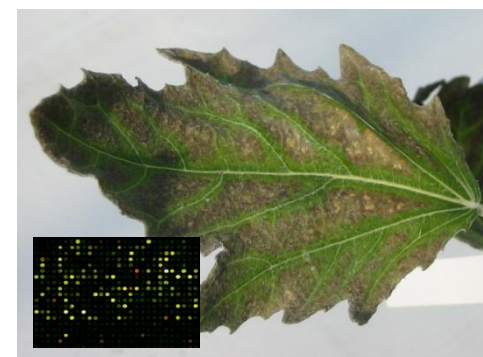


Risposta adattativa di alberi forestali e miglioramento genetico



Sabatti M.¹, Beritognolo I.², Villani F. ², Malvolti M. E. ²

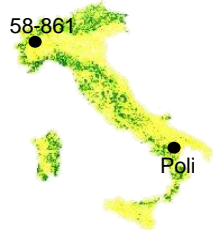
¹ Dipartimento per l'Innovazione nei sistemi Biologici Agroalimentari e Forestali. Università degli Studi della Toscana. Viterbo

² Istituto di Biologia Agro-ambientale e Forestale. Consiglio Nazionale delle Ricerche. Porano (TR)

Esperienze e prospettive del miglioramento genetico degli alberi forestali in Italia
Consiglio per la Ricerca e l'analisi dell'Economia Agraria – Azienda Ovale
Roma, 30 maggio 2018

Experiences on fast growth and multipurpose forest species

F₁ *Populus nigra* family (POP5)



162 clones planted in:
Cavallermaggiore (CAV, 2004)
Savigliano (SAV, 2008)
Viterbo (VT, 2008)
Montelibretti (MT, 2004)

Poplars
Fast growth,
Biomass,
Wood,
Agro-forestry systems
Ecosystem services

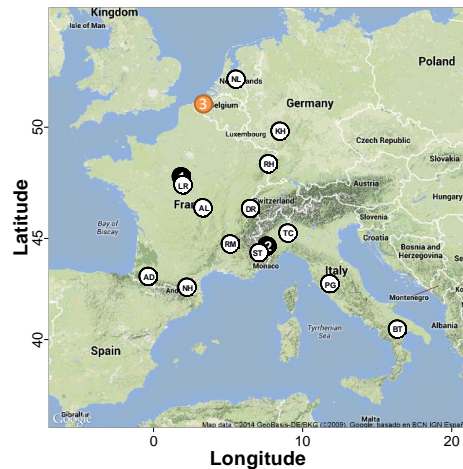
Chestnut
Multipurpose,
Fruits,
Food,
Agro-forestry systems
Ecosystem services

Walnut
Multipurpose
Fruits,
Wood,
Agro-forestry systems
Ecosystem services

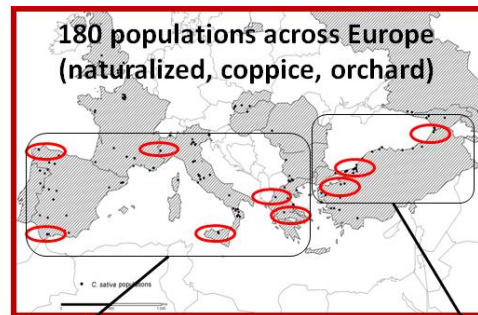


J. regia ecotypes

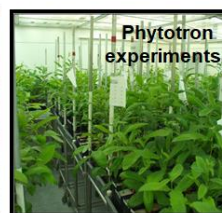
P. nigra association population



Castanea sativa populations & full sib families



Provenance
field trials

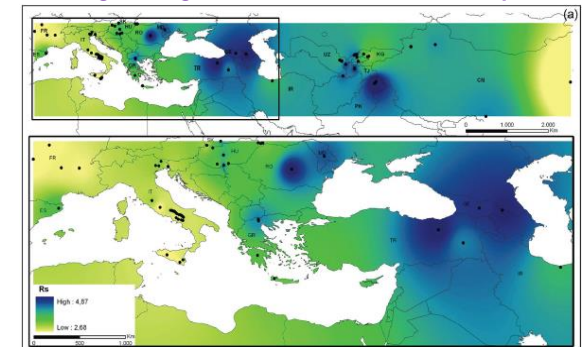


Phytotron
experiments



Full sib families

Juglans regia allelic richness across Europe



Full sib families



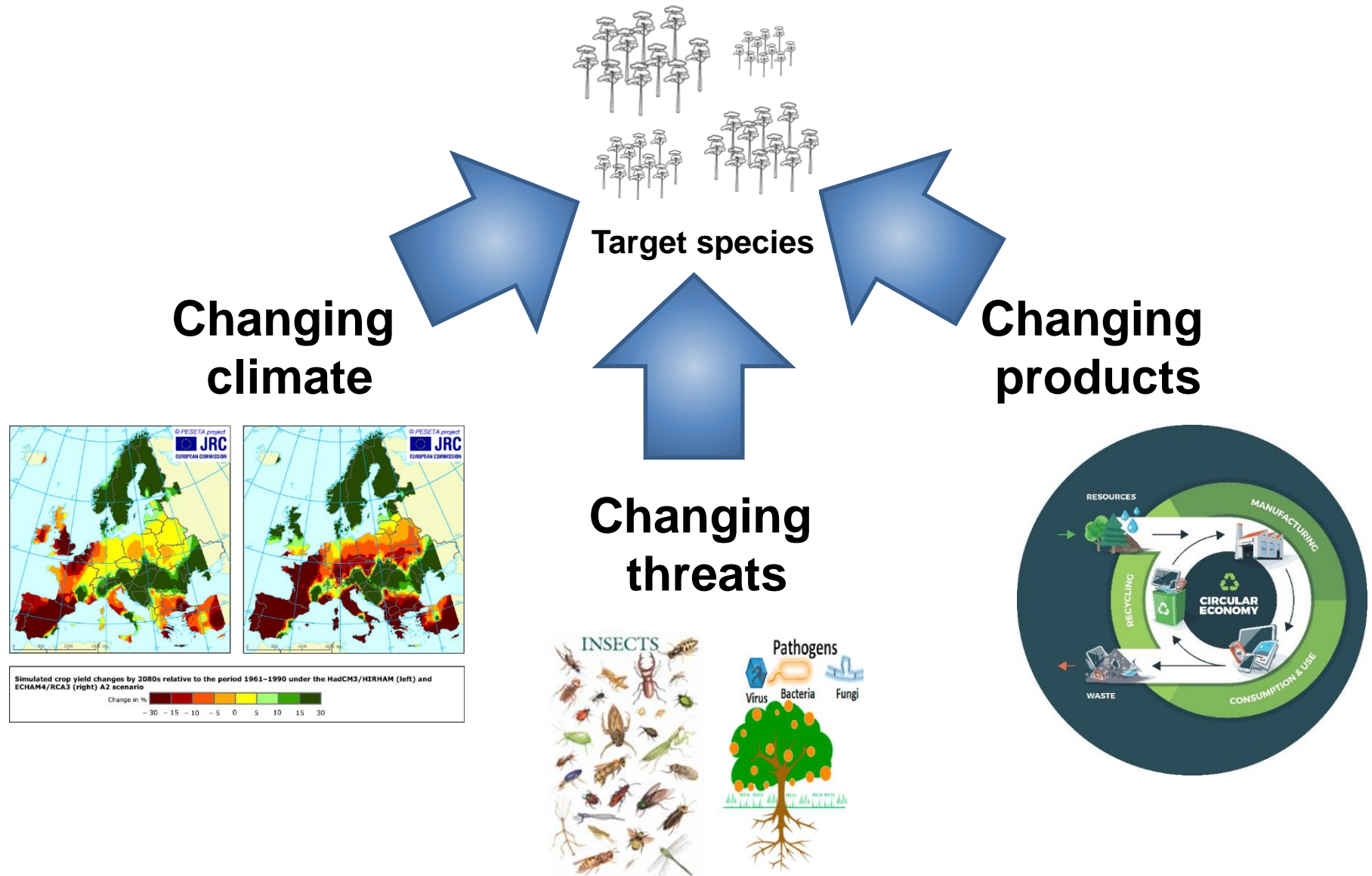
Grafting



J.regia varieties

Geographic origin of 13 metapopulations of the *Populus nigra* collection (white circles) and geographic position of the three field trials: ORL (1) and SAV (2) (black circles), GDB (3) (red circle).

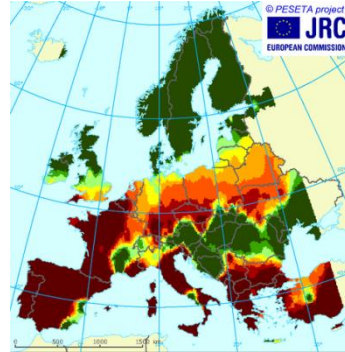
Future challenges of tree genetic improvement



Breeding for long lived species

Requests driven by changing environmental, productive and societal needs

General objectives:

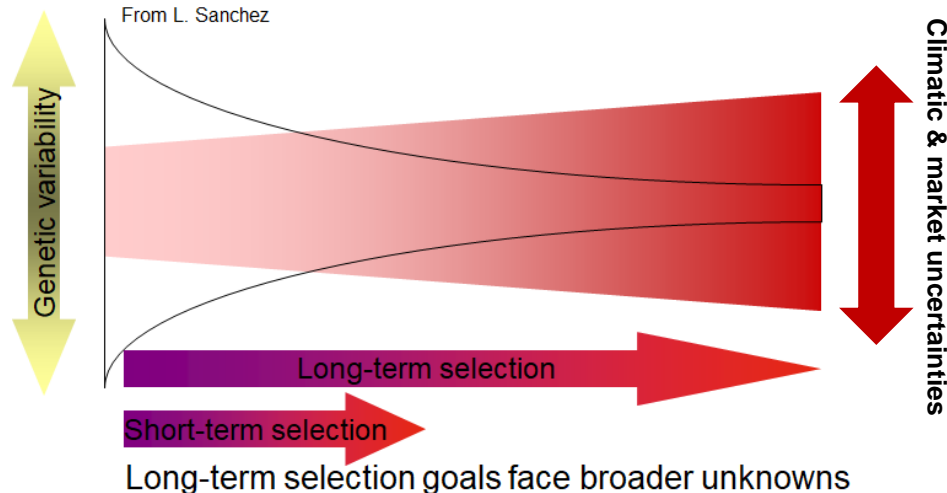


**Management and
conservation
of genetic resources**

Adaptability
stability of performances in
changing environment

Specific added values
(new markets, circular
economy)

**Continuous genetic
gains**
for wood and fruit
production



➤ **Multidisciplinary research activities at different scales: gene, individual, population, landscape**

A common parameter: Genetic Variability

Wide geographic range and high adaptive potential

Populus nigra



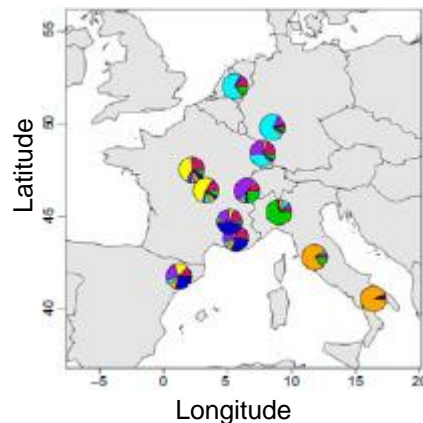
MOLECULAR ECOLOGY RESOURCES

Molecular Ecology Resources (2016) 16, 1023–1036

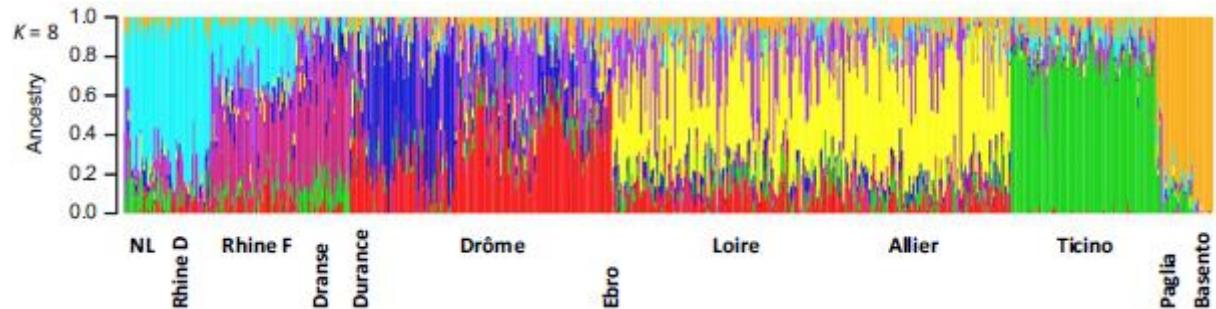
doi: 10.1111/1755-0998.12513

New resources for genetic studies in *Populus nigra*:
genome-wide SNP discovery and development of a 12k
Infinium array

P. Faivre Rampant et al. 2016



Geographical distribution of the
populations and the genetic
structure revealed by ADMIXTURE.



Population structure analysis estimated from 600 SNPs distributed
throughout the *Populus nigra* genome in validated genotypes.

- Seven gene pools across the *P. nigra* Western range
- High level of admixture, clear genetic differentiation between populations
- Italian populations structured along a latitudinal gradient, low level of admixture.

A common parameter: Genetic Variability

Wide geographic range and high adaptive potential

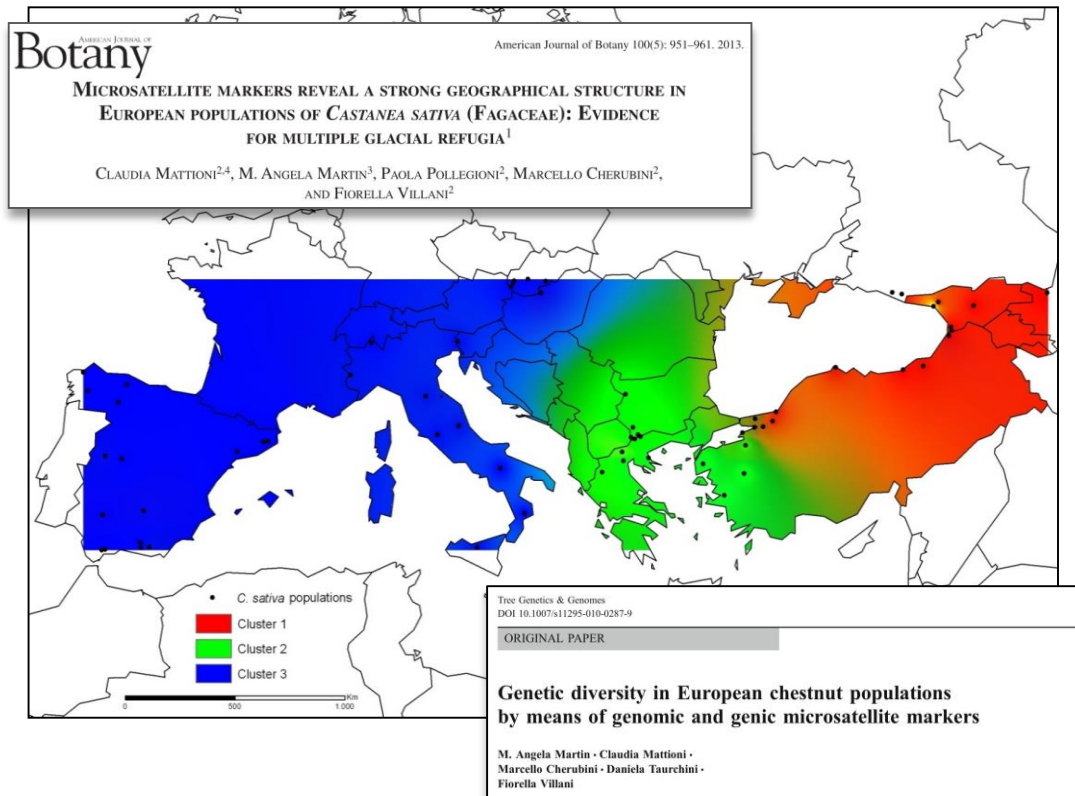
Castanea sativa



Genetic resources: genetic diversity

Genetic structure of populations

Main gene pools of *C. sativa* natural populations in Europe
Neutral diversity (SSRs)



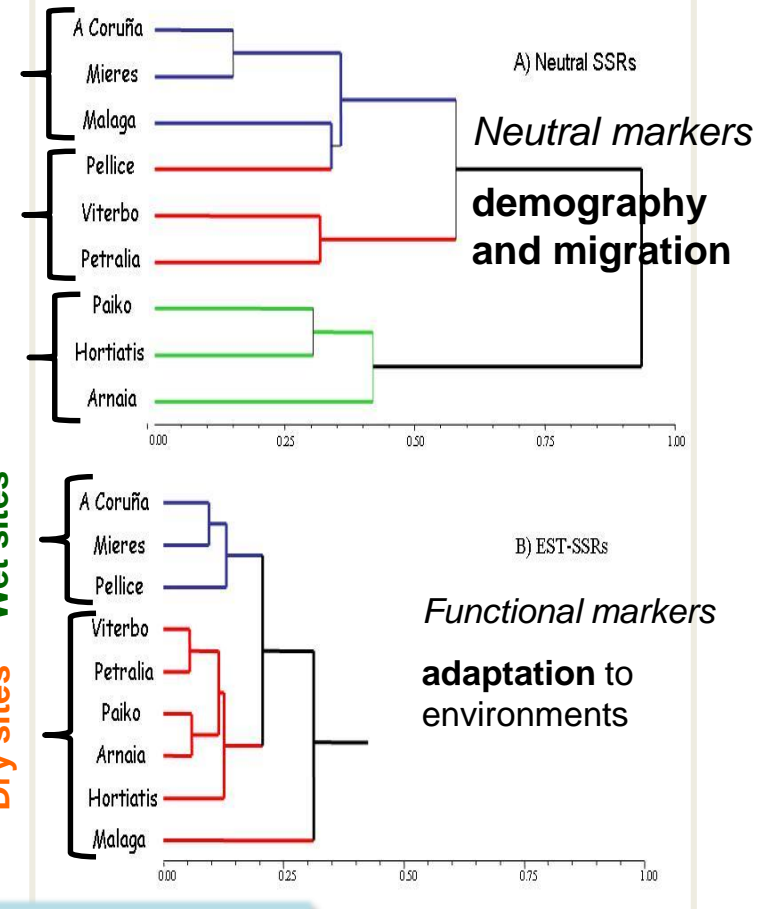
Spain

Italy

Greece

Wet sites

Dry sites



- Three main gene pools across *C. sativa* natural range
- Different population genetic structuring: history vs. adaptation

A common parameter: Genetic Variability

Juglans regia



Wide geographic range and high adaptive potential

Natural and human-mediated evolutionary history

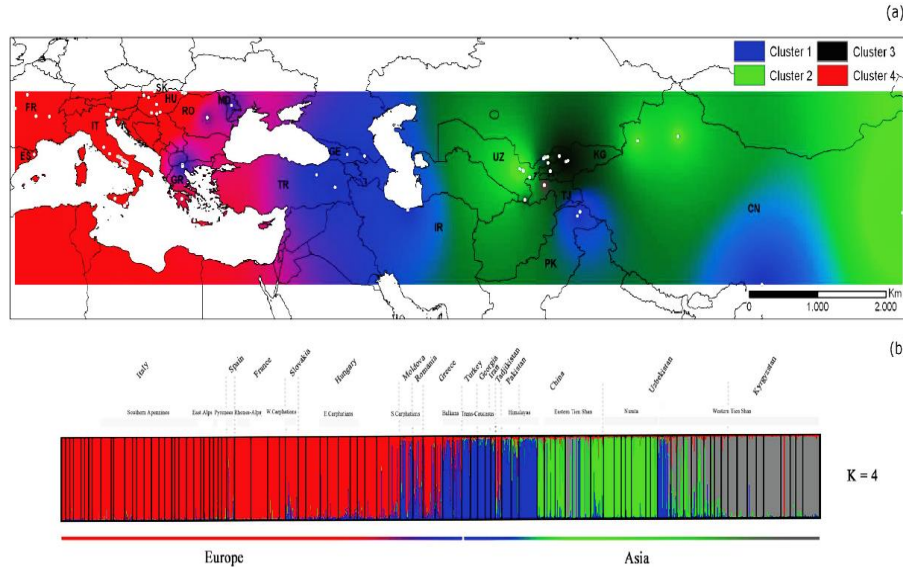


Fig 2. Spatial genetic structure of 91 walnut populations in Eurasia. Population structure inference for 91 walnut populations by Bayesian assignment using STRUCTURE for $K = 4$. Synthetic map of Inverse Distance Weighted (IDW) interpolations of the estimated mean population membership values (\hat{Q}) (a) and bar plot showing assignment probabilities of individuals to K clusters (b). Abbreviations: CN = China, UZ = Uzbekistan, KG = Kyrgyzstan, TJ = Tajikistan, PK = Pakistan, IR = Iran, GE = Georgia, TR = Turkey, MD = Moldova, RO = Romania, HU = Hungary, SK = Slovakia, GR = Greece, IT = Italy, FR = France, ES = Spain.

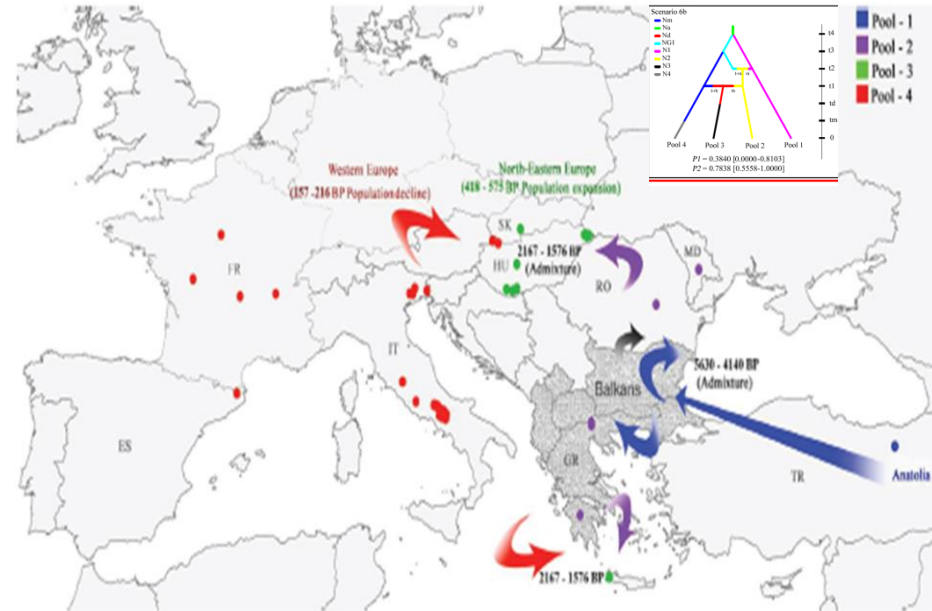
Tree Genetics & Genomes
DOI 10.1007/s11295-011-40740-2

ORIGINAL PAPER

Landscape genetics of Persian walnut (*Juglans regia* L.) across its Asian range

Paola Pollegioni¹ • Keith E. Woeste² • Francesca Chiochini¹ • Irene Olimpieri¹ • Virginia Tortolano¹ • Jo Clark³ • Gabriel E. Hemery² • Sergio Mapelli¹ • Maria Emilia Malvolti¹

Acta Hort. 1190. ISHS 2018. DOI 10.17660/ActaHortic.2018.1190.5
Proc. Int. Symp. on the Role of Plant Genetic Resources in Reclaiming
Lands and Environment Deteriorated by Human and Natural Actions
Eds.: A. Gharaghani and M. Khosh-Khui



Human-mediated dispersal routes of walnut during the Late Holocene as inferred by approximate Bayesian computation. Arrows represent the relationships between population pools used in DIYABC analysis (Pool 1, Pool 2, Pool 3, Pool 4) as inferred from stage 2, scenario 6b.

PLOS ONE

PLOS ONE

RESEARCH ARTICLE

Ancient Humans Influenced the Current Spatial Genetic Structure of Common Walnut Populations in Asia

Paola Pollegioni^{1*}, Keith E. Woeste², Francesca Chiochini¹, Stefano Del Lungo³, Irene Olimpieri¹, Virginia Tortolano¹, Jo Clark³, Gabriel E. Hemery², Sergio Mapelli¹, Maria Emilia Malvolti¹

RESEARCH ARTICLE

Rethinking the history of common walnut (*Juglans regia* L.) in Europe: Its origins and human interactions

Paola Pollegioni^{1,2*}, Keith Woeste³, Francesca Chiochini¹, Stefano Del Lungo³, Marco Cioffi¹, Irene Olimpieri¹, Virginia Tortolano¹, Jo Clark³, Gabriel E. Hemery², Sergio Mapelli¹, Maria Emilia Malvolti¹



Bud flush in the *P. alba* common garden study at Viterbo – Italy

Timing of **bud flush** and **bud set** represent a critical ecological and evolutionary tradeoff between survival and growth.

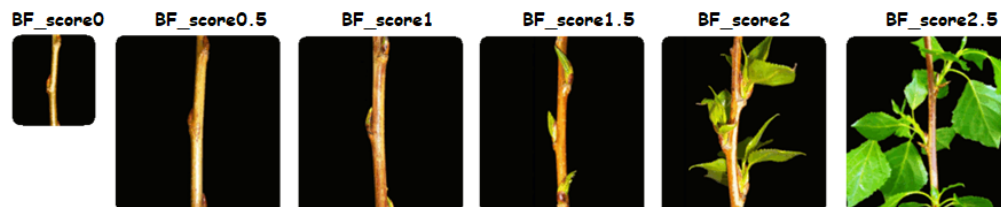


Figure 2. High-resolution scoring system for bud flush process in *Populus nigra*. Pictures were taken from individuals of *P. nigra* (Ludovisi, 2014). The scoring system for bud flush process was adapted and modified from Castellani et al. (1967).

Bud flush: depends mainly on the cumulative **temperature** sum in spring

Bud set: day length is the critical signal for **growth cessation** even if the timing of bud formation is also influenced by other factors such as:

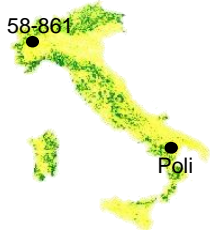
- temperature
- temperature × photoperiod
- nutrition
- drought
- biotic stresses.



Figure 3. High-resolution scoring system for bud set process in *Populus nigra*. Pictures were taken from individuals of *P. nigra* (Fabbrini et al., 2012). The scoring system for bud set process was adapted and modified from Rohde et al. (2011).



F₁ *Populus nigra* family (POP5)



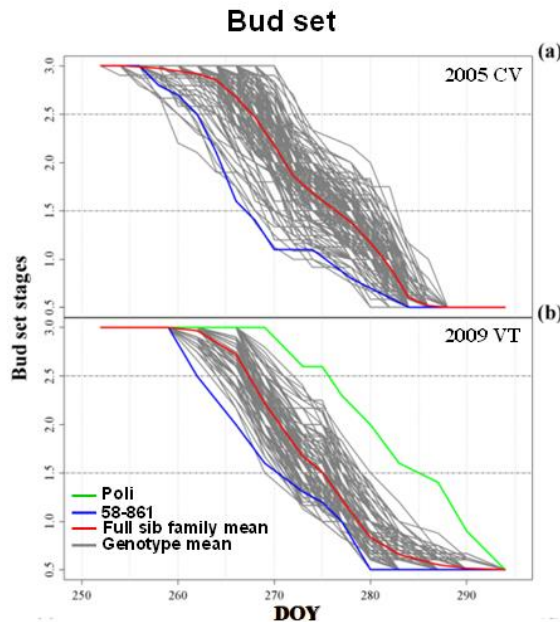
Segregation confirmed important additive effects inherited from 58-861 and Poli parents
→ a favourable situation to identify QTL

Fabbrini et al. *BMC Plant Biology* 2012, **12**:47
<http://www.biomedcentral.com/1471-2229/12/47>

RESEARCH ARTICLE

Open Access

Phenotypic plasticity, QTL mapping and genomic characterization of bud set in black poplar

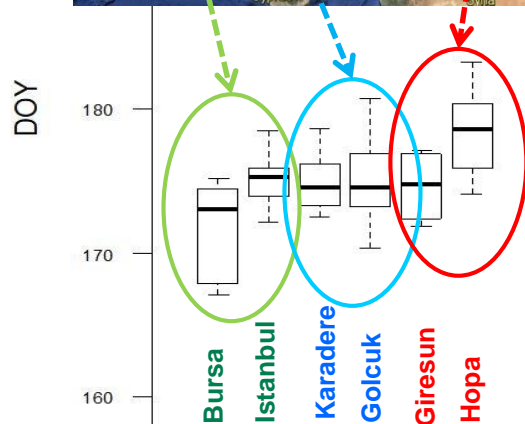
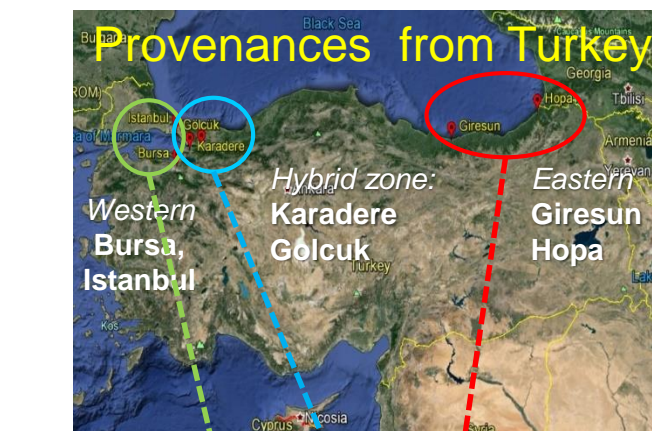


- ✓ High phenotypic variation in F₁ family (range 20 days) and moderate heritability ($H^2 = 0.45$).
- ✓ Considerable genotype × environment (G × E) interaction in all phenological stages, low temperatures influence the sensitivity of the most plastic genotypes.

Adaptive variation: floral phenology in chestnut



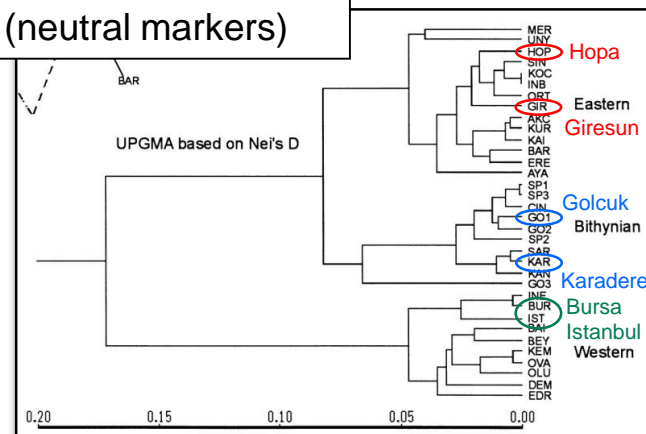
Common garden studies on natural germplasm



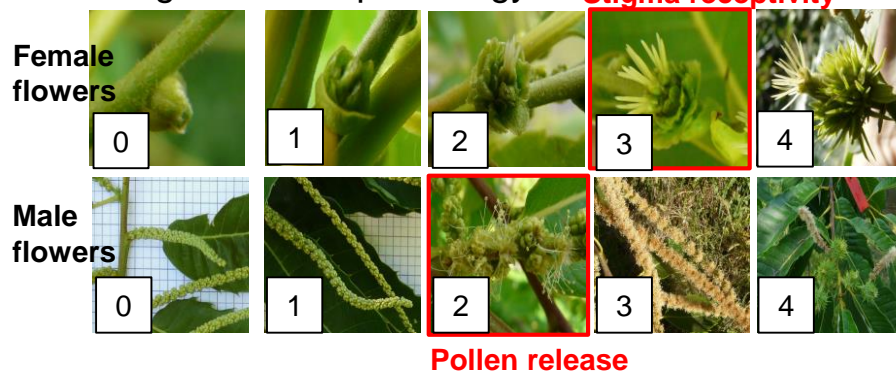
Date of pollen release
in common garden trial
CNR IBAF



Genetic distance (neutral markers)



Stages of floral phenology



- Phenology variation consistent with geographic and genetic distance: western provenances express earlier flowering than eastern provenances

Adaptive variation: floral phenology in walnut

Juglans regia



Genetic improvement and breeding

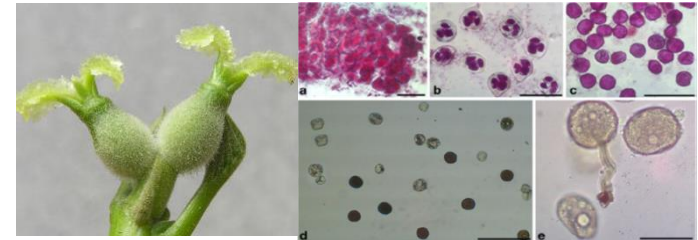
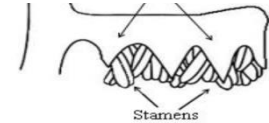
(Twenty years of cooperation with CREA FL and CD)

Tree Genetics & Genomes (2013) 9:291–305
DOI 10.1007/s11295-012-0555-y

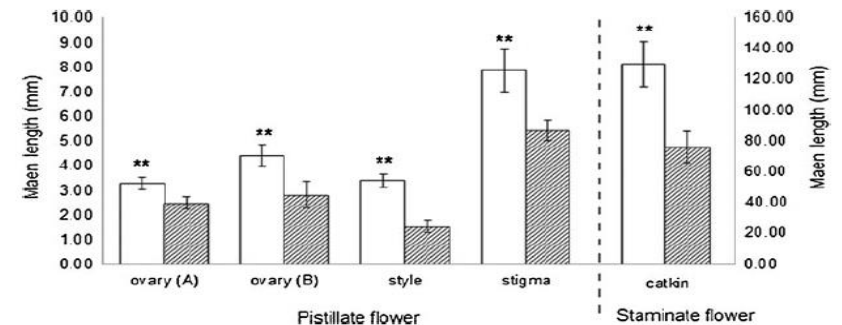
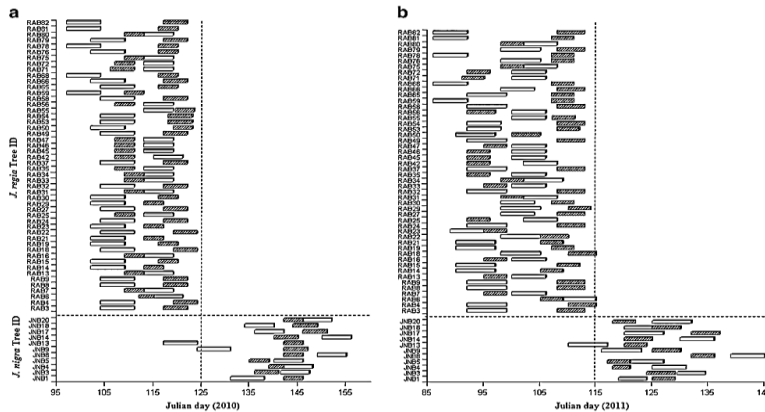
ORIGINAL PAPER

Barriers to interspecific hybridization between *Juglans nigra* L. and *J. regia* L species

Paola Pollegioni · Irene Olimpieri · Keith E. Woeste ·
Giovanni De Simoni · Maria Gras · Maria E. Malvolti



Tree Genetics & Genomes



Flowering period of female (white bars) and male (dashed bars) inflorescences for 11 *Juglans nigra* (JNB1-JNB20) and 50 *Juglans regia* (RAB3-RAB82) adult trees in 2010 (a) and 2011 (b).

Morphological characters of receptive pistillate flowers and mature catkins collected from 11 *Juglans nigra* (empty square) and 50 *Juglans regia* (filled square) adult trees ovary size (mean length of equatorial diameter A and polar diameter B \pm SD), mean length of style (\pm SD), stigma (\pm SD), and catkin (\pm SD). Statistical significance was tested by Student's t test with**p<0.0001



Abiotic stress: response to salinity in *P. alba*

- *P. alba* provenances from contrasting sites
- Comparative experiments (ex situ)

Differential mechanisms control variability in salinity tolerance

- Salt exclusion in roots
- Salt compartmentalization in cells
- Plasticity of leaf micromorphology
- Stomatal regulation
- Transcriptome regulation

Candidate genes

Tonoplast proton pumps
Heat Shock Proteins (HSP70)
Trehalose-6-phosphate synthase

- Natural *P. alba* genetic resources to improve salinity tolerance
- Selection of functional molecular markers to aid breeding



Tree Physiology 31, 1335–1355
doi:10.1093/treephys/tpo083

Research paper: Part of a special section on poplars and the environment

Comparative study of transcriptional and physiological responses to salinity stress in two contrasting *Populus alba* L. genotypes

Isacco Beritognolo^{1,2}, Antoine Harfouche¹, Federico Brilli³, Gianluca Prosperini⁴, Muriel Gaudet¹, Mikael Brosché⁵, Francesco Salani¹, Elena Kuzminsky¹, Petri Auvinen⁶, Lars Paulin⁶, Jaakko Kangasjärvi⁵, Francesco Loreto⁷, Riccardo Valentini¹, Giuseppe Scarascia Mugnozza⁸ and Maurizio Sabatti^{1,9}

Environmental and Experimental Botany 66 (2009) 381–388



ELSEVIER

Contents lists available at ScienceDirect

Environmental and Experimental Botany

journal homepage: www.elsevier.com/locate/envexpbot



Leaf morphological plasticity and stomatal conductance in three *Populus alba* L. genotypes subjected to salt stress

Grazia Abbruzzese^a, Isacco Beritognolo^a, Rosario Muleo^b, Moica Piazzai^a, Maurizio Sabatti^a, Giuseppe Scarascia Mugnozza^a, Elena Kuzminsky^{a,*}

Trees (2007) 21:465–477
DOI 10.1007/s00468-007-0139-x

ORIGINAL PAPER

Functional characterisation of three Italian *Populus alba* L. genotypes under salinity stress

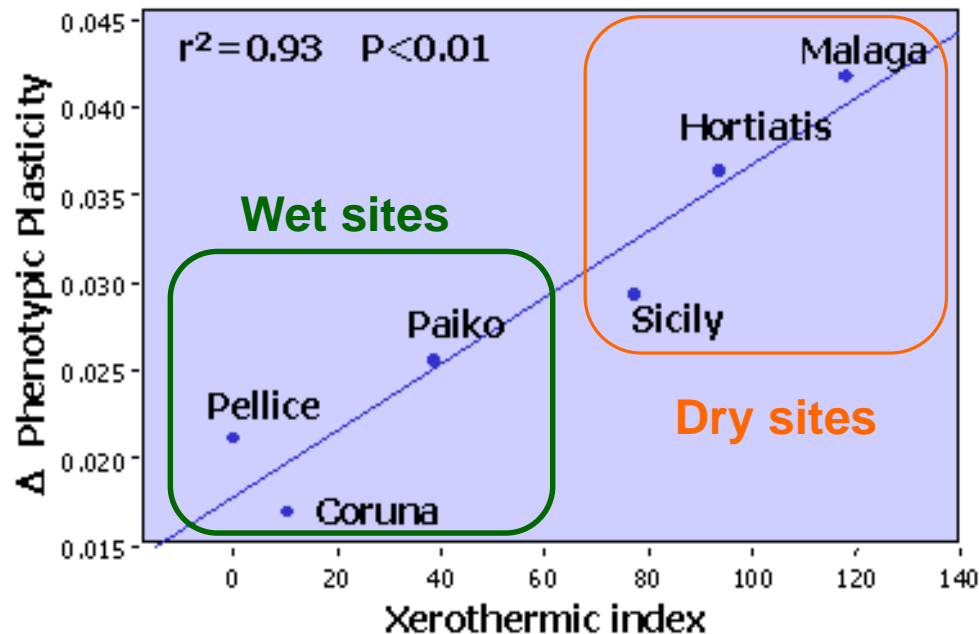
Isacco Beritognolo · Moica Piazzai · Simona Benucci · Elena Kuzminsky · Maurizio Sabatti · Giuseppe Scarascia Mugnozza · Rosario Muleo

Abiotic stress: drought tolerance in *C. sativa*

Castanea sativa



Phenotypic plasticity of Water Use Efficiency correlates with Xerothermic Index



Genetic variation in carbon isotope discrimination in six European populations of *Castanea sativa* Mill. originating from contrasting localities

M. LAUTERI,* A. PLIURA,† M. C. MONTEVERDI,* E. BRUGNOLI,* F. VILLANI* & G. ERIKSSON‡

*Consiglio Nazionale delle Ricerche, Istituto di Biologia Agroambientale e Forestale, Porano, Italy

†Department of Forest Genetics and Reforestation, Lithuanian Forest Research Institute, Girionys, Lithuania

‡Department of Forest Genetics, SLU, Uppsala, Sweden

J. EVOL. BIOL. 17 (2004) 1286–1296

- Δ $^{13}\text{C} / ^{12}\text{C}$ isotope discrimination of leaves

- Provenances from dry sites express a higher phenotypic plasticity in response to drought
- Phenotypic plasticity contributes to environmental adaptation

Biotic stress: chestnut gall wasp

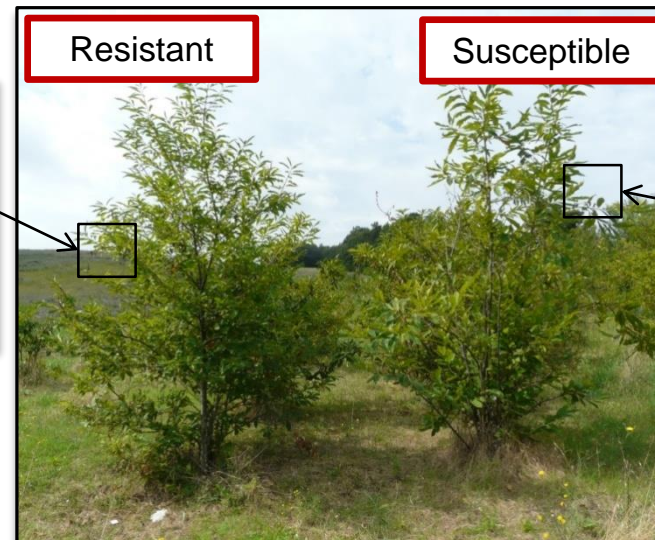
Castanea sativa



Susceptibility to *Dryocosmus kuriphylus* in common garden provenance trial (CNR IBAF)



Infestation of buds 2016



Provenance	Sampled plants	Resistant plants	% of resistant
Coruna (SPAIN) wet	19	0	0
Malaga (SPAIN) dry	21	1	4.8
Pellice (ITALY) wet	17	0	0
Petralia Sottana (ITALY) dry	20	0	0
Paiko (GREECE) wet	20	3	15
Hortiatis (GREECE) dry	20	6	30

- Greek provenances:
 - high percentage of resistant plants
 - Valuable genetic resources for research and breeding
- Genomic studies are in progress

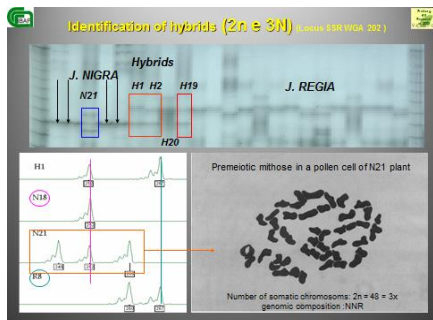
Infestation potential of *Dryocosmus kuriphylus* Yasumatsu, 1951 (Hymenoptera: Cynipidae) in different natural populations of *Castanea sativa* Miller: an experimental ex-situ test
Bombi, Fedi, Zapparoli, Cammarano, Guidolotti, Pallozzi, Gaudet, Mattioni, Cherubini, Beritognolo, Villani

International Journal of Pest Management. Accepted for publication

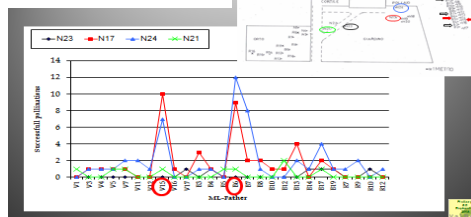
Genetic improvement for interspecific walnut hybrids

(cooperation with CREA)

Selection of hybrid walnut plants in common garden.
Identification of maternal hybridogenic trees, putative fathers, and hybrid progenies.



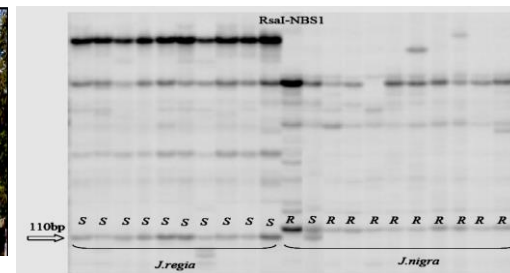
Reproductive success of male parents



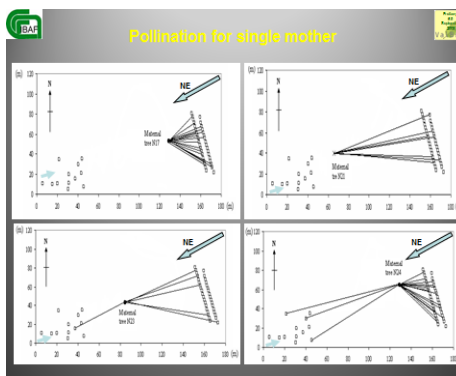
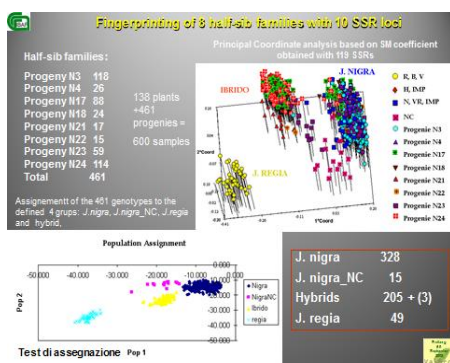
Selection of *J. nigra* and *J. regia* genotypes susceptible/tolerant to biotic stress (*Gnomonia leptostyla*) by neutral and functional markers



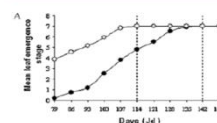
Juglandetum CREA



Functional molecular markers associated with Resistant or Resistant-analogs genes used for the early selection of genotypes tolerant/susceptible to diseases. (NBS profiling approach)



Avoidance by late flushing



Pathogenicity test in vivo



Controlled interspecific crosses

Mol Breeding (2009) 24:321–335
DOI 10.1007/s11032-009-9294-7

Retrospective identification of hybridogenic walnut plants by SSR fingerprinting and parentage analysis



Mechanisms governing the responses to anthracnose pathogen in *Juglans* spp.

P. Pollegioni^{a,*}, G. Van der Linden^b, A. Belisario^c, M. Gras^d, N. Anselmi^e, I. Olimpieri^a, L. Luongo^c, A. Santini^g, F. Turco^g, G. Scarascia Mugnozza^f, M.F. Malvoltri^a



Phenotypic characterization of productive traits (fruit quality) in walnut

Improvement of local varieties and ecotypes: molecular, morphological, biochemical markers

Bioremediation, Biodiversity and Bioavailability ©2010 Global Science Books



Juglans regia Provenance Research by Molecular, Morphological and Biochemical Markers: A Case Study in Italy

Maria Emilia Malvolti¹ • Paola Pollegioni¹ • Alcide Bertani¹ • Sergio Mapelli^{2*}



Ecotipi/varietà locali poco valorizzati e a rischio estinzione

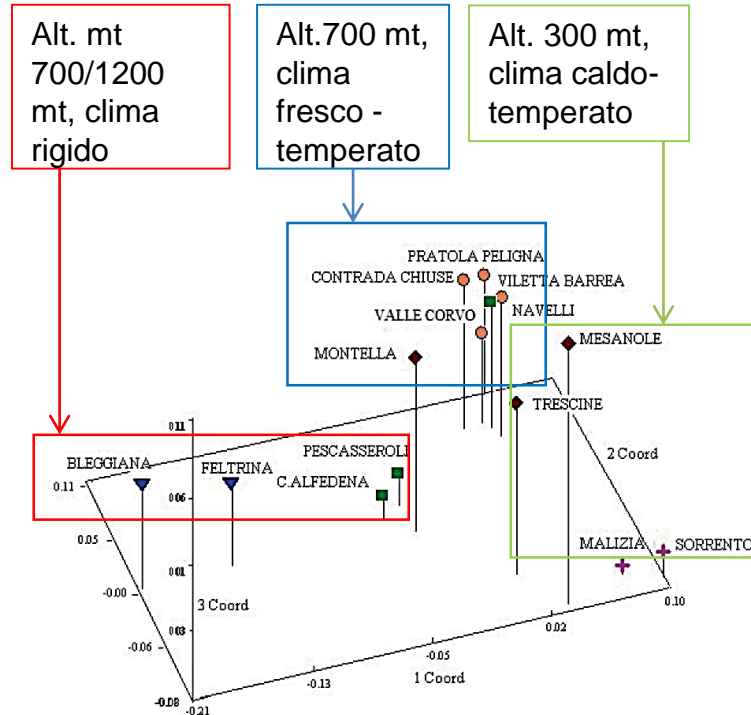
Pericolo di erosione genetica

Caratteristiche di rusticità, adatto per riforestazione e sistemazione idrogeologica

Mancanza di strategie di valorizzazione

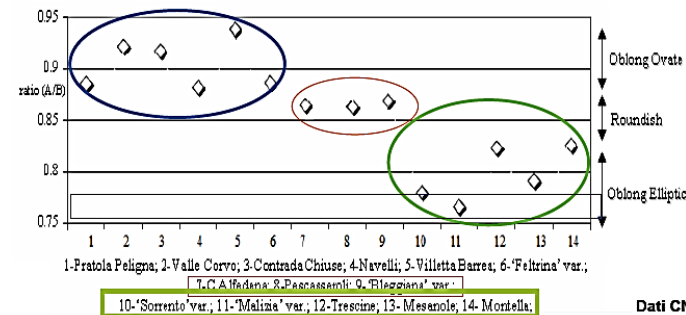
Produzioni e superfici in continuo calo

- 1) Individuazione di una macroarea-studio
- 2) Analisi della nocicoltura di montagna
- 3) Uso di metodologie analitiche per il controllo e la caratterizzazione
- 4) Costruzione di strategie territoriali per la valorizzazione



Analisi delle coordinate principali basata su 10 SSR

Varietà conosciute di confronto: Sorrento e Malizia (Campania), Bleggiana (Trentino) e Feltrina (Veneto)

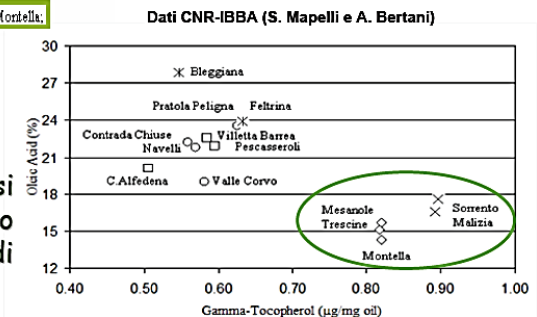


RISULTATI

Forma frutto: rapporto diametro Equatoriale / Polare

Regione Campania
Regione Abruzzo
Regione Molise

Tocoferolo, acidi grassi poliinsaturi ($\omega 3$; $\omega 6$) ed acido oleico contenuti nei frutti di noce.



Towards more collaborative *multidisciplinary research activities* for tree breeding in Italy?

- Agro-forestry species are multipurpose and ecologically relevant, but with low economic impact
- Tree species require permanent infrastructures for long-term research and breeding
- Characterised tree genetic resources available in collections are valuable for breeding but underexploited
- Part of the novel scientific knowledge not yet used in innovative Italian tree breeding programmes
- How to improve the system?



Thanks for your attention!