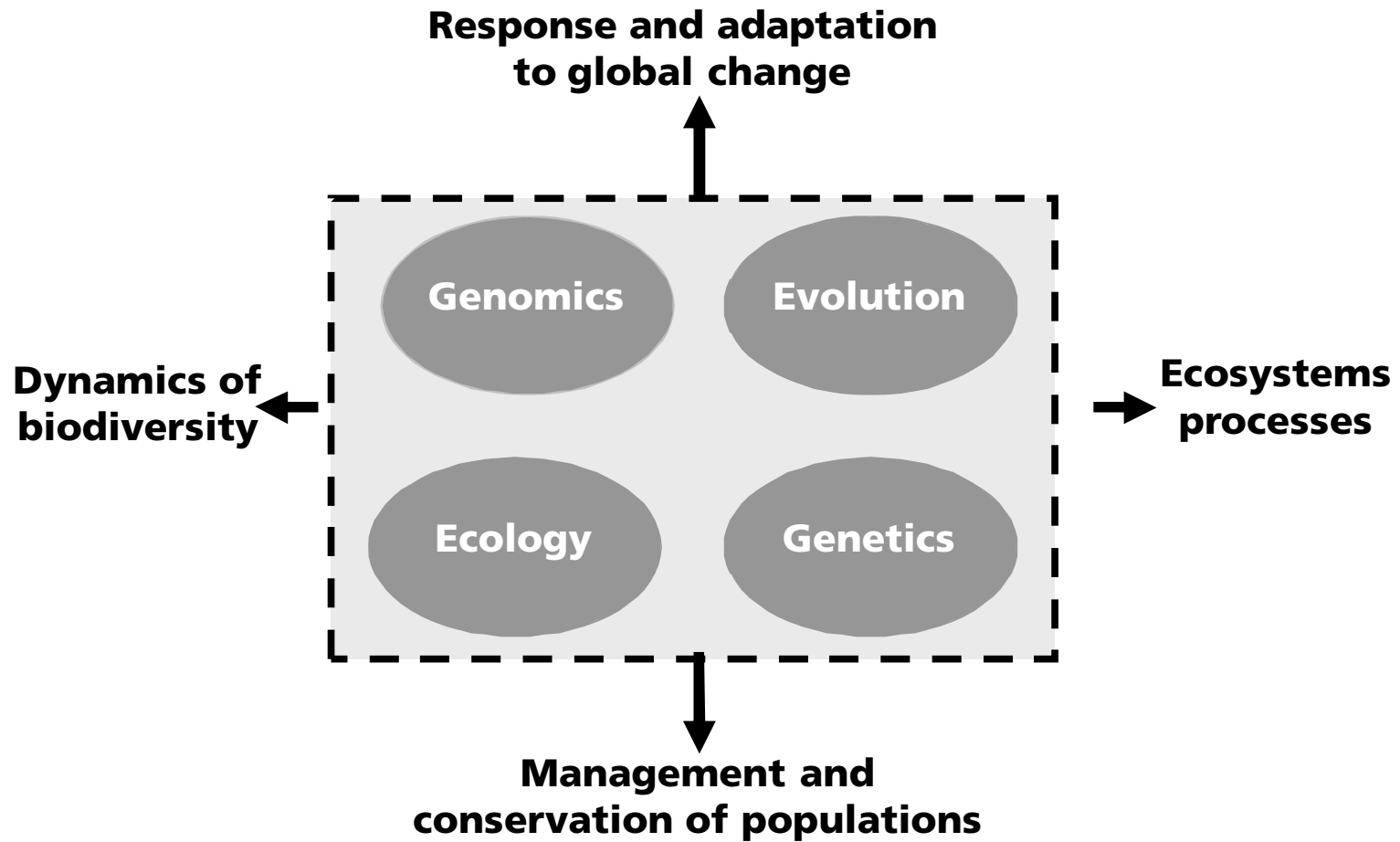
Duration: 4 years (+ 6 month no-cost extension)



© INRA



EVOLTREE: disciplines



The Repository Centre



- **Central material & data access point**
- **Automated management and supply of plant-genes & gDNA**
- **centralized and standardized resources**
- **Online access to distributed information**
- **embedded in an European research network (NoE)**

13.09.10, Vienna

Forest 'gene resources'



- Material available
 - Leaf tissue (37.000)
 - gDNA (21.000)
 - BACs (60.000)
 - ESTs (400.000)
- From Gene Banks & Intensive Study Sites (ISS)
- Data base, LIMS, web portal
- Objective: Distribution of forest 'gene resources' for ecosystem research

13.09.10, Vienna

ISS-material sampled



	Blizyn	Landes	Loire	Punkaharju	Solling	Valais	Ventoux	Koralm	Roudsea	Veluwe	Total
<i>Abies alba</i>	500						1800				2300
<i>Abies, Fagus, Pinus</i>							200				200
<i>Alnus</i>	50										50
<i>Betula</i>	50										50
<i>Fagus sylvatica</i>	200				629		1400			150	2379
<i>Fagus, Pinus, Quercus</i>							190				190
<i>Larix</i>	50										50
<i>Picea abies</i>	50			500	200			720			1470
<i>Pinus cembra</i>								720			720
<i>Pinus mugo</i>								720			720
<i>Pinus sylvestris</i>	500			770		800					2070
<i>Populus nigra</i>			400								400
<i>Quercus sp.</i>	100	1500		50	500	800	500		500		3950
Total	1500	1500	400	1320	1329	1600	4090	2160	500	150	14549

Gene Bank Network

28 mapping populations (9500 genotypes)

★ Centres of maintenance (referenced Jan2008)
(Pedigrees + Association populations)

Fagaceae :

- 4 *Quercus*,
- 1 *Fagus*,
- 1 *Castanea*

Pinaceae: 5 *P. pinaster*,

- 2 *P. sylvestris*,
- 5 *Picea abies*,
- 2 *Larix*,
- 1 *Cedrus*

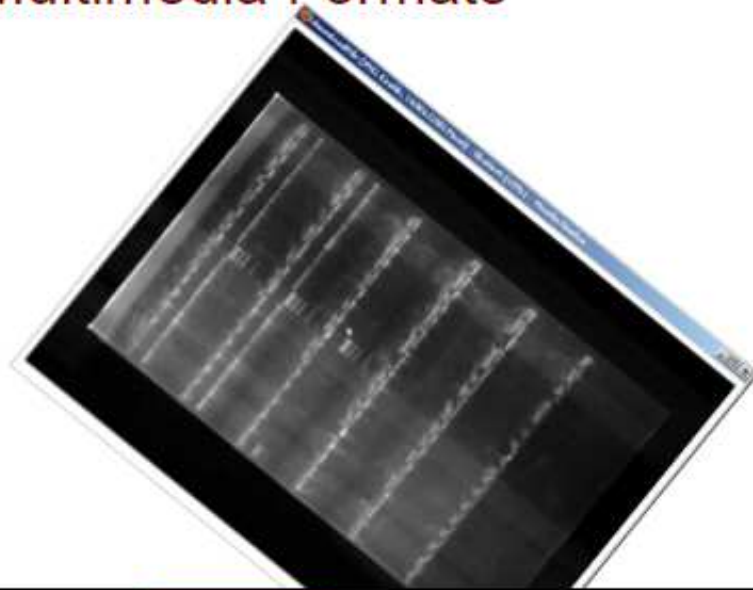
Salicaceae:

- 6 *Populus sp.*
- 2 *Salix*



9 within Evoltree, 10 outside

Documentation DBs: Georeferenced Data Multimedia Formats



Well Parameter Material Parameter

Material Default Collection

Name	ISS-PU-A770
Kommentar	
Externe ID	
Beweis	gDNA
Spezies	Scots pine (<i>Pinus sylvestris</i>)
Erstellungsdatum	2009-12-02 00:00:00
Gruppierungen	

Delen Drucken Ansicht Chronik Leseschichten Extras Hilfe

MAPS

1 2 3 4 5 6 7 8 9 10 11 12

A B C D E F G H

Container Default Gels

gDNA Gel

Actin PCR Gel

Well Parameter Material Parameter

Material Default Collection

Developmental Stage: adult

Sampling Date: 25.05.2010

Accuracy	0.0
Longitude	29.305
Latitude	61.8075

Karte Satellit Hybrid

Sampling Location

Name Sampler: Taina Salminen

e-mail Sampler: taina.salminen@metla.fi

Pedigree

ISS: Pankahägi

Tree ID: A770

Material at Provider:

Additional Information

13.09.10, Vienna

Integration: Distributed data sets & data access

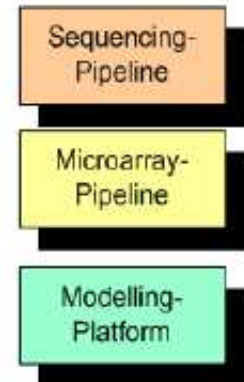
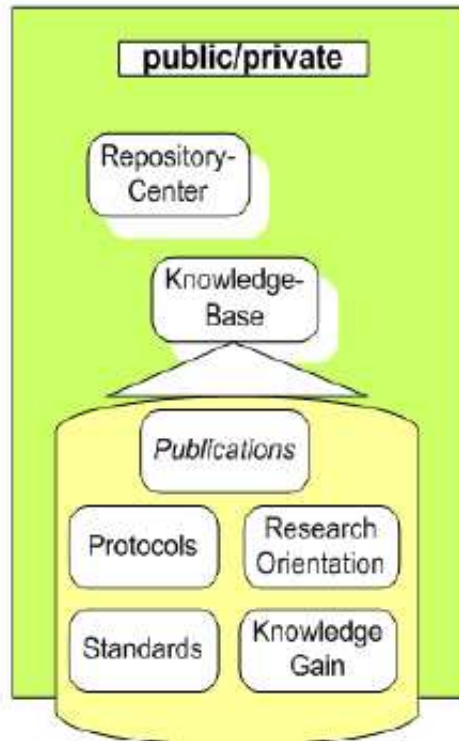
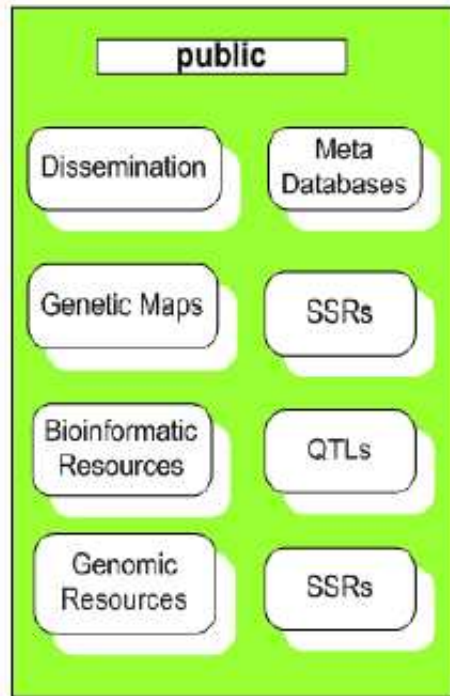


13.09.10, Vienna

www.evoltree.eu
www.picme.at
www.ait.ac.at



EvoTREE Portal: Access to data via web-site



Jointly executed research activities



- Identify genes of adaptive significance in regards to global change in three model species of trees (*Pinus*, *Populus* and *Quercus*), phytophagous insects (*Limantria*) and mycorrhizal fungi (*Laccaria* and *Glomus*)
- Assess the level and distribution of nucleotide diversity in genes of adaptive significance in trees, insects and mycorrhizal fungi
- Assess the impact of trees on the composition of communities by studying interactions between trees and their associated species
- Investigate the evolutionary processes in trees by reconstructing their past history and predicting their future response to global change



Dissemination

Results presented at an international scientific conference on "Forest Ecosystem Genomics and Adaptation" (Spain, June 2010), organized by EVOLTREE

Establishment of an EVOLTREE Stakeholder Group (science-policy interface)

Science-based contributions to policy processes (MCPFE, PEBLDS)

Public awareness campaign and products

Evoltree continuation

Network will be carried as a new research programme within the European Forestry Institute (EFI)

The following networking activities will be maintained and supported by either cash or in kind contributions of the EVOLTREE partners:

1. maintenance and upgrading of EVOLTREE Portal and databases,
2. maintenance and access to common infrastructures and resources established,
3. dissemination (newsletters and others dissemination products),
4. training (organization of 4 to 6 summer schools or workshops per year),
5. coordination of activities



Evoltree continuation



Different levels of membership have been defined, based on a governance model to allow for the different levels of financial and in-kind

Research activities per se will be continued only if funds for collaborative projects are available (either FP7 or FP8 projects, ERA-NETs or other international funding opportunities)





<http://www.evoltree.eu/>

