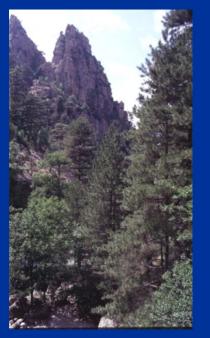


COMMISSION DES RESSOURCES GENETIQUES FORESTIERES Bruno Fady, INRA – URFM, Ecologie des Forêts Méditerranéennes Avignon, France bruno.fady@avignon.inra.fr

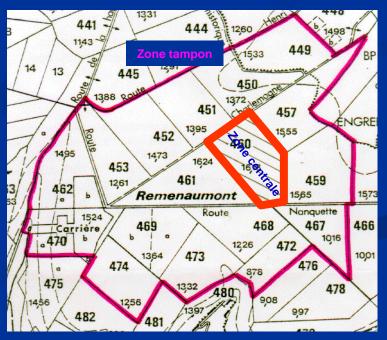


Conservation of genetic resources: the European strategy and a case study





Genomics and the conservation of conifer genetic resources



ProCoGen dissemination workshop – Szombathely, Hungary, 1-3 September 2014

The goals of in situ conservation

A strategy for safeguarding keystone or emblematic species against natural and made-made ecological catastrophies

=> Networks must contain all the genetic diversity of a species within its entire distribution range

=> Conservation units must be made to garantee local adaptation under diverse selection pressures



How to correctly sample the genetic diversity of a species? Considering evolutionary history

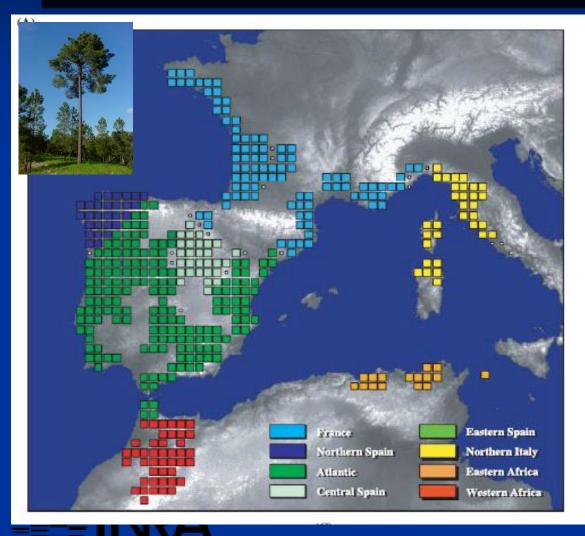
ESU : Evolutionary Significant Unit (Moritz 1994)

- = group of populations deriving from a common ancestor(lineage) and significantly different from other lineages within the species
- = signature of long term evolutionary history (mt/cpDNA).

MU : Management Unit (Palsbøll et al 2007)

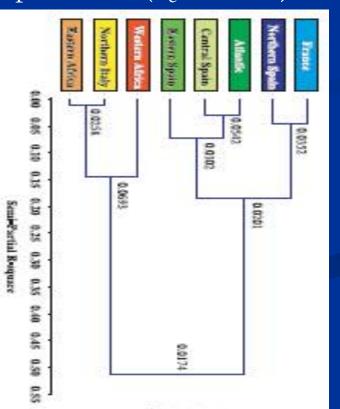
= group of populations that differs from another by significant differences in genetic markers (reduced gene flow)
= signature of short term evolutionary history (nDNA ou SSRs)

Considering evolutionary history (demography): an example of data availability in Pinus pinaster



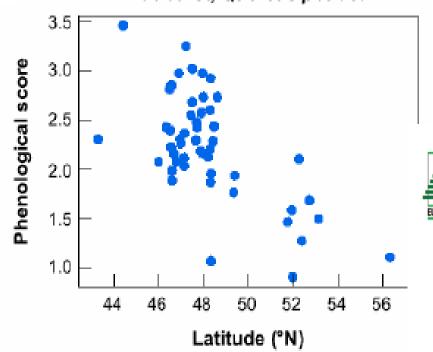
Bucci et al. (Mol. Ecol.) 2007

3 lineages and 8 genetic groups from 16 (most common) haplotypes at 5 cpSSR loci ($h_e = 0.825$).



Populations have different adaptive properties in addition to different evolutionary histories

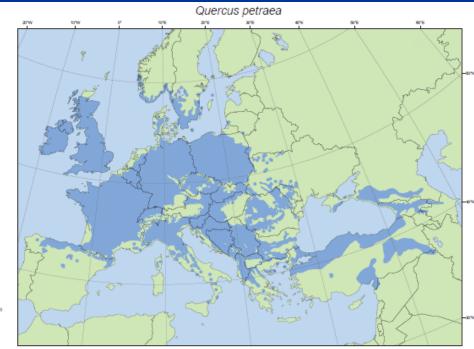
Budburst, Quercus petraea



Ducousso et al. (AFS) 1996

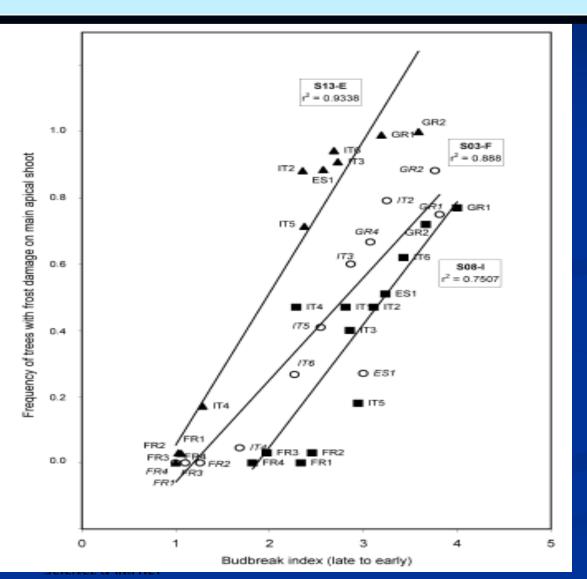


RUFCROEN Secondate do Risovarily International Via del The Denard, 4721a 00057 Macazenee (Funktion Rome, tray Tai, (-abgottent/Rufer et __secondaria@gotten.org Mose Information and other maps at: where antipages.org A strong link between geographic origin and bud break date in the European oak *Quercus petraea* (4 common garden experiments)



This distibution map, showing the natural distribution area of Quercus petrace was compiled by members of the EUFORGEN Networks Citation: Distribution map of Sesalle cek (Quercus petrace) EUFORGEN 2009, www.eutorgen.org.

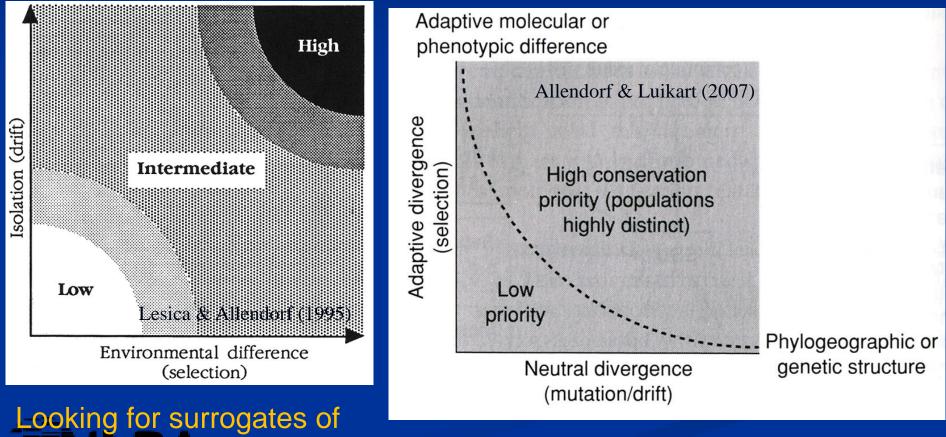
Considering adaptive properties in the evolutionary history of populations and species



A strong link between geographic origin, bud break date and susceptibility to late frost damage in walnut (*Juglans regia*) in 3 common gardens

Fady et al. (NeFo) 2003

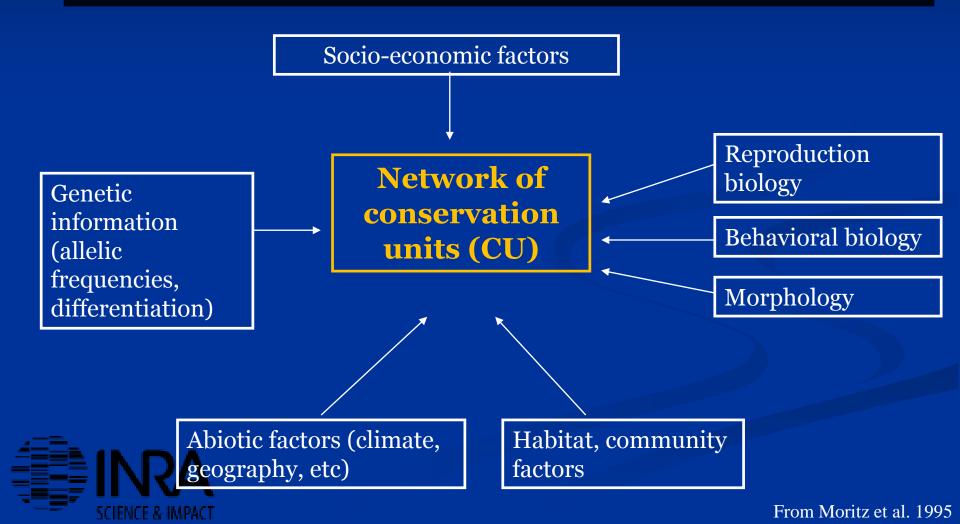
How to correctly sample the genetic diversity of a species? Considering adaptation in addition to demography and evolutionary history



adaptation: environmental Lookir gradients: & IMPACT pheno

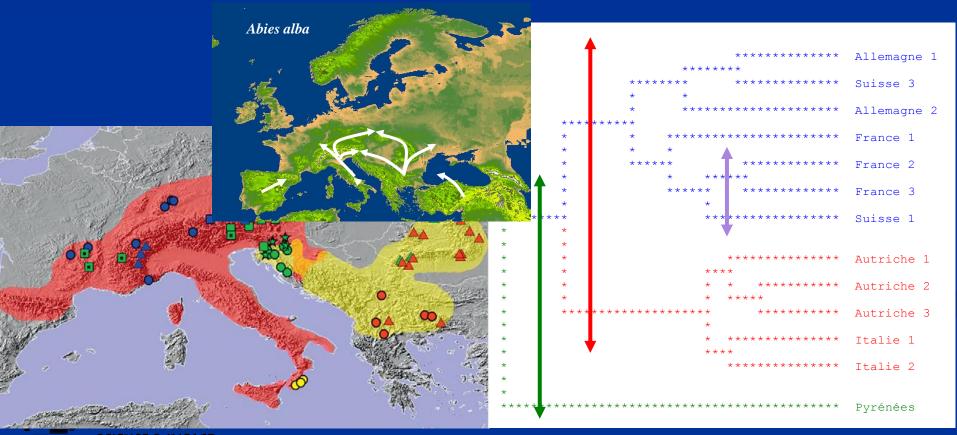
Looking for surrogates of adaptation: phenotypic / genotypic trait variability

Integrating approaches for a science-based sustainable in situ conservation strategy



One example of genetic resource conservation network in France: Abies alba

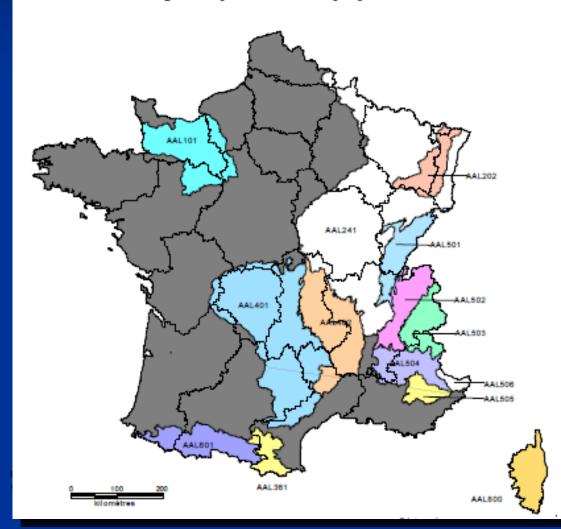
An ancient common origin for western lineages At least 2 Quaternary western lineages (Pyrenees + Alps) => 2 ESUs At least 2 genetic clusters within the Alps => 2 MUs



Fady et al. (Forest Genetics) 1999. Liepelt et al. (Rev. PaleoBot. Palynol.) 2009

One example of genetic resource conservation network in France: Abies alba

Régions de provenance du Sapin pectiné



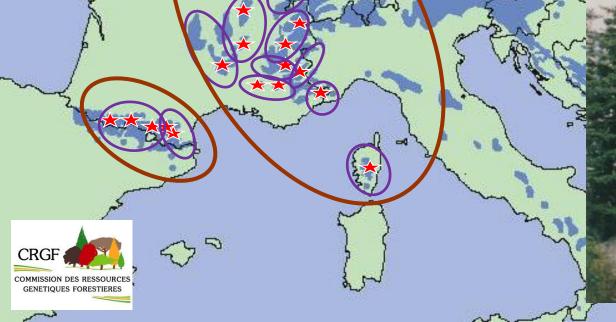
Little analytical work from provenance trials available / no data from genomic tools yet

14 regions of provenance : an estimator of ecological structuration in France

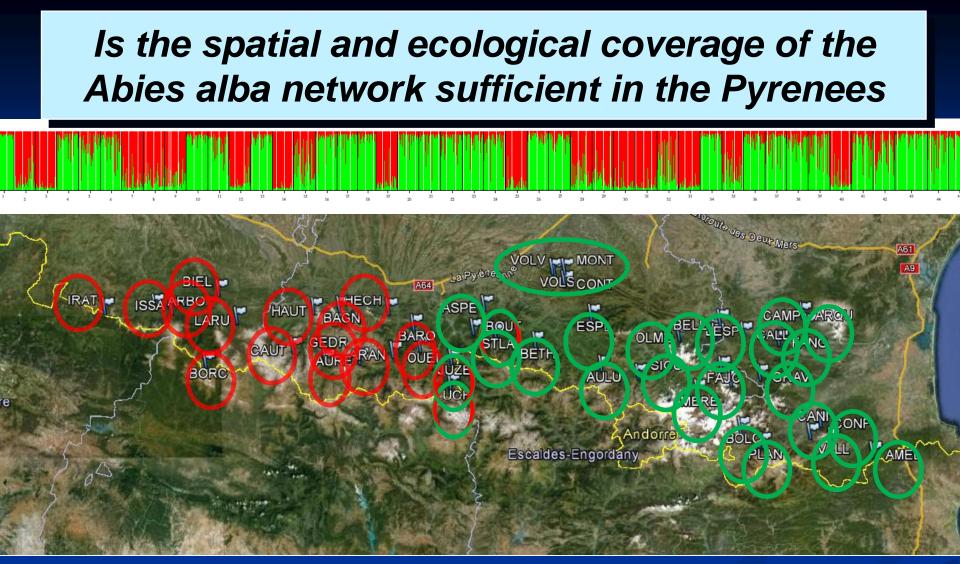
=> At least 14 CUs (emphasizing local adaptation) in France

The actual and current network of FGR conservation of Abies alba in France

In practice, a combined approach: ESU + MU + ecological structure + marginal populations = 21 CUs officially registered

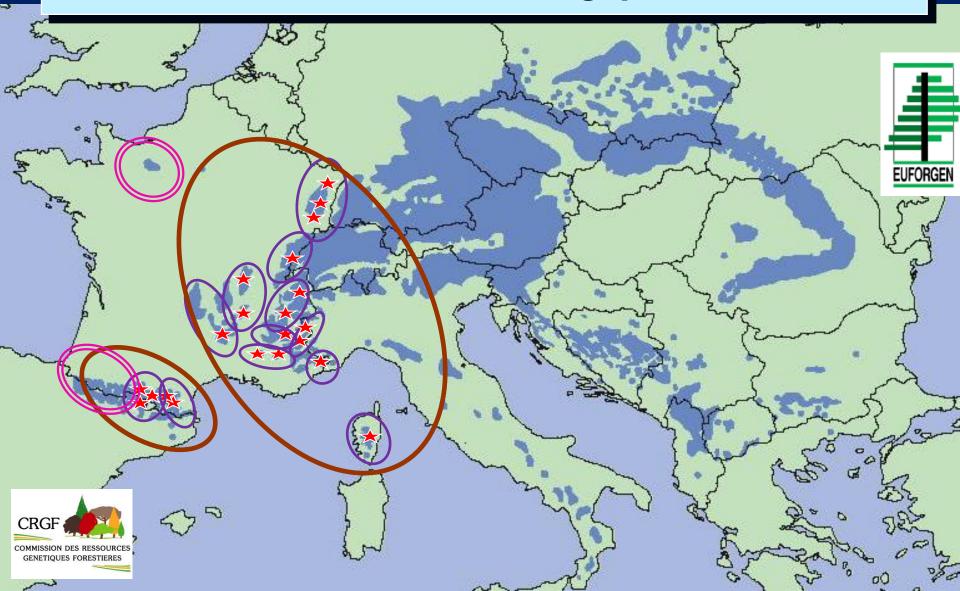


CU of Beaumont de Ventoux (84) – a marginal population

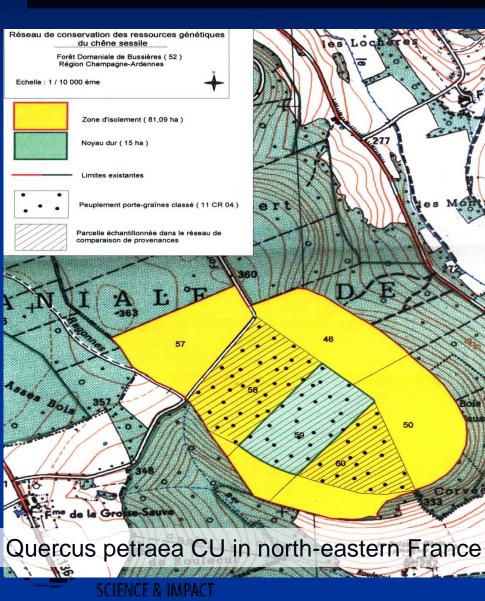


A significant geographic structure, 2 evolutionary lineages (10 SSR loci), no effect of marginality (VOLV)
A clear transition admixed zone at lineage boundary
Detection of a planted population (ASPE)

The current network of FGR conservation of Abies alba in France: gaps to fill!



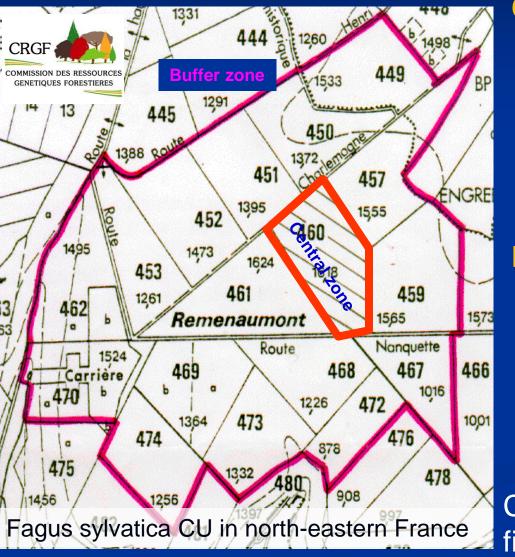
What must a Conservation Unit guarantee?



Ongoing local adaptation under natural selection must be maintained

Management must: ==> maintain reproduction and seedling recruitment; ==> maintain high adult density to avoid drift / inbreeding during reproduction; ==> prevent unwanted gene flow. A need for monitoring

How must a Conservation Unit be managed? A legally binding charter



Central zone

- > Autochthonous forest
- > 500 seed trees minimum
- > 60 seed trees/ ha
- Natural regeneration only (potentially assisted using local seeds)

Buffer zone

> No introduction of hybridogenous exotic species / populations
> Regeneration after the central zone

Control of game species, wild fire protection, monitoring, etc.

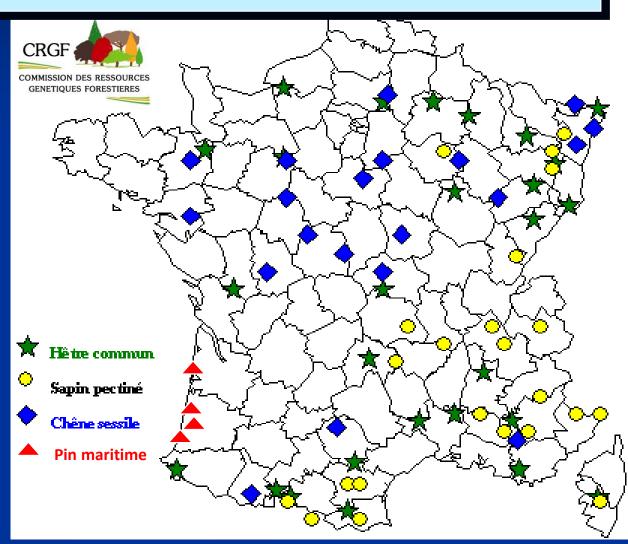
The French register of conservation units for widely occurring species

Abies alba: 21 CUs ~ 3500 ha (160 ha / CU)

F. sylvatica: 28 CUs ~ 3950 ha (140 ha / CU)

Pinus pinaster: 4 CUs ~ 900 ha (225 ha / CU)

Q. petraea: 20 CUs ~ 2400 ha (120 ha / CU)



http://agriculture.gouv.fr/conservation-des-ressources

In situ conservation of forest genetic resources (FGR): the pan-European dimension

A major political player: Euforgen

- Created in 1994 under Forest Europe
- Secretariat in Rome at Bioversity International
- Promote and streamline national FGR conservation strategies at European level (e.g. minimum requirements for CUs)

EUFORGE

- Raise awareness on FGR conservation of forest habitat

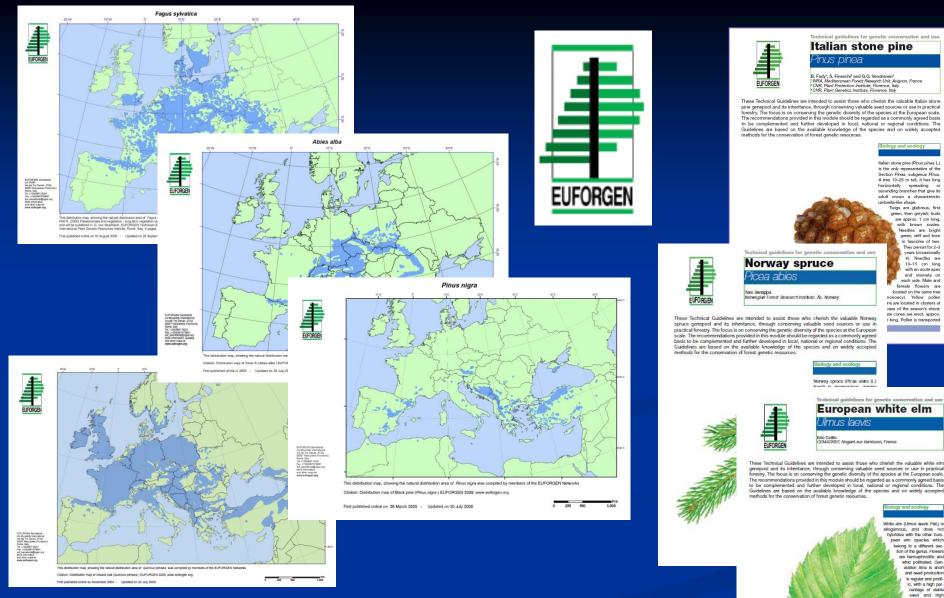
Euforgen Phase IV (2010-2014)



Scope: facilitating implementation of practical gene conservation by developing pan-European gene conservation strategies for forest trees (under the umbrella of Forest Europe)

- Working group 1: a European gene conservation strategy
- Working group 2: genetic monitoring methods
- Working group 3: guidelines for transfer and use of FRM
- Working group 4: incorporation of FGR in national policies

- Working group 5: managing CUs under climate change

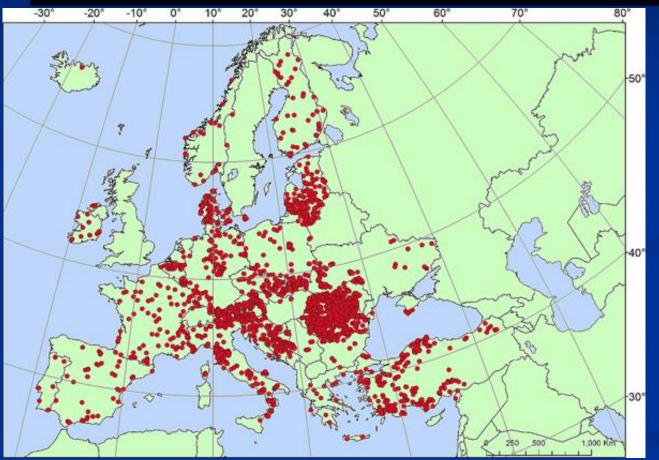


Seeds disperse by wind or came downstream b rivers enable th colonisation of new sites. Root suckering may play a role in th

regeneration of establishe stands whereas stool sucke ing is thought to be poor. The typical habitat of th while eim is ripartian diaciduce forest, where it can tolerate pri ionsel flooriding for longer period

Euforgen : more than technical guidelines and distribution maps widely used by researchers and managers alike

In situ conservation of forest genetic resources (FGR): the pan-European dimension



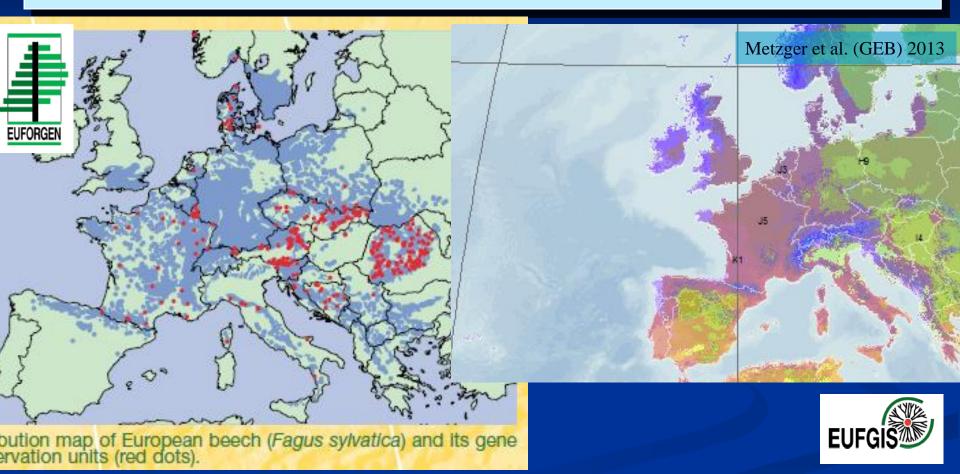




Koskela et al. (Biol Cons) 2013 Lefèvre et al. (Cons Biol) 2013

The EUFGIS database: **2774** CUs and **98** tree species in **31** countries. Each unit is managed for genetic conservation of one or more target tree species under a set of minimum requirements.

In situ conservation of forest genetic resources (FGR): the pan-European dimension



A tool for identifying gaps in pan-European strategies, raising awareness on conservation needs, particularly at range margins

Why an ex situ approach ?

A pragmatic approach:

- When in situ methods are not possible (extirpation)
- Create and reinforce collections of genetic material
- Methodological development (gene banks, cryo-conservation, etc)

Appropriate models:

- Pandemic risks (insects, pathogens)
- Habitat destruction (e.g. river banks, land use change)
- Disseminated «noble» species at risk from sylviculture
- Genetic pollution
- => Marginal populations



An example of species under pandemic risk: Ulmus sp.

Espèces menacées par une pandémie

orme champêtre, o. de montagne, o. lisse



20 ans de la CRGF, colloque 16 nov. 2011, Paris

conservation <u>dynamique</u>

reconstitution de haies champêtres







An example of species where risks lie on habitat: field hedges



Species in field hedges such as wild cherry (*Prunus avium*), service tree (*Sorbus domestica*) and English walnut (*Juglans regia*) need ex situ conservation



An example of species where risks lie on habitat: English walnut

A conservation site for remarkable genotypes in southern France



English walnut (Juglans regia) faces extirpation in many forests as it is over logged for its highly valuable cabinet-making wood

Example of a species where risks lie on habitat degradation and genetic pollution



Example of a species where risks lie on habitat degradation and genetic pollution

CRGF

Salzmann pine (P. nigra salzmanni)



An integrated FGR conservation and sustainable use program for Salzmann pine in France

Funding : DRAAF LR 2007-2008 (38 kEuros HT excluding permanent staff) ONF LR 2009 – 2015 (500 kEuros HT excluding permanent staff) Partners : ONF LR - ONF CGAF Orléans – Pépinière des Milles

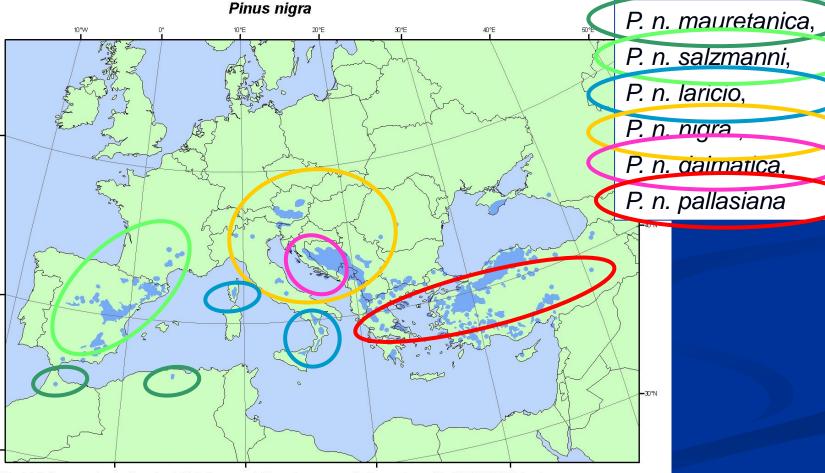




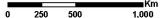
Geography and taxonomy of black pine







This distribution map, showing the natural distribution area of *Pinus nigra* was compiled by members of the EUFORGEN Networks and was published in: Isajev, V., B. Fady, H. Semerci and V. Andonovski. 2004. EUFORGEN Technical Guidelines for genetic conservation and use of European black pine (*Pinus nigra*). International Plant Genetic Resources Institute, Rome, Italy. 6 pages



Geography and taxonomy of black pine in France: marginal populations



Uniqueness, risks and protection needs

- Wild fires and climate change: risks on habitat (loss)
- Hybridization: potential risk (benefit?) for genetic resources

- Priority habitat under Habitats directive (D. 92/43 CEE of 21 May 1992) : « Endemic (sub-) Mediterranean black pine forests : Salzmann pine ».

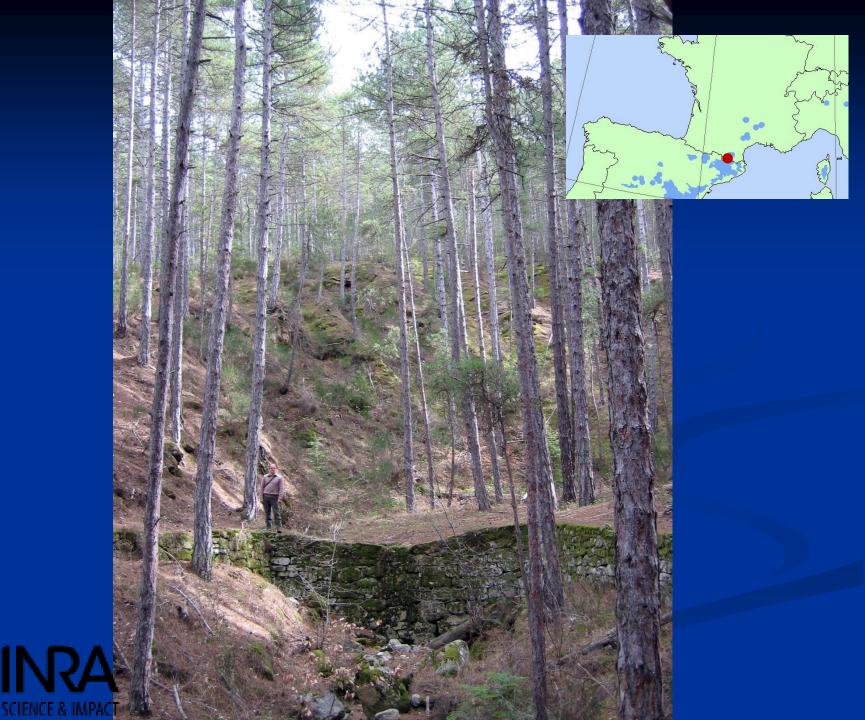
- European, national and regional issue: conservation and sustainable use within the national and regional biodiversity strategies.





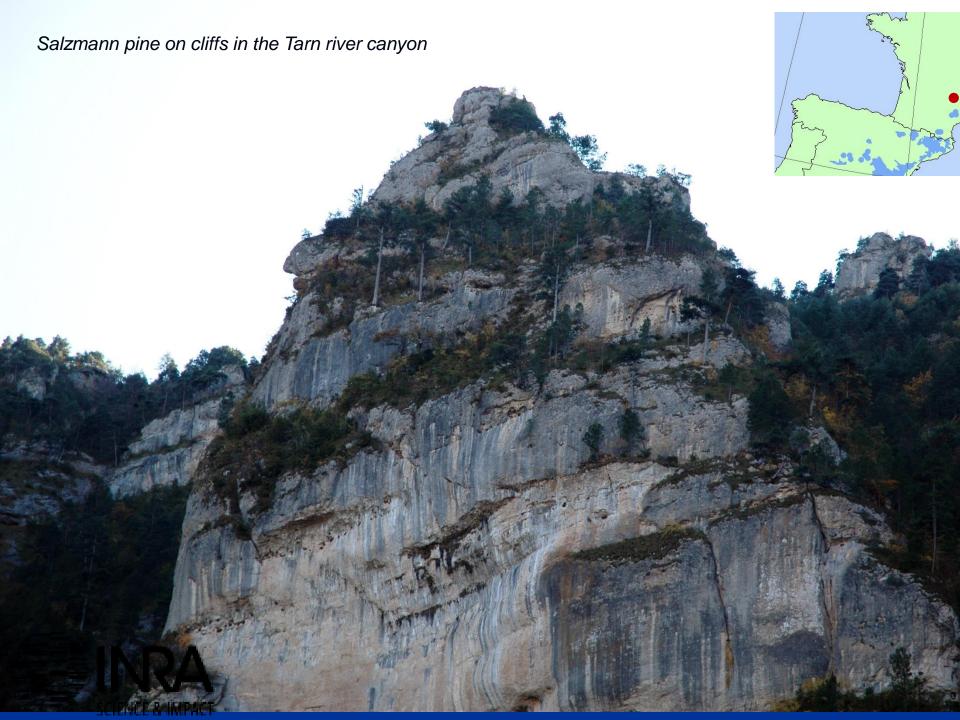












The objectives of the « Salzmann » project: ex situ collection of autochthonous pines

Finding autochthonous Salzmann pines in France.

Sampling grafts for ex-situ conservation in clonal plantations.

Sampling (leaves and seeds) for the study of genetic diversity:

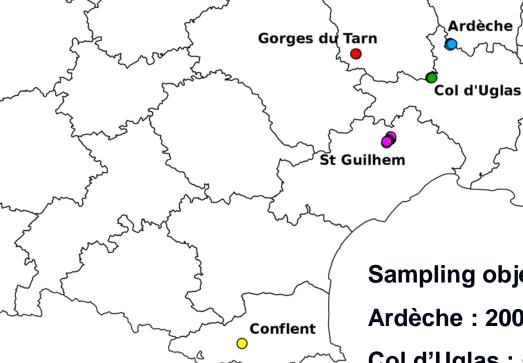
 Evolutionary history, uniqueness of populations for conservation

- Contemporary gene flow and mating system

Candidate genes of adaptive significance

SCIENCE & IMPA

Finding autochthonous Salzmann pines in France



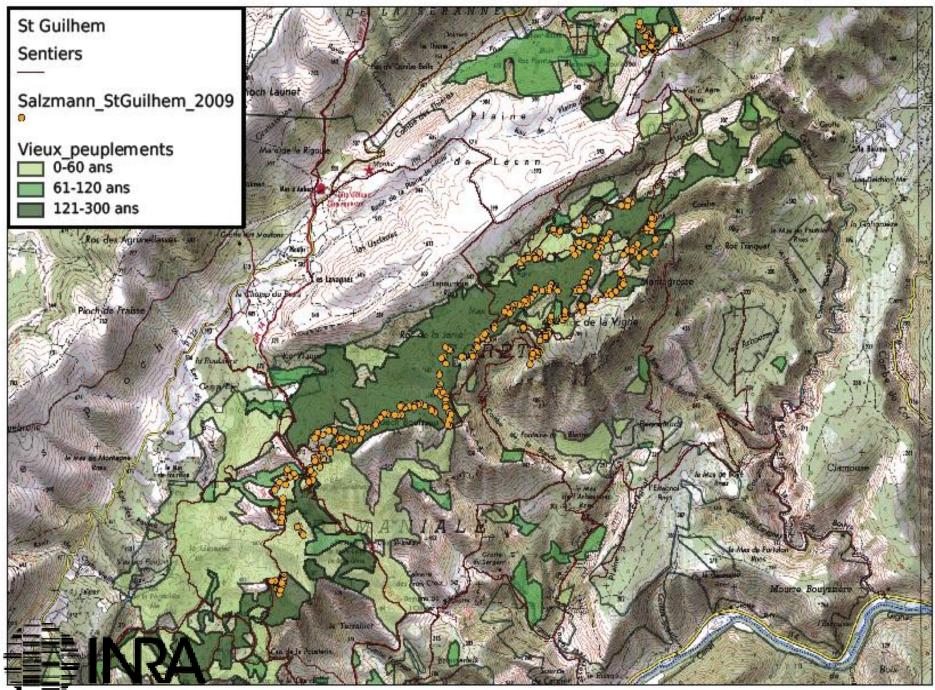
Sampling objectives for ex situ conservation: Ardèche : 200 adult individuals Col d'Uglas : 50 adult individuals Gorges du Tarn : 50 adult individuals Saint Guilhem le Désert : 300 adult individuals Conflent : 200 adult individuals

Finding autochthonous Salzmann pines in France

Wood cores from a candidate tree: age is only practical guaranty of autochthony

Finding autochthonous Salzmann pines in France





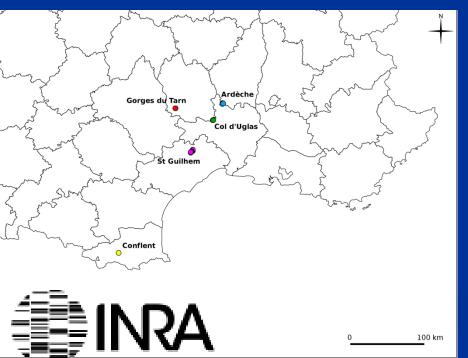
SCIENCE & IMPACT

Sampling for grafting and genetic monitoring

SCIENCE & IMPAC

Sampling for grafting and genetic monitoring

Sampling requires strong field experience. It is timeconsuming and expensive





A.C.



Plant material for grafting and conservation.





- Extracting seeds from cones Weighing X Rays for seed quality Data base and meta-data



Grafting (March – April)





Grafting: a high performance cloning technique with uneven success rates for old material

Success rate in 2008 : 20 genotypes out of 244 (8%)

Success rate in 2009 : 139 génotypes sur 260 (53%)

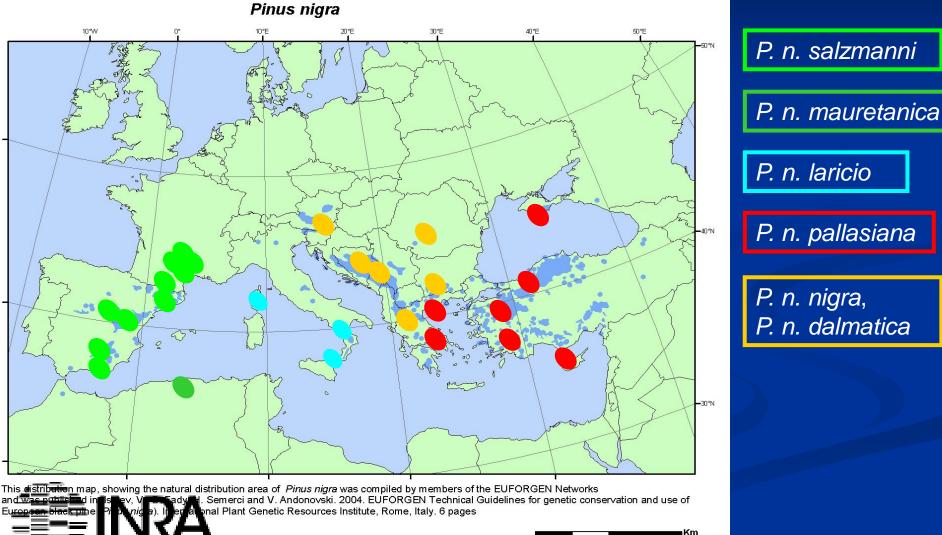
Success rate in 2012 : 741 génotypes sur 800 (92%)

Objective: 800 genotypes

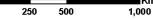




Retracing the evolutionary history of Salzmann pine and black pine

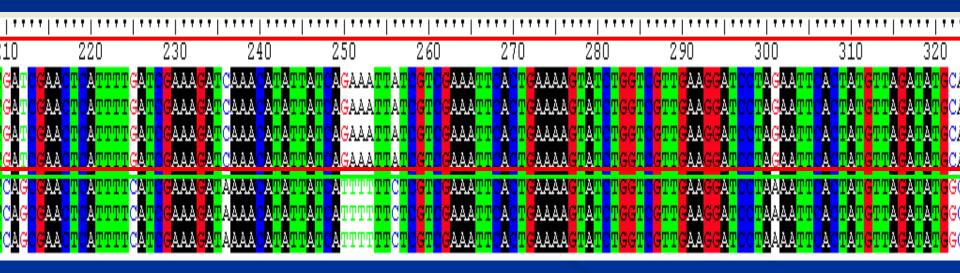


rst-outline on 26 March 2005 - Updated on 30 July 2008



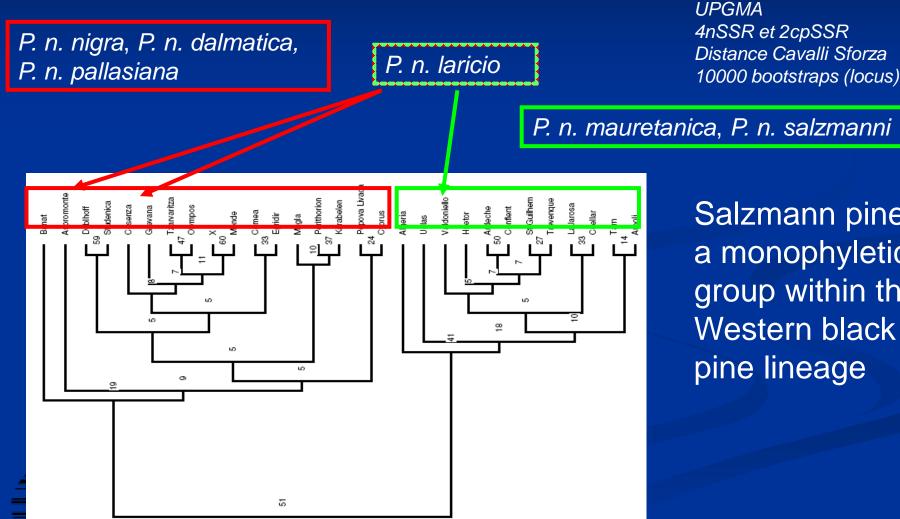
The DNA barcode approach: black pines are a homogeneous genetic group

Abies vs Pinus nigra : 10 genes, 6 amplify in conifers



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	460	470	480	490	500	510	520	530	540	550	560	570
AA <mark>C</mark>	A <mark>ac</mark> aaa <mark>c</mark> a-	- <mark>T</mark> GA <mark>AA</mark> TT <mark>G</mark> AA	CAG <mark>T</mark> A <mark>CC</mark> AAA	CCTTT <mark>CA</mark> TT:	rttt <mark>t</mark> ttt <mark>t</mark>	<mark>G</mark> AAA <mark>GG</mark> CTT	I <mark>gg</mark> tactg <mark>ttc</mark>	AG <mark>TT</mark> AA <mark>GC</mark> .	AA <mark>TAC</mark> AC <mark>AGT</mark> AA	CAA <mark>T</mark> CCAT <mark>C</mark> A	CAA <mark>TATT</mark> AT <mark>C</mark>	G <mark>CCTATT</mark> G
AA <mark>C</mark> /	AA <mark>C</mark> AAA <mark>C</mark> A-	- <mark>T</mark> GAAA <mark>TT</mark> GAA(CAG <mark>T</mark> A <mark>CC</mark> AAA	CC <mark>TTT</mark> CATT:	PTTT <mark>T</mark> TTT	<mark>GAAA</mark> GG <mark>CT</mark>	P <mark>GG</mark> TACTG <mark>TTC</mark> .	AG <mark>TT</mark> AA <mark>CC</mark> .	AA <mark>TAC</mark> AC <mark>AG</mark> TAA	CAA <mark>TCC</mark> A <mark>T</mark> CA	GAA <mark>T, ITT</mark> AT <mark>CO</mark>	G <mark>CC</mark> TATT <mark>G</mark>
AA <mark>C</mark> /	AA <mark>C</mark> AAA <mark>C</mark> A-	- <mark>T</mark> GAAA <mark>TT</mark> GAA	CAG <mark>T</mark> ACCAAAA	CCTTT <mark>CA</mark> TT:	PTTT <mark>T</mark> TTT <mark>TT</mark>	PT <mark>GAAA<mark>gg</mark>oti</mark>	I <mark>GG</mark> TACTG <mark>TTC</mark>	AG <mark>TT</mark> AA <mark>CC</mark> .	AA <mark>TAC</mark> AC <mark>AGT</mark> AA	CAA <mark>TCC</mark> A <mark>SC</mark> A	AAA <mark>TATT</mark> AT <mark>C(</mark>	G <mark>CC</mark> TATT <mark>G</mark>
AA <mark>C</mark> Z	AA <mark>C</mark> AAA <mark>C</mark> A-	- <mark>T</mark> GAAA <mark>TTG</mark> AA(CA <mark>GTACC</mark> AAA	CCTTTCATT:	PTTT <mark>T</mark> TTT <mark>TT</mark>	<mark>CAAAGC</mark> TT	I <mark>GGTACTGTTC</mark>	A <mark>GTT</mark> AA <mark>C</mark> C.	AA <mark>TACAC</mark> AG <mark>T</mark> AA	CAAT <mark>CC</mark> AT <mark>C</mark> A	GAATATTATCO	C <mark>GCC</mark> TATT <mark>C</mark>
4A <mark>C</mark> A	A <mark>C</mark> CAAA <mark>C</mark> AA	A <mark>T</mark> ACAA <mark>TT</mark> CAA	GCA <mark>TACC</mark> AAG	CCTTT <mark>CA</mark> ATA	AAAA <mark>T</mark> GAA <mark>T</mark>	<mark>GAAA</mark> GG <mark>CT</mark>	F <mark>GG</mark> TATGC <mark>TTC</mark>	AA <mark>TT</mark> AA <mark>C</mark> C.	AA <mark>TAC</mark> ACGA <mark>T</mark> AA	CAA <mark>TCC</mark> AT <mark>C</mark> A	GAG <mark>TATTAT</mark> C(CA <mark>CC</mark> TATT <mark>C</mark>
AA <mark>C</mark> A	A <mark>C</mark> CAAA <mark>C</mark> AA	A <mark>T</mark> ACAA <mark>TT</mark> GAA(GCA <mark>TACC</mark> AAG	CCTTT <mark>CA</mark> AT/	AAAA <mark>T</mark> GAA <mark>T</mark>	<mark>GAAA</mark> GG <mark>CT</mark>	F <mark>GG</mark> TATGC <mark>TTC</mark>	AA <mark>TT</mark> AA <mark>GC</mark> .	AA <mark>TAC</mark> ACGA <mark>T</mark> AA	CAA <mark>TCC</mark> AT <mark>C</mark> A	GAG <mark>TATTAT</mark> C(CA <mark>CC</mark> TATT <mark>G</mark>
AA <mark>C</mark> /	A <mark>C</mark> CAAA <mark>C</mark> AA	A <mark>T</mark> ACAA <mark>TT</mark> CAA	GCA <mark>TACC</mark> AAG	CCTTT <mark>CA</mark> AT/	AAAA <mark>T</mark> GAA <mark>T</mark>	<mark>GAAA</mark> GG <mark>CT</mark> I	T <mark>GG</mark> TATGC <mark>TTC</mark> .	AA <mark>TT</mark> AA <mark>GC</mark>	AA <mark>TAC</mark> ACGA <mark>T</mark> AA	CAA <mark>TCC</mark> AT <mark>C</mark> A	GAG <mark>TATTAT</mark> C(CA <mark>CC</mark> TATT <mark>G</mark>
AA <mark>C</mark> /	A <mark>C</mark> CAAA <mark>C</mark> AA	A <mark>T</mark> ACAA <mark>TTC</mark> AA(GCA <mark>T</mark> A <mark>CC</mark> AAG	CCTTT <mark>CA</mark> AT/	AAAA <mark>T</mark> GAA <mark>T</mark>	<mark>G</mark> AAA <mark>GGC</mark> T!	I <mark>GG</mark> TATGC <mark>TTC</mark>	AA <mark>TT</mark> AA <mark>CC</mark> .	AA <mark>T</mark> AC <mark>AC</mark> GA <mark>T</mark> AA	CAA <mark>TCC</mark> AT <mark>C</mark> A	GAG <mark>TATT</mark> AT <mark>CO</mark>	CA <mark>CC</mark> TATT <mark>G</mark>

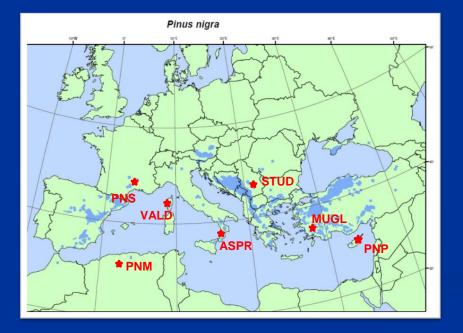
The genetic structure of black pines: two main phylogenetic groups... and strong gene flow



Salzmann pine: a monophyletic group within the Western black pine lineage

Focusing on 7 populations using nSSRs and SNPs at adaptive genes

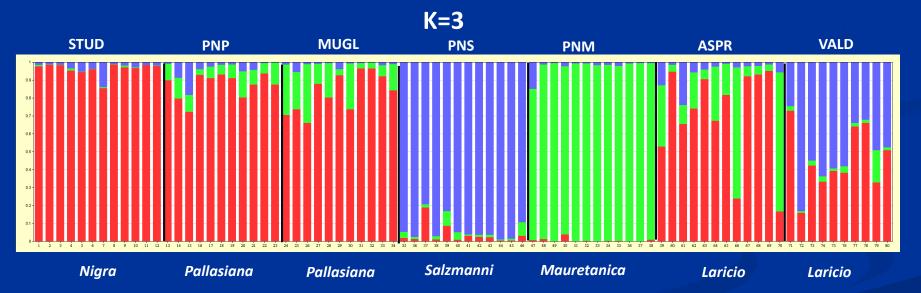
- Selected 12 genes (0_10162_01; 0_10384_02; 0_10667_02; 0_12216_02; 0_14221_01; 0_16810_02; 0_18101_02; 0_2078_01; 0_6293_01; 0_7916_01; 0_8479_01; 2_1405_01; CL4470Ct1_01)
- Optimization of PCR condition on a subset of individuals
- Amplification of 86 individuals from 7 populations
- **Editing of the sequences**



•PNM_P.nigra mauretanica (Algeria, Djurdjura moutains)
•ASPR_P.nigra laricio (Italy, Calabria, Aspromonte)
•VALD_P.nigra laricio (France, Corsica, Valdonielo)
•PNS_P.nigra salzmanni (France, Saint Guilhem)
•STUD_P.nigra nigra (Serbia, Studenica)
•MUGL_P.nigra pallasiana (Turkey, Mugla)
•PNP_P.nigra pallasiana (Cyprus)

The genetic structure of black pines: two main phylogenetic groups... but strong gene flow

TESS Bayesian clustering: Admix(BYM) model_nuclear SSR_nigra (4 nSSRs, 80 individuals); K from 2 to 10, 5 iteractions (Durand et al. MBE 2009)

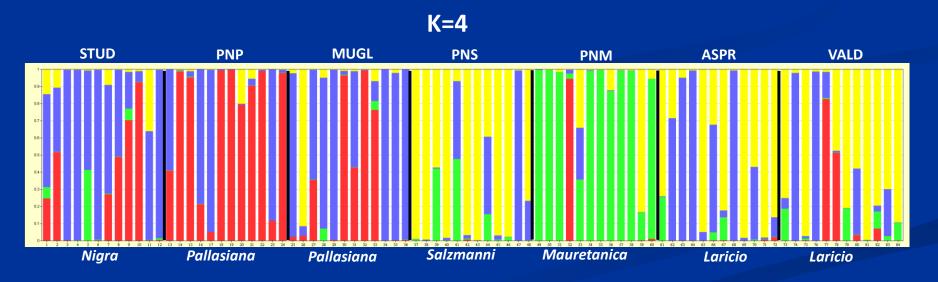




Courtesy of GG Vendramin

The genetic structure of black pines: two main phylogenetic groups... but strong gene flow

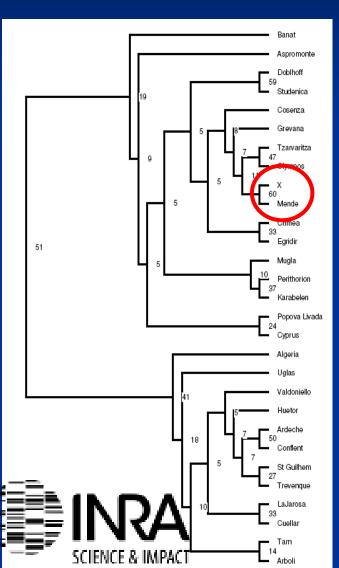
TESS: Admix(BYM) model_SNP_nigra (265 loci, 84 individuals) 10000 burns 50000 sweep; K from 2 to 10, 5 iteraction



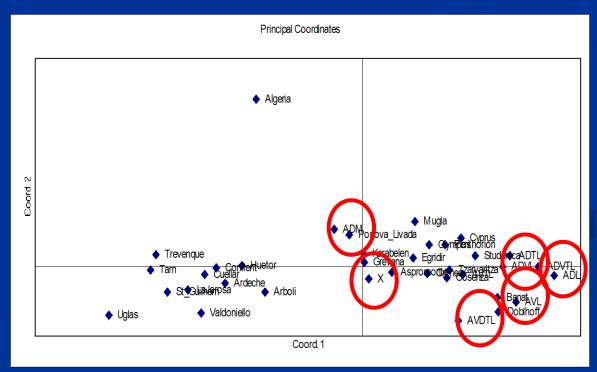


Courtesy of GG Vendramin

Hybridization in France : fiction or reality?

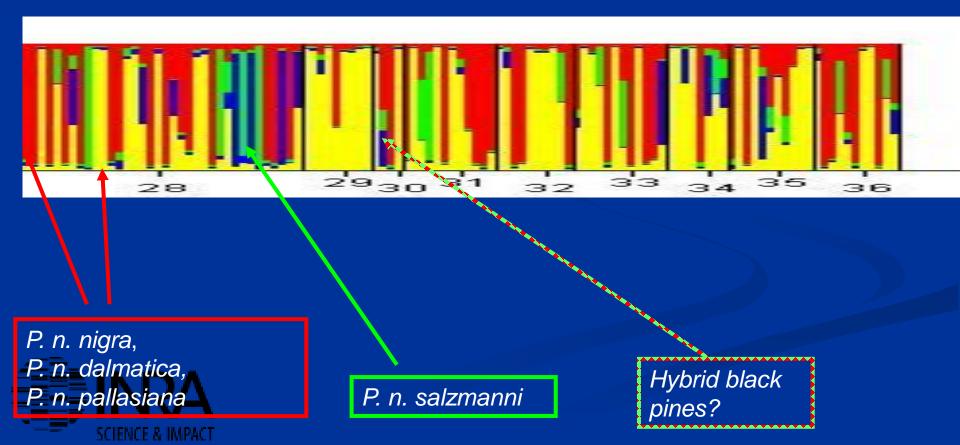


One phenotypically suspect population (Mende, 8 morphological types) et one potentially hybrid from gene flow (Parlatges = X)



Hybridization in France : fiction or reality?

Detecting hybrids at K = 4. To be continued using more markers



Considering adaptation in addition to demography and evolutionary history. Drift or local adaptation?

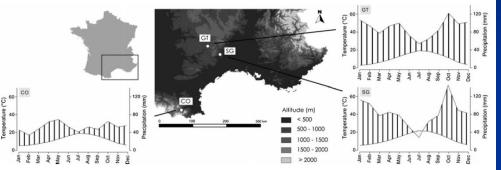


Fig. 1 Location of studied sites. Gaussen diagrams of each site are presented. Climatic data are calculated for the period 1961–1990. Bar plots represent the sum of monthly precipitations, and white-background plot, monthly temperatures. The intercept of two curves represents a dry period

• *Results* The Conflent population showed a strong sensitivity to spring precipitation deficits (March to June), while at Gorges du Tarn and Saint-Guilhem sites, autumn (October) of the previous year and winter (February) temperatures explained more variance than precipitations and were, respectively, negatively and positively correlated to radial growth.



Amodei et al. (AFSc) 2013

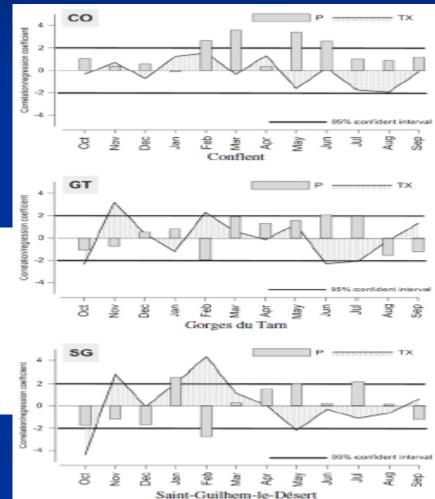


Fig. 2 Correlation/regression coefficients between monthly maximum temperatures (TX) and monthly precipitations (P) and standardized

HÉRAULT Une pouponnière s pour les pins de Salzmann



→ Une grande attention est portée à cette espèce menacée

RÉGION

→ 4 000 greffons ont été récupérés dans le massif do Saint-Cuilhom

Photo Gh

Le pin Salzmann menacée par le...

→ L'objectif : atteindre

les 800 individus

clonés ou greffés

RÉGION

FAITS DIVERS Montpellier Un forcené maîtrisé

me de 29 ans qui, de

ain a été éva

→ L'Office national

des forêts pilote

l'espèce en voie

en cette belle journée, des techniciens forestiers de l'In-ra perchés sur les plus hautes branches des pins de Saiz-mann du massif de Saint-Guil-hem-le-Désert (Hérault), sont

te de greffons et des cônes. Quelque 4 000 greffons et 1 000 cônes ont été récupérés

dans la région à des fins de re-cherche et de reproduction. Les greffons sont destinés à

Les greffons sont destinés une pépinière d'État charg de la multiplication végéta ve. Quant aux graines, elles i téressent les généticiens l'Inra d'Avignon et ceux e

conservatoire génétique des arbres forestiers d'Orléans.

Des opérations qui entren

pour mieux la protéger, mais

aussi pour son intérêt sylvico-

lassi poin son antre spino le, « Le più de Satzmann qui n'était jusqu'à présent qu'une curiosité botanique est de-puis quelques années portaur d'espoir », explique Daniel Cambon, responsable techni-que ONF du projet. Cette espè-

espèce locale

nous touche »

one arrivée à provi

« Cette

d'extinction

un programme de conservation de

par le GIPN mmes du Groupe d'in ion de la police natio e (GIPN) sont intervenus en fin d'après-midi, dans auartier de Port-Mariann toellier Ils ont maîtris

s le CHU

rontignan Un

puis se rend

vers 18 h, les po

sont intervenus à la cité

tte, à Frontignan, Dan

cait de me

ée mais f

ns. s'est rendu à la po

duit à l'hônital de Sèt

gardiens agressés

maison d'arrê

and serait lege

nes, lors d'une prome

re du nez et de dent

ent touché. Les syndicat

appelle « qu'il n'y a pa

he dans les c

sau'à 40°C

rpignan

incendié

Un immeuble

de L'un couffrirait d'un

Jîmes Deux

à la prison

orcené menace

Hérault L'ONF au chevet des derniers pins de Salzmann



Le technicien forestier de l'Inra, Norbert Turion, cueille o Des operations qui entrent dans le cadre d'un programme de conservation (2008-2012) des populations françaises de pin de Salzmann et doté d'un budget de 500 000 €. L'objec-tif est d'étudier cette espèce

ce autochtone a traversé les siècles. Autre qualité : elle fournit un excellent bois d'œu-vre. Dans le massif de Saint-Guilhem, l'étude des charbons (*) (résidus de feux) vestre. « La problématique c'est que cet arbre n'est pas forcément pur sur le plan géndtique car nos prédéces seurs ont reboisé à l'époque charbons (*) (riskultus de froz), seurer out revisei de Noue-septen visi-arpeia U, exploratent déja construite de Soura-septen visi-arpeia U, exploratent déja con-d'Autriches. Le pris de Saltz-construit de Soura-construite de la construite de la construi-construite de la construite de la construite de la construi-construite de la construite de la construite de la construite construite de la construite de la construite de la construite construite de la construite de la construite de la construite construite de la construite de la construite de la construite construite de la construite de la construite de la construite construite de la construite de la construite de la construite construite de la construite de la construite de la construite de la construite construite de la construite de la construite de la construite de la construite construite de la constru

approché cette centaine d'ar-bres reliques qui n'ont survé-le terrain, sa technicité, elle con que parce qu'ils étaient ac-constitue le premier maillon crochés à la falaise et donc de la chaîne des programmes de ces trois sont multiples et souvent pla forestiers, rattanifiées sur plusieurs and hés à l'unité expérimentale Le travail requiert d'exceller terranéenne

Accéder au site | Toutes les vidéos

les graines destinées au reboi-sement solent issues de souches pures. Aussi afin d'éviter les sujets hybrides, le projet a débuté avec le recensement des vieux arbres ceux de plus de 140 ans, présents avant les reboisements. Ces ancêtres, baptisés à l'origine pins de Montpellier, ont été sondés, marqués, positionnés par GPS ouis cartographiés, 280 arbres (dont 120 à Saint-Guilhem) ont été sélectionnés sur différents peuplements. Et ce n'est pas terminé, l'objectif est de 800 individus clonés, greffés ou mis en collection. • Texte et photos : Ghislaine GUIBAUD

Copier pour protéger

1 400 sujets vont être treffés, explique Patrice

Brahic responsable de la

Pépinière forestière d'État

nière. On table sur 50

de pertes environ, dues à la difficulté technique de

l'opération. A raison de

six areffons par sujet, nous

ons créer trois

par pin sauvegardé. » Récupérés par l'ONF, les plants sont destinés à la

réation d'un verger à

graines dans le Sud-Ouest

Recherche par ADN

travaillé à l'extraction d'AD

Sentification génétique

ensuite nous étudierons le

graines. Il s'agit de connaîtr

u du feuillage et à

énétique de chaque unlement afin d'établi une carte d'identité des populations françaises de

ces pins et de sélections

boisement « explique

Bruno Fady, directeur de recherches à l'unité Ecologi

des forêts méditerranéer

d'Avignon. Les premiers

résultats montrent que les lorêts de Saint-Guilhem et

de l'Ardèche proches sur le plan génétique, présenten également une grande

diversité génétique et sont,

de ce fait, intéres

graine pour le

ceux qui seront les plus à même de produire de la

d'Aix-en-Provence, Les

plants resteront a

(*) Etude de Jean-Louis Verner (CNRS Montpellier)

Patrimoine

Autrefois répandu jusqu'en plaine, le pin de Salzmann noir découverte en 1810 par Salzmann, botaniste alle-mand, dans le massif de Saint-Guilhem (Hérault), et présente a l'état naturel dans le sud de la France et en Espanne. Elle a révressé en Espagne. Elle a régress jusqu'à devenir une des es sences les plus rares de Fran ce avec 3 000 hectares (100 fois moins qu'en Espa-gne) situés majoritairement dans les forêts publiques du Languedoc-Roussillon.

les pins de S en voie de dist



n est au cœur de

cela nous tor

umt Et





The story continues... The evolutionary history and the genetic diversity of key functional traits in Pinus nigra: consequences for the conservation of Pinus nigra salzmannii

- The PhD thesis of Guia Giovannelli
- New genomic resources from transcriptome sequencing: collaboration with CNR Florence
- Is there local adaptation for radial growth?
- A new niche model for Pinus nigra subspecies



Conserving genetic resources of marginal populations: challenges that remain

- A need for conservation strategies that include a consideration of genetic diversity
- A need to prove the « value » of marginal populations (demography and adaptation)
- A need to link with habitat conservation and adaptive management strategies under global change
- A need for both large scale political agreements and coordination and for locally sound implementation
- A need to convince society!



A need to include marginal populations in genetic and habitat conservation networks

Thank you for your attention!



H ANG

ProCoGen dissemination workshop "Genomics and the conservation of conifer genetic resources" -Szombathely, Hungary, 1-3 Septem