

Demographic and evolutionary history of the European black pine (Pinus nigra Arn)

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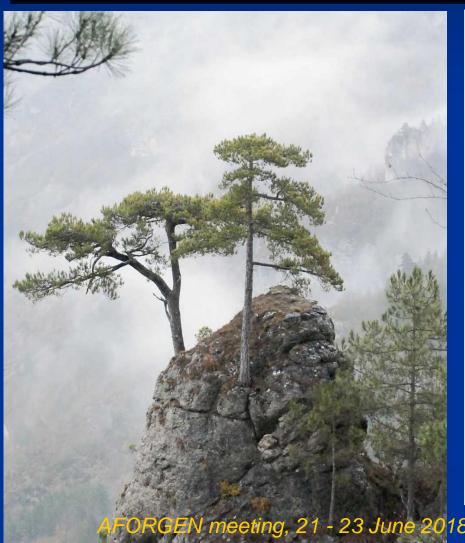
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Demographic and evolutionary history of the European black pine (Pinus nigra Arn)



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GEN méeting, 21 - 23 June 2018, Kranjska Gora, Slovenia

Do you know the European black pine?

The conifer tree that everyone loves to hate, and which foresters plant everywhere...



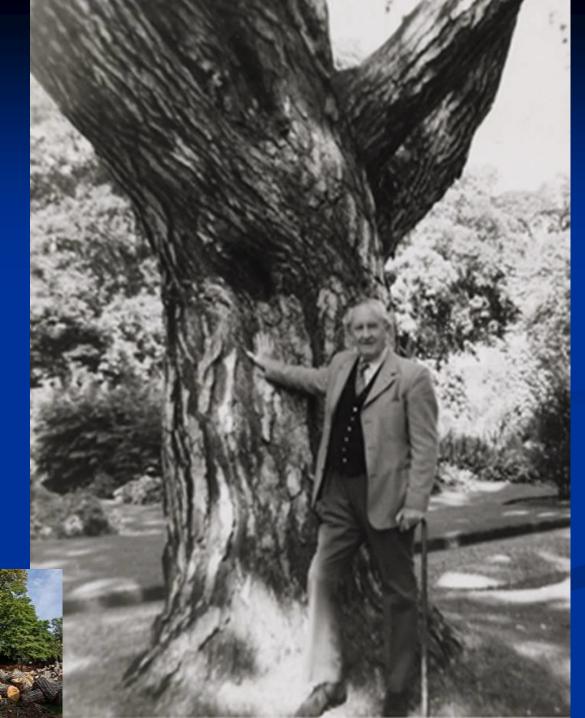
This is a beauty of dissonance, this resonance of stony strand, this smoky cry curled over a black pine like a broken and wind-battered branch when the wind bends the tops of the pines and curdles the sky from the north.

> A.J.M. Smith Canadian poet (1902-1980)

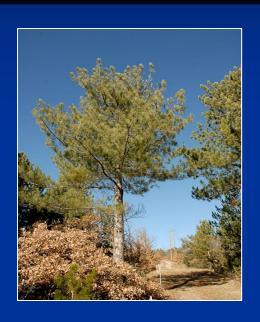


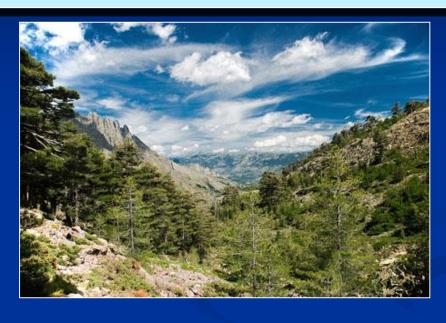
One tree to rule them all: JRR Tolkien with the black pine in Oxford Botanic Garden



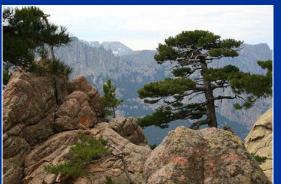


A pioneer species with low ecological requirements





Resistance to drought



Low soil pH and substrate specificity

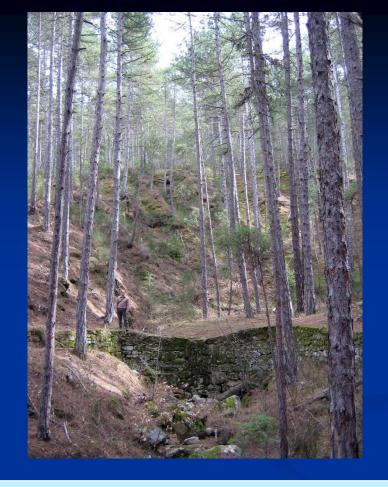
One of the most productive, multiple purpose, native European conifers

- > Over 3.5 million hectares of native forests
- ➤ Mean productivity from 6-10 m³/ha/year (natural conditions) to 8-20 m³/ha/year (monoculture)
- > Durable wood, high resin content, easy to work





- Ecological restoration
- Erosion control
- Reforestation
- Parks and recreation



... But whose evolutionary history we actually know little about!

A habitat at stakes: uniqueness, risks and protection needs

Priority habitat under "Habitats directive D. 92/43 CEE"
 (21 May 1992): « (Sub-) Mediterranean pine forests with endemic black pines »

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



Also an ecological asset: uniqueness, risks and protection needs

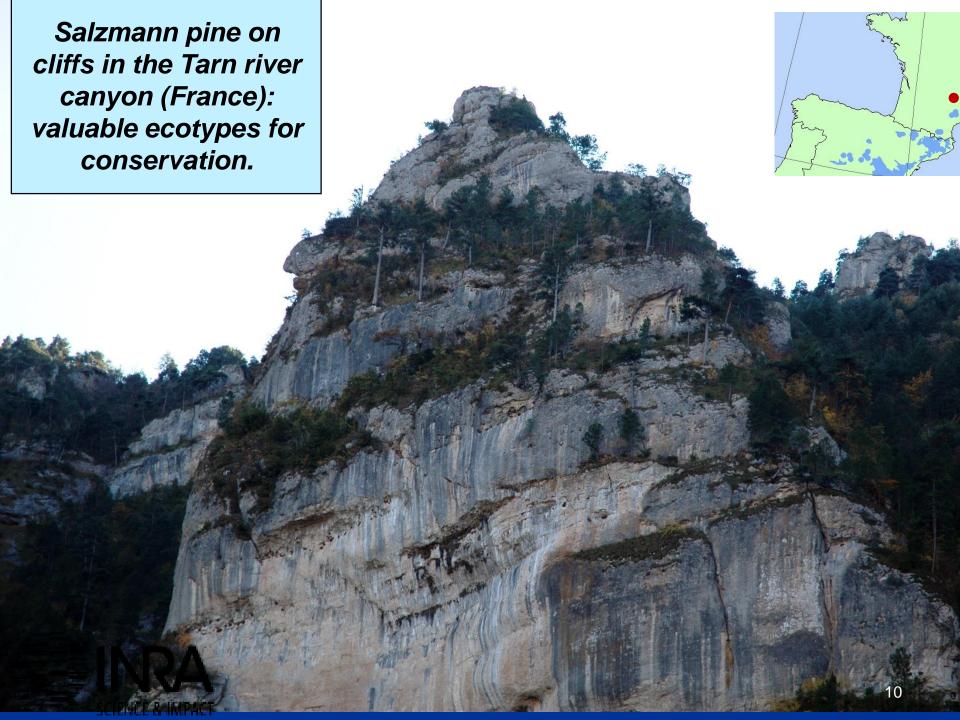
Wild fires and climate change: risks on habitat (loss)

 Hybridization: potential risk (benefit?) for genetic resources. In France, 5 000 ha native forests versus 300 000 ha planted Austrian and Corsican black pine

forests







Chorology of black pine and its 3.5 million ha of forests: fragmentation rules!

Pinus nigra



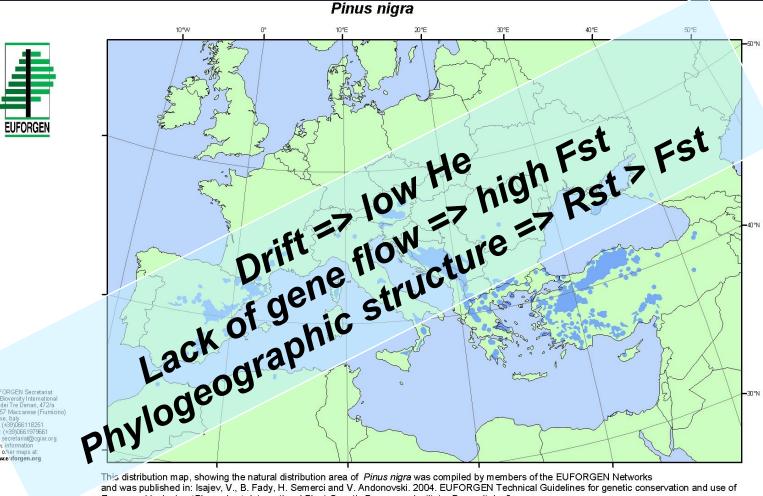


EUFORGEN Secretariat cob Bioversity International Via dei Tre Denari, 472/a 00057 Maccarese (Fiurnicino) Rome, Italy Tel. (+39)066118251 Fax: (+39)0661979561 euf_secretaria@cgiar.org More information and other maps at: www.euforgen.org

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

Expectations under long-term, strong fragmentation and a common ancestor

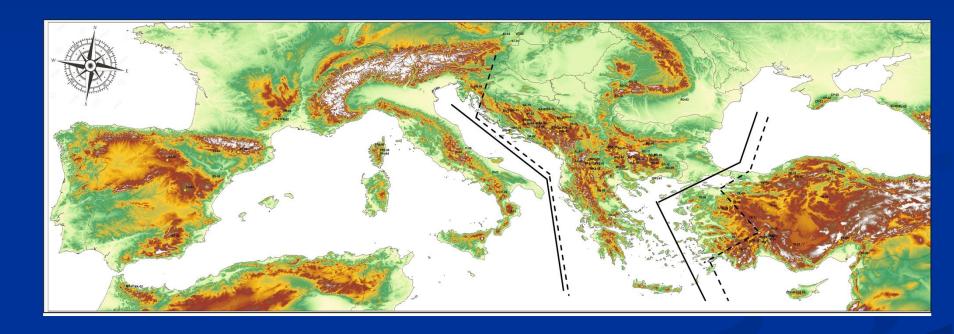




EUFORGEN Secretariat c/o Bioversity International Via dei Tre Denari, 472/a 00057 Maccarese (Fiumicino) Fax: (+39)0661979661 eut_secretariat@cgiar.org and other maps at

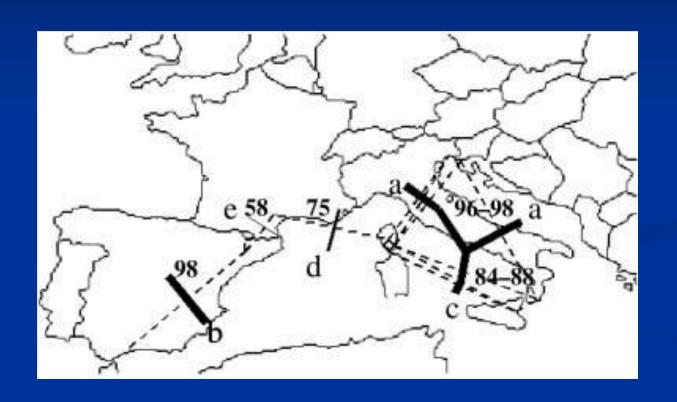
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Contradictory demographic scenarios (1)



Naydenov et al., 2016, 2017 (cpSSR sequences): Ancient split into 3 lineages, between 4,37 and 0,93 My ago 272 haplotypes, Gst ~0.177, H~0.29

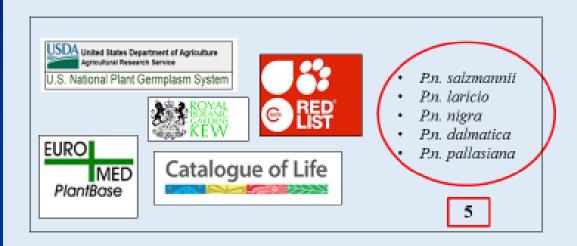
Contradictory demographic scenarios (2)



Fady et al., 2010 (cpSSR length polymorphism): 229 haplotypes, Fst ~ 0.11, He ~ 0.95

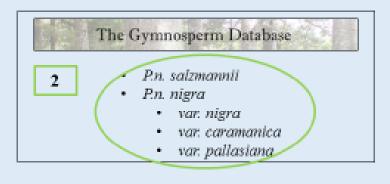
Afzal-Rafii and Dodd, 2007 (cpSSR length polymorphism): Recent split into 6 lineages, between 150 and 30 kY 235 haplotypes, PHIct ~ 0.35, He ~ 0.95

Taxonomic uncertainties





- P.n. salzmannii
- P.n. laricio
 - var. calabrica
 - var. corsicana
- P.n. nigra
 - var. nigra
 - · var. pallasiana







3

10

Differentiation and the evolutionary history of black pine

Sampling:

19 populations (12 trees per population)





DNA markers:

4 barcoding (organelle) genes

4 cpSSR (Vendramin & al. 1996),

14 nSSR (Giovannelli & al. 2017),

14 « candidate » genes (SNP) (Mosca & al. 2012),

Differentiation and the evolutionary history of black pine

Sampling:

19 populations (12 trees per population)





Results:

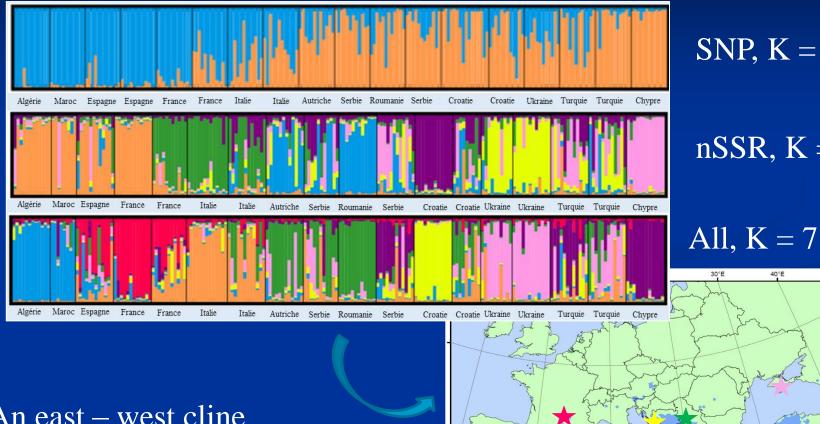
Barcoding genes: no diversity

cpSSRs: 143 haplotypes, He~0,86, Fst~0,05

nSSRs: 0,30<He<0,85, Fst~0,138

"Candidate genes": 0,53<He<0,75, Fst~0,12

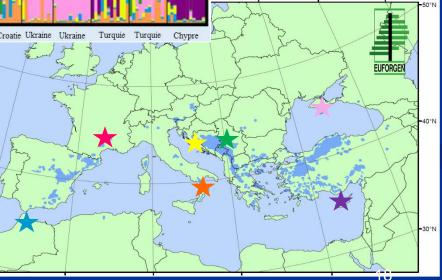
The most likely genetic structure of black pine range-wide



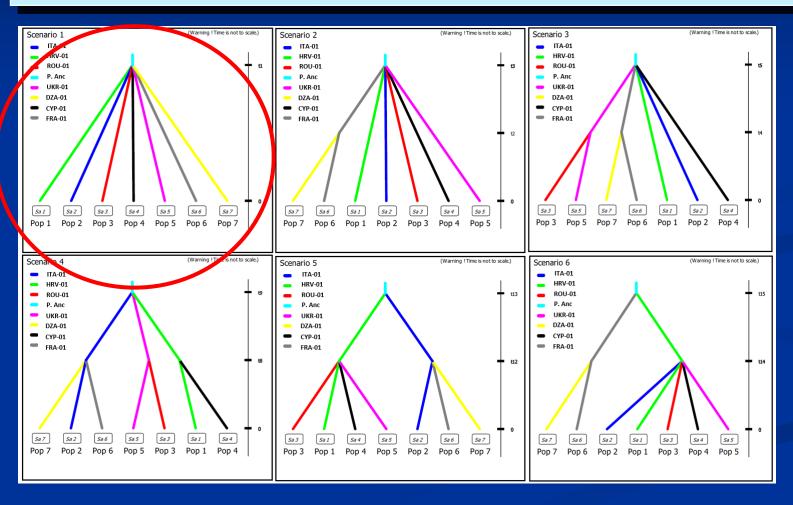
- An east west cline
- \triangleright (2 to) 7 different lineages
- High gene flow among populations

SNP, K = 2

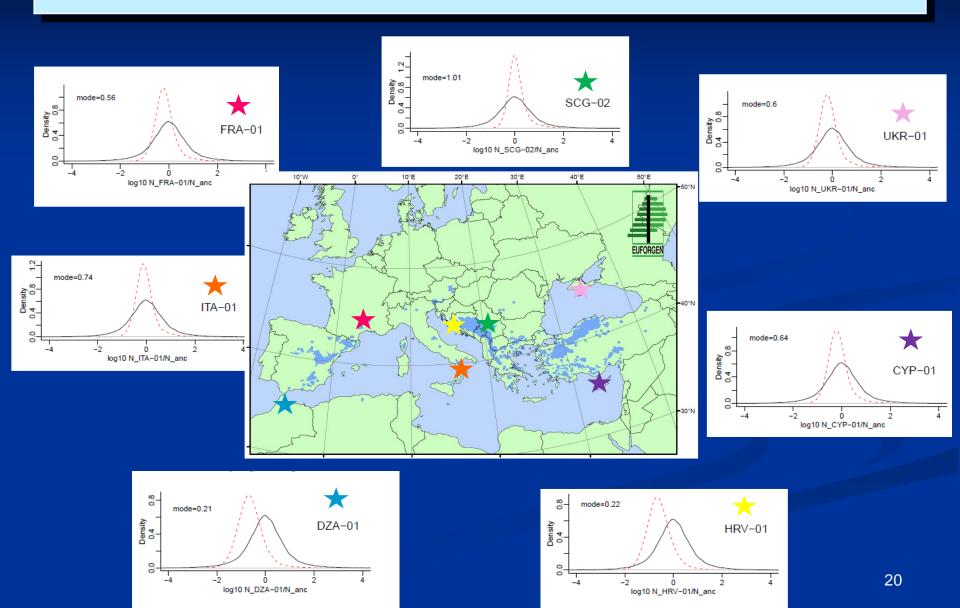
nSSR, K = 6



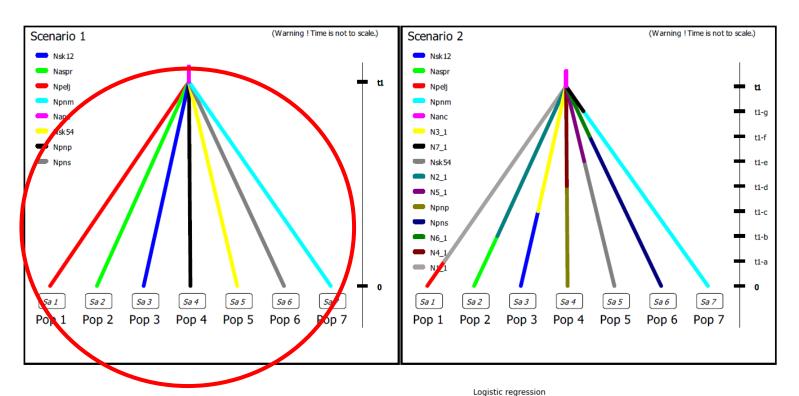
The genetic structure of black pine is best explained by a last-glaciation-time demographic split into 7 lineages

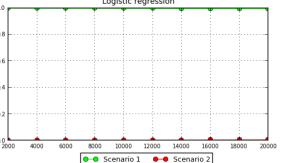


Expansion or contraction in the 7 lineages

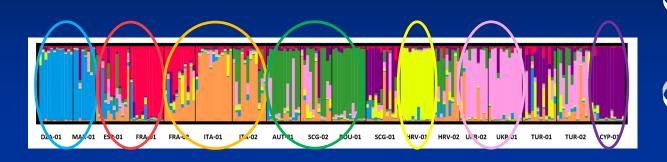


... And no bottlenecks





Identification of 7 different lineages



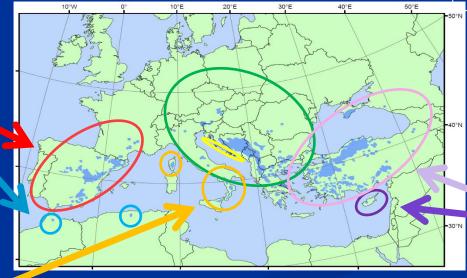
Black pine and its sub-species (?):

- ► P.n. mauretanica
- ► P.n. salzmannii
- > P.n. laricio
- ➤ P.n. nigra
- > P. n. pallasiana
- ► P. n. dalmatica
- ► P.n. caramanica

Algeria, Morocco (P.n. mauretanica)

 \neq or =

France, Spain (P.n. salzmannii)



Turkey/Ukraine (*P.n. pallasiana*)

#

Cyprus (P.n.

caramanica)

Corsica (P.n. laricio)

=

Calabria (P.n. laricio)

Giovannelli 2017, Giovannelli et al. (in prep)

The evolutionary history of the European black pine

- High levels of gene flow despite fragmentation and moderate gene diversity (Fst~0.06-0.10; He~0.8);
- Weak phylogeographic and past climate signals;
- A relatively recently (Holocene) structured genetic diversity of 7 lineages;
- A few genetically depauperate lineages



The evolutionary history of the European black pine

- Some opportunities for taxonomic fingerprinting:
 => detecting man-made translocation events;
 => selecting gene conservation units outside of man-made hybrid zones;
- A need to revise taxonomy and possibly legal (forest reproductive material) documents;
- Outlook: SNP from transcriptome sequences.



The European black pine: there is more than meets the eye...

A structured diversity across its distribution range that can guide gene conservation strategies and forest management

