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How can we harness the genetic diversity for genetic improvement, taking into account reproductive traits acquired through the evolution of the citrus species?

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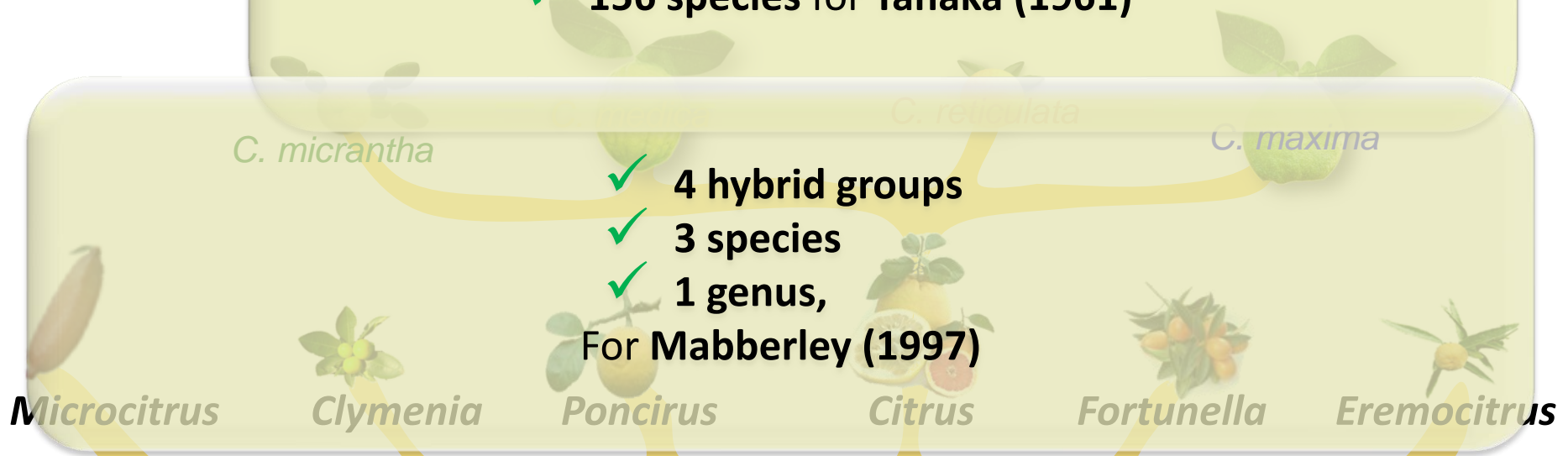
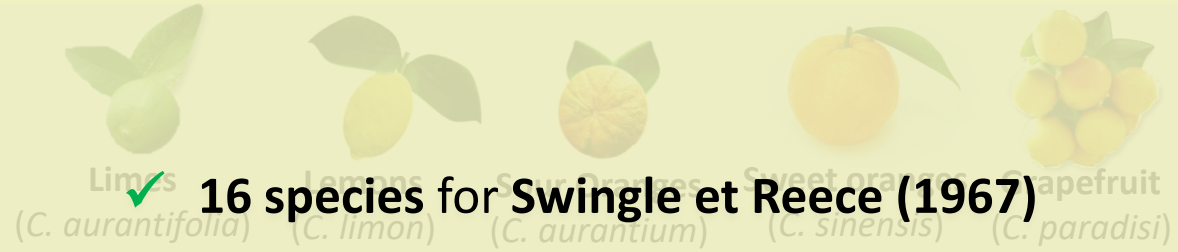
How can we harness the genetic diversity for genetic improvement, taking into account reproductive traits acquired through the evolution of the citrus species?

François Luro

INRAe-Cirad, Agap Corse, France



How to classify citrus species



True citrus
(6 genus)

Sub tribes ***Citrinae***

Tribe ***Citreae***

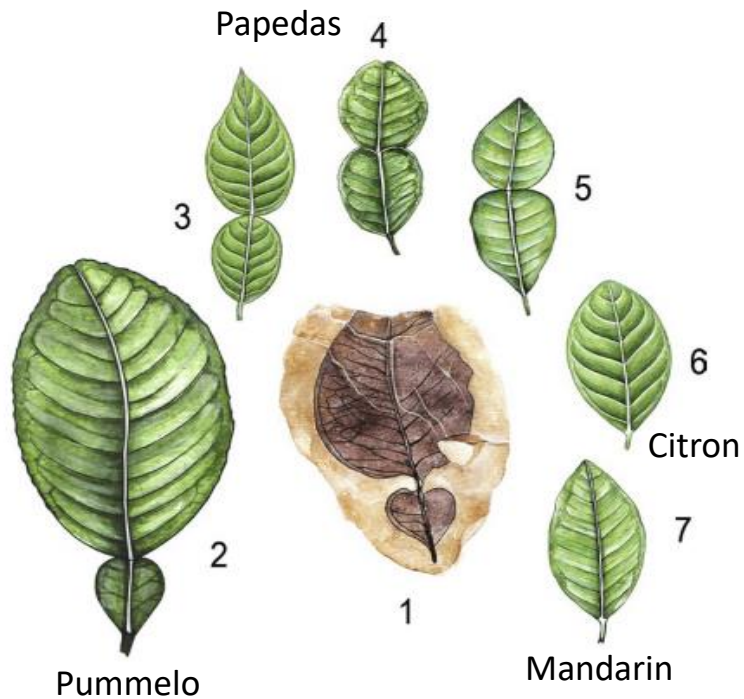
Sub Family ***Aurantioideae***

Family ***Rutaceae***

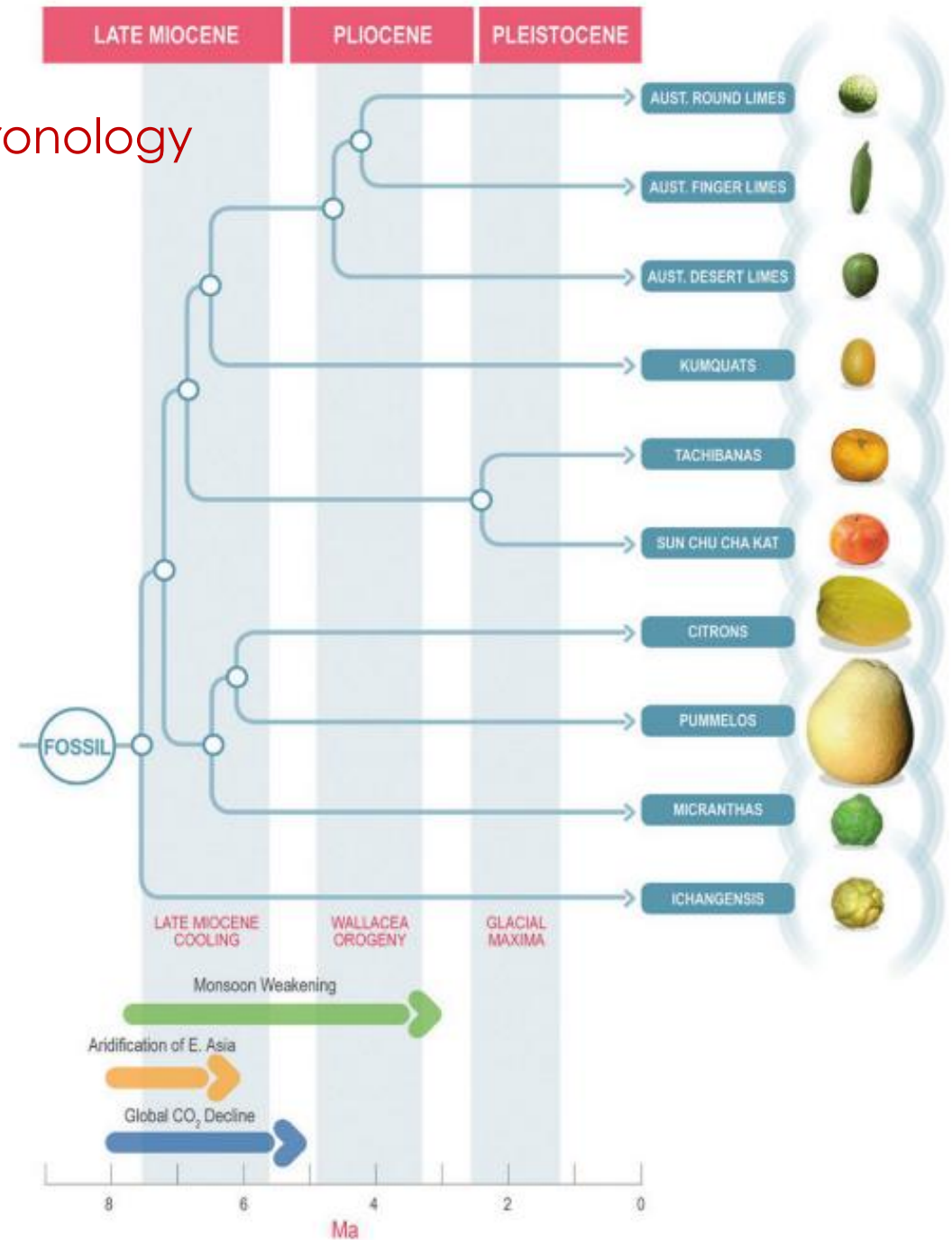
Evolution of citrus ancestors

Speciation chronology

Characteristics of a fossil leaf of *C. linczangensis* (1) compared with those of present-day species

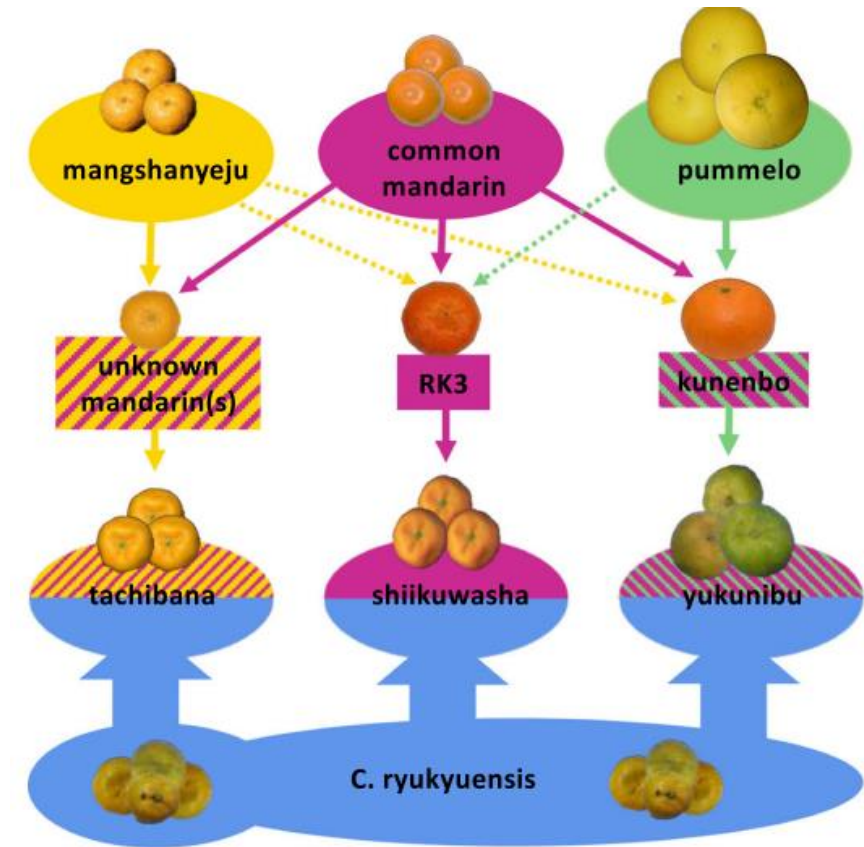
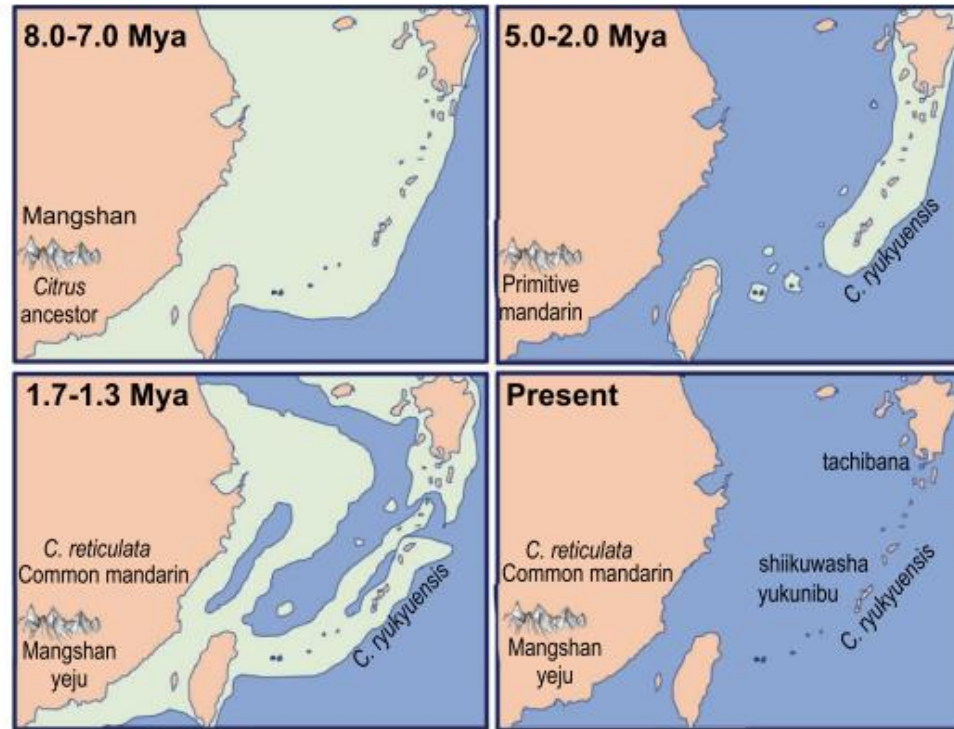
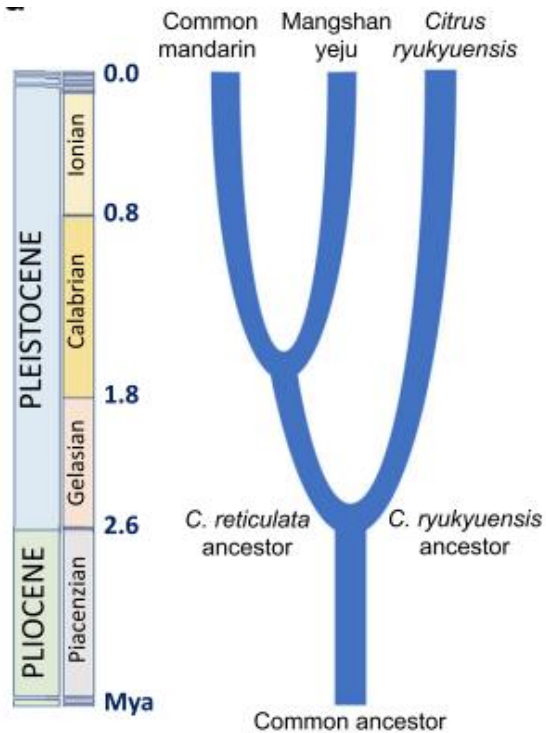


Xie et al. 2013



Mandarin evolution: allopatric evolution and interspecific crosses

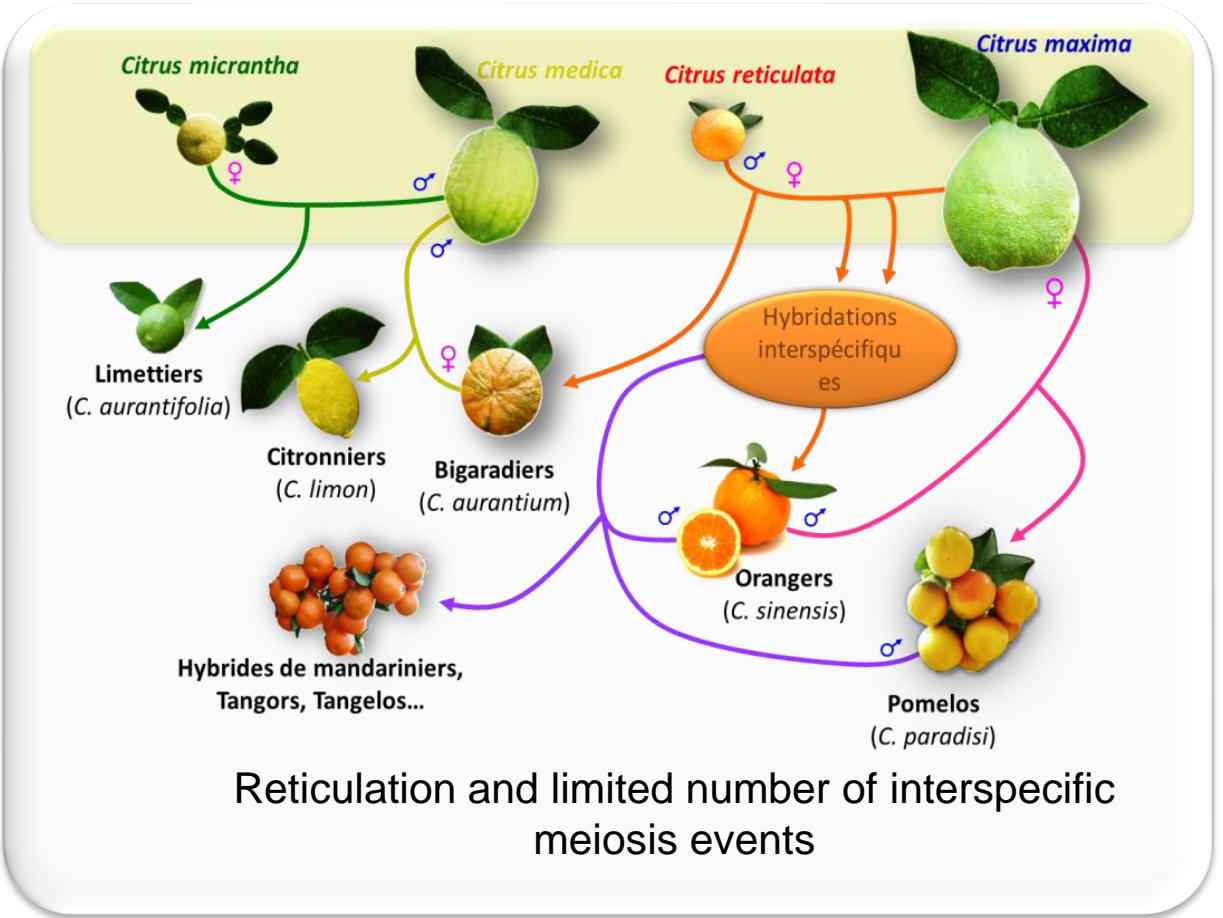
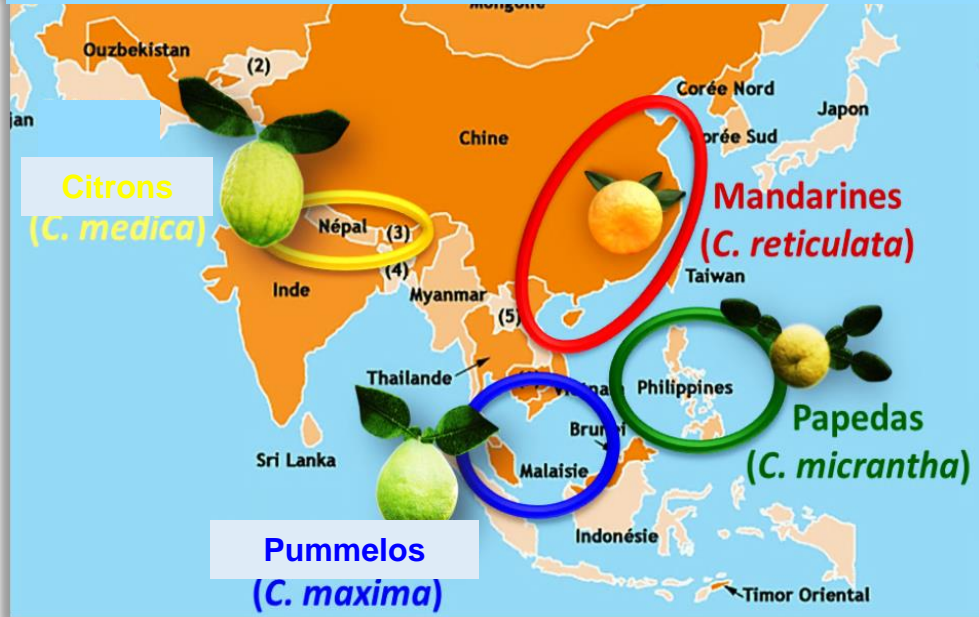
Chronogram of speciation and biogeography of the East Asian mandarin tree in the Ryukyu arc and mainland Japan



Crosses between mandarin subpopulations and with *C. maxima* are the source of *C. reticulata* diversification

Reticulate evolution of citrus species

Differentiation between basic taxa
initial allopatric evolution



The genomes of cultivated *Citrus* are mosaics of large chromosome fragments from the basic taxa in frequent interspecific heterozygosity

Structures maintained by apomixis or horticultural vegetative propagation practices



Apomictic reproduction in citrus

Variation of development step between embryos

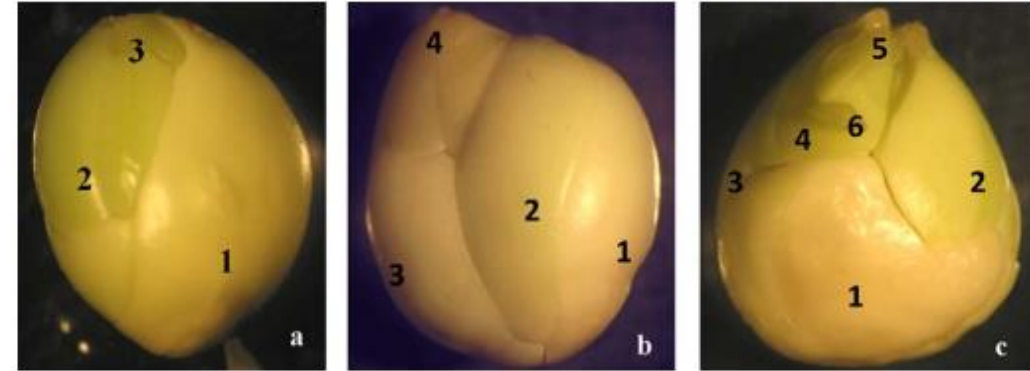
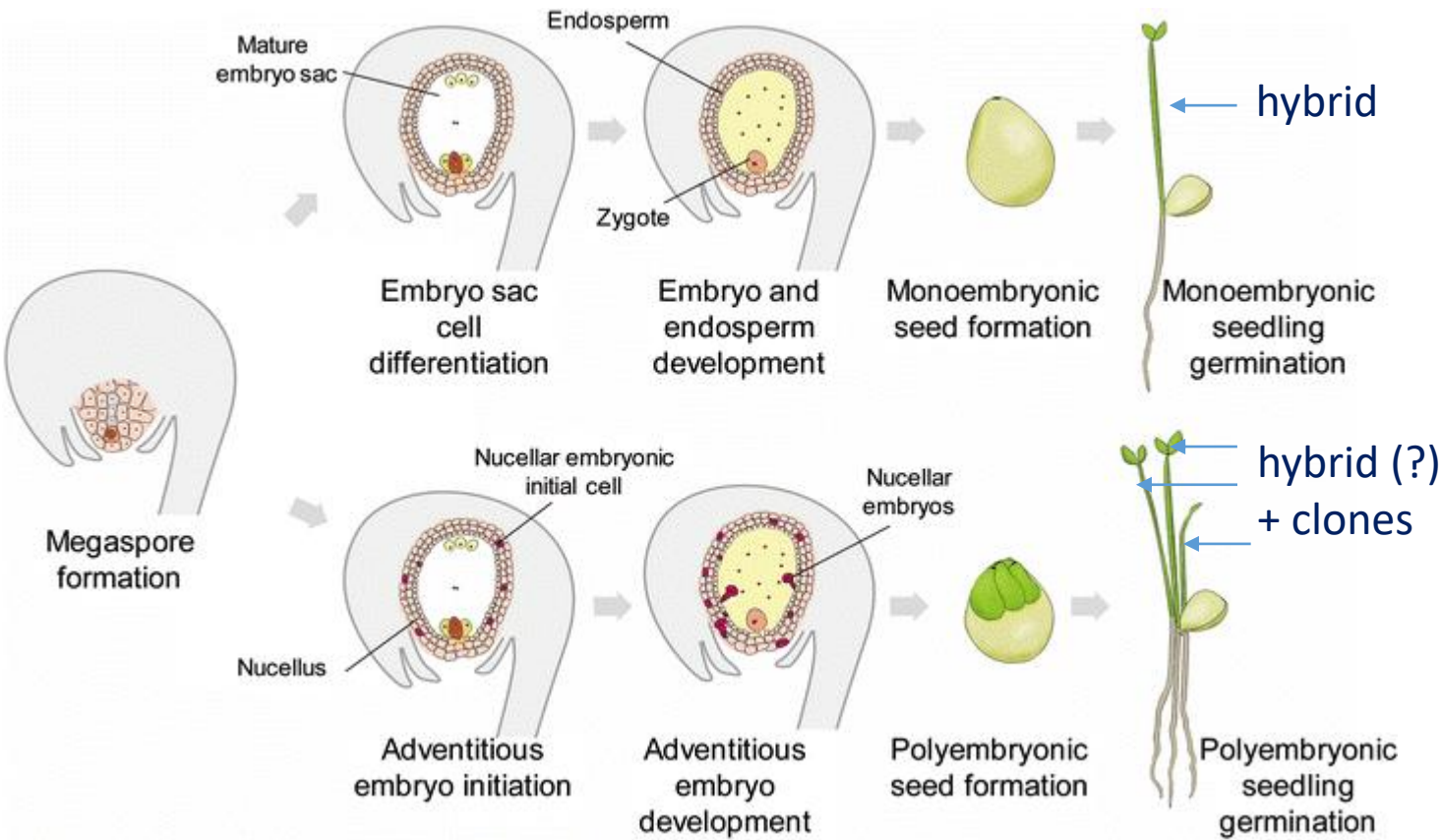


Fig. 4. Variation in biomass partitioning among embryos. (a) Triplet; (b) quadruplet; and (c) sextuplet.



Zygote formation is necessary for adventitious embryogenesis

Emergence

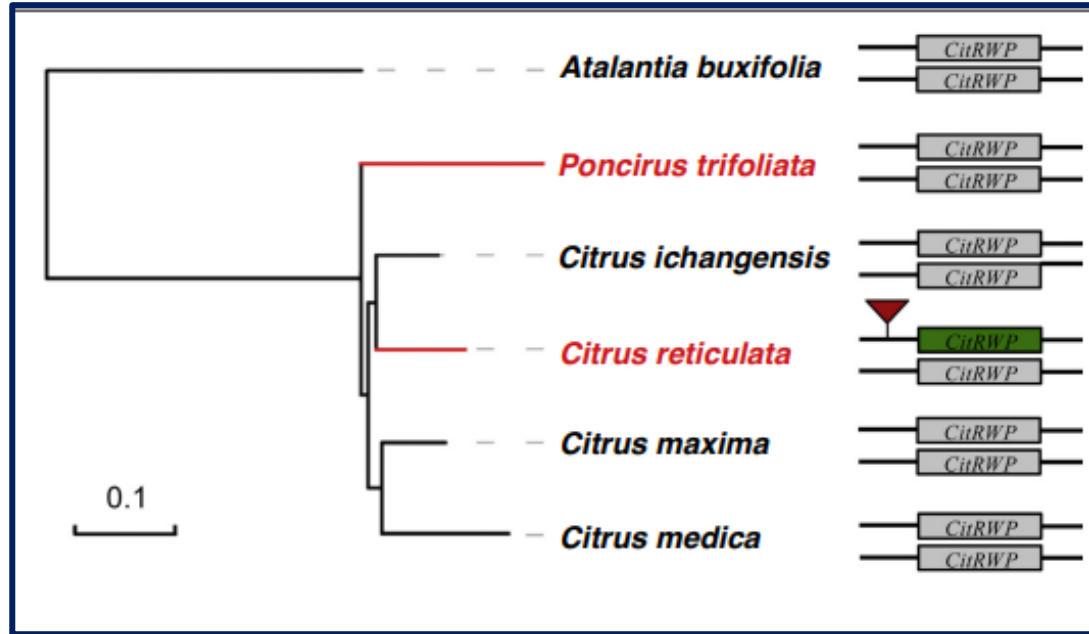
asynchrone

synchrone

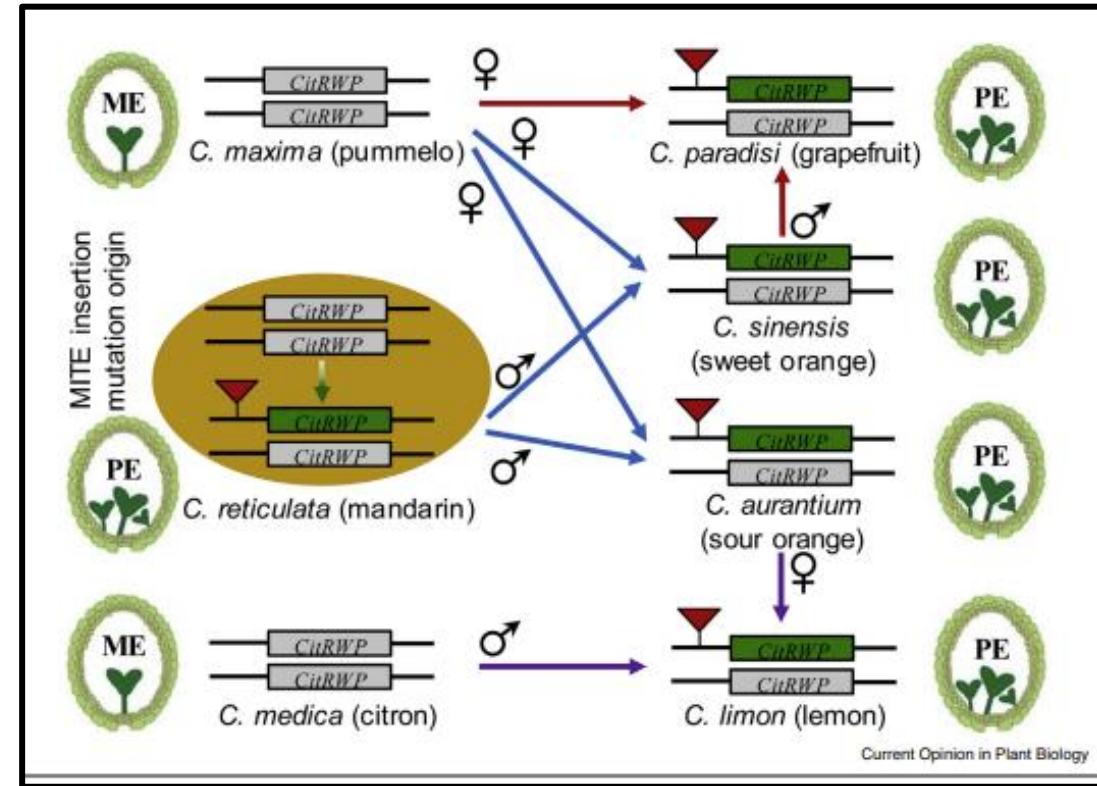
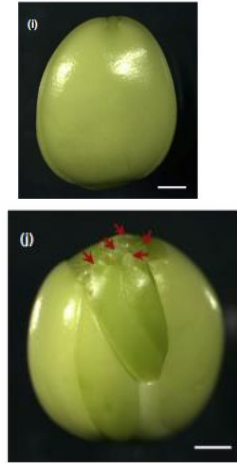


Competition between embryos

Origin and heredity of somatic embryogenesis



(Zhang et al. 2018)



- Nucellar embryo has appeared in mandarins after insertion of a transposable element (MITE) upstream of the *CitRWP* gene.
- The mechanism of nucellar embryony is unknown in *Poncirus trifoliata*

- All cultivated taxa have inherited the apomictic reproduction from mandarin.
- Apomixis contributed to the selection of descendants who inherited this trait

The citrus genome



A « small » genome (**360** Mpb. - **398** Mpb.) [Ollitrault *et al.* **1994**]



Arabidopsis thaliana **119** Mpb



Zea mays **5.000** Mbp



Populus trichocarpa **485** Mbp



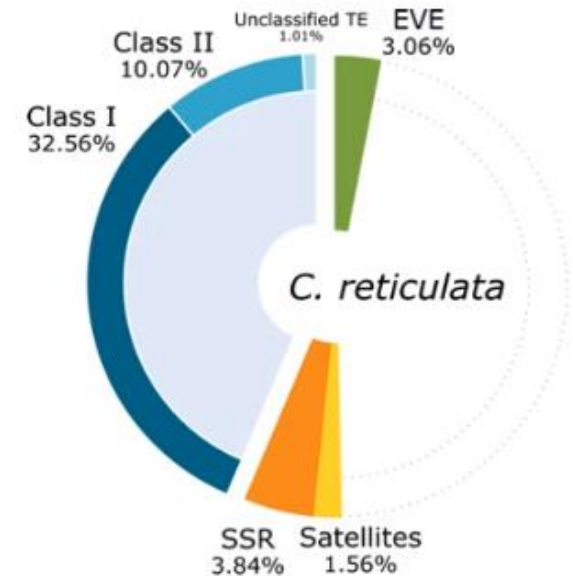
Triticum **17.000** Mbp



n = 9 chromosomes

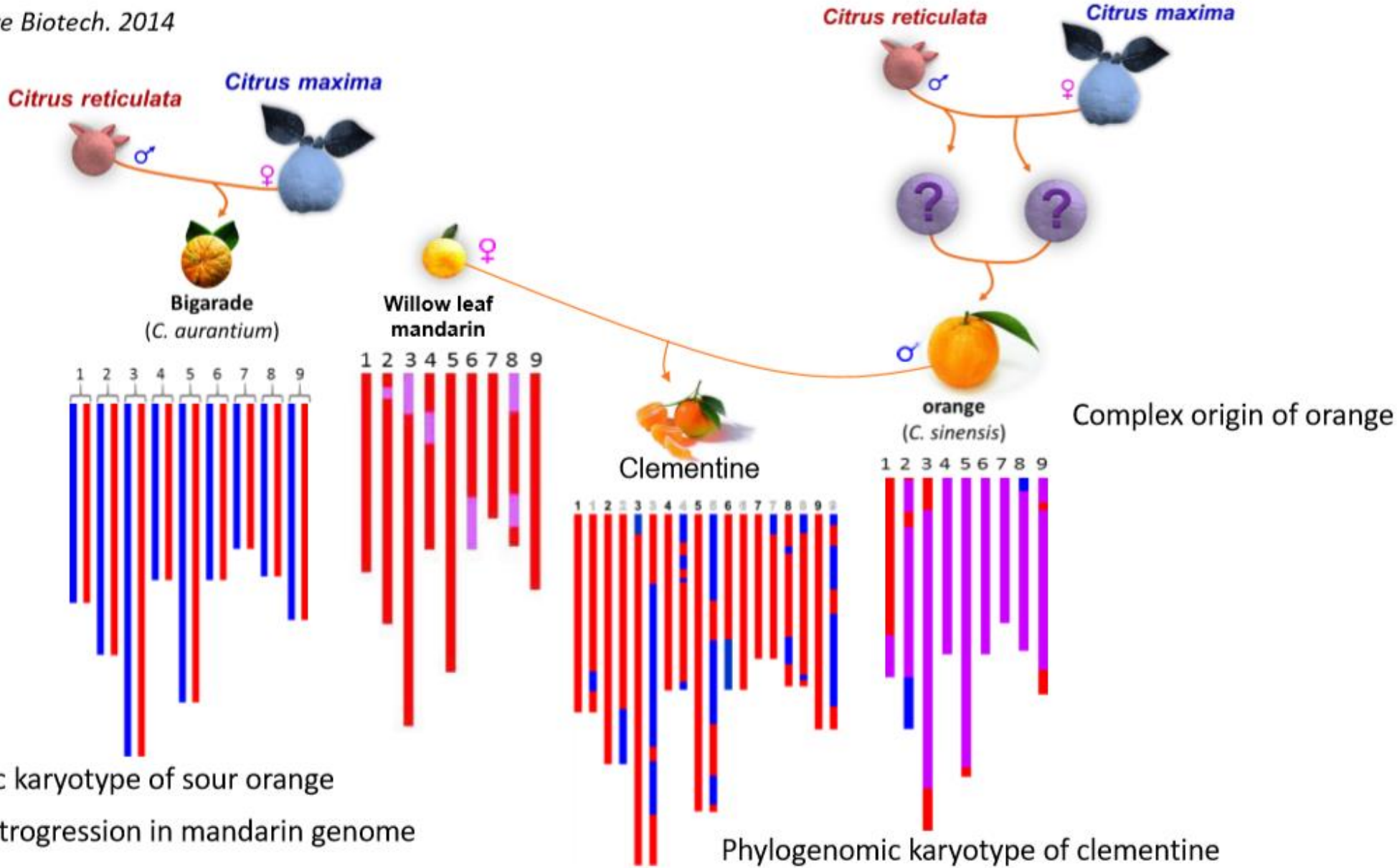
About 37 000 genes

Repeated sequences ~50% genome
with 42% of transposable elements



Genomic structures of cultivated citrus by Genome Sequencing (WGS)

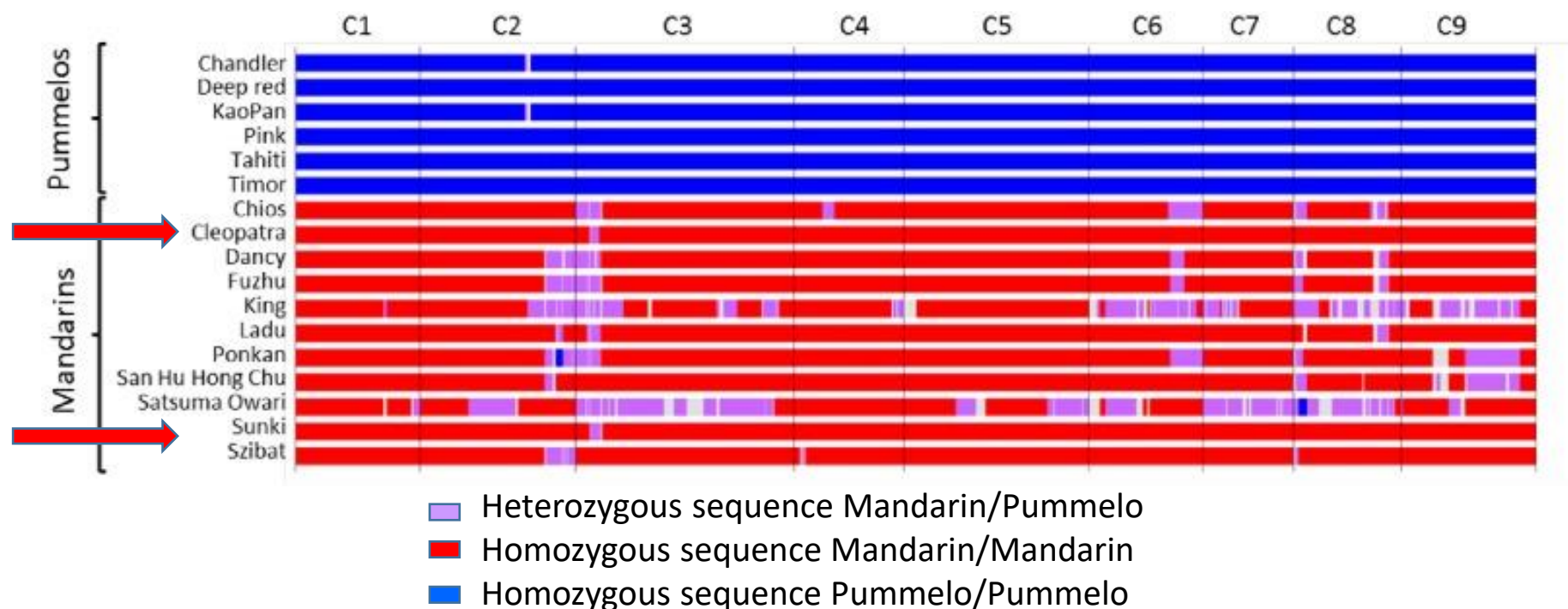
Wu et al. Nature Biotech. 2014



Introgression of pummelo into modern cultivars of mandarin genome

Are representatives of *C. reticulata* ancestral taxa existing ?

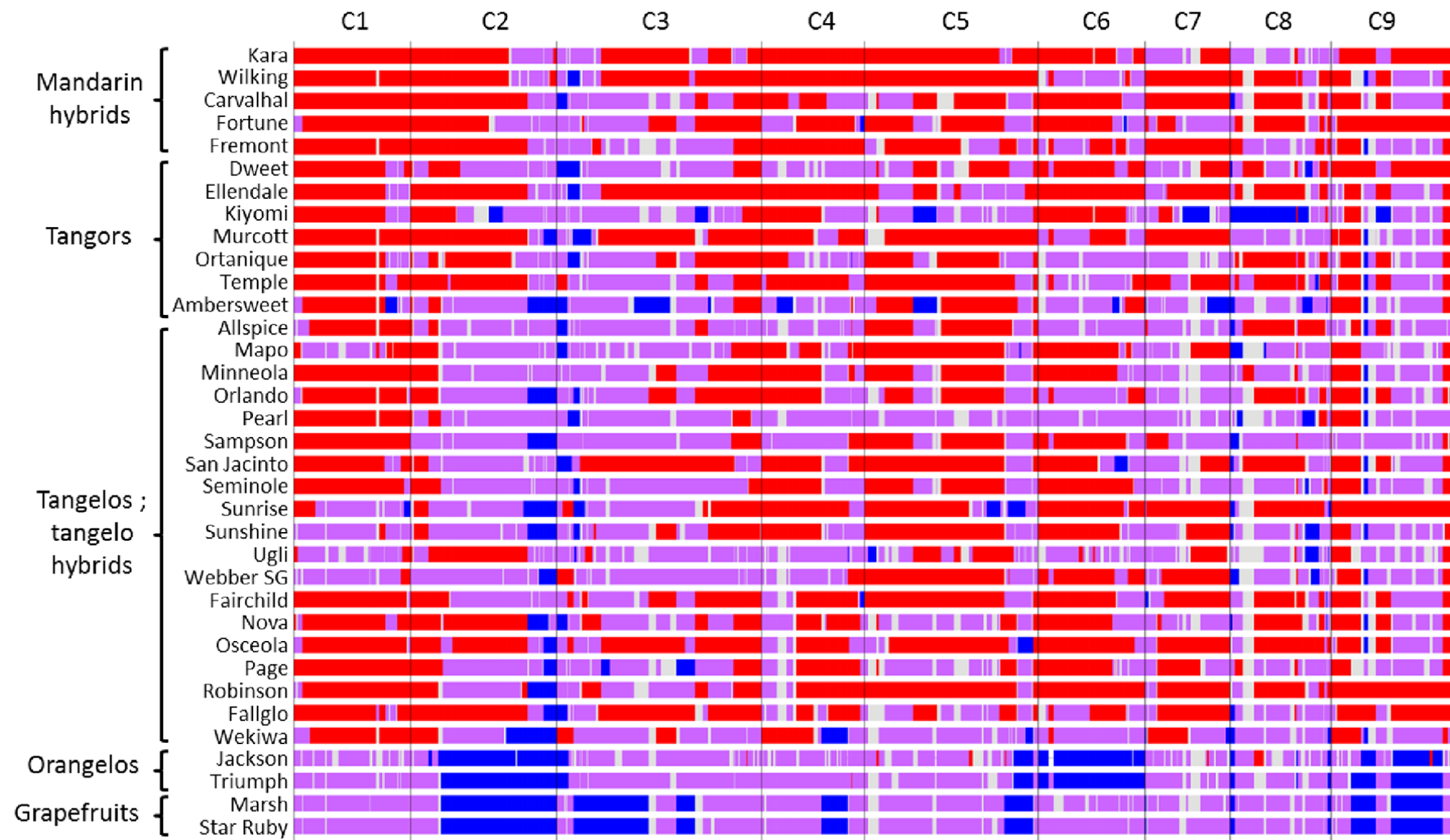
Phylogenomic karyotype based on GBS data of varieties used as references of pummelos and mandarins



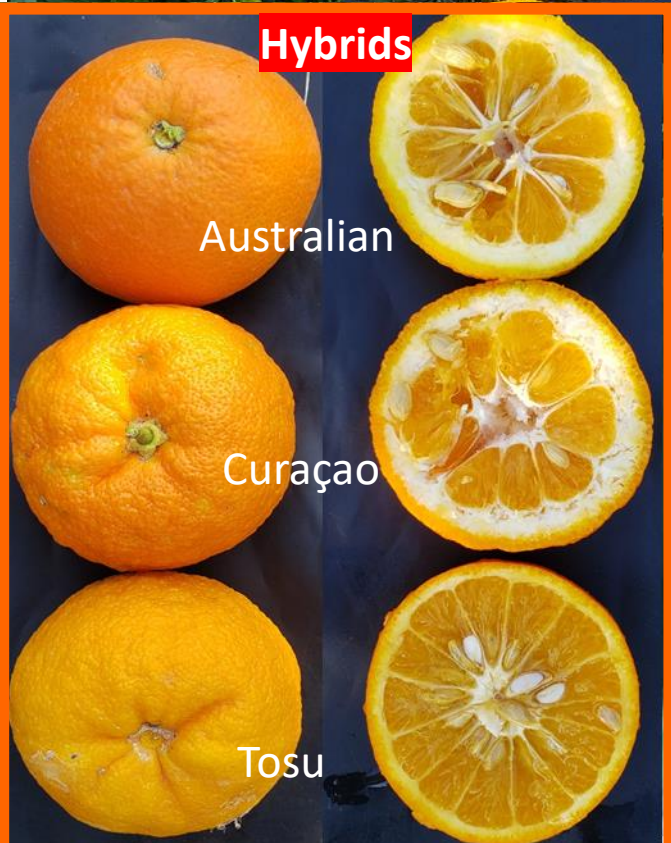
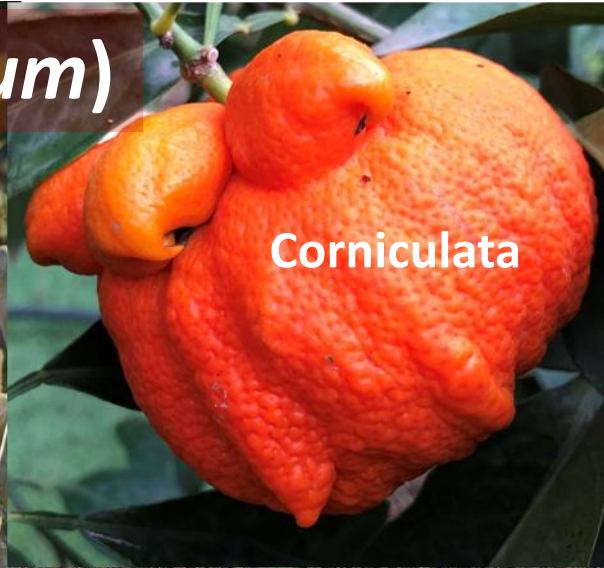
Cleopatra and Sunki are the closest to wild mandarins

Other mandarins share more or less introgressions of pummelo genome

Analysis of phylogenomic structures and interspecific recombination for a population derived from the *C. reticulata* / *C. maxima* gene pools...



Mutation as source of diversity in secondary species



Mutation as source of diversity in secondary species

Anthocyanin synthesis in bloody oranges is induced by an insertion of a transposable element in *Ruby* promoter region

The Plant Cell, Vol. 24: 1242–1255, March 2012, www.plantcell.org © 2012 American Society of Plant Biologists. All rights reserved.

Retrotransposons Control Fruit-Specific, Cold-Dependent Accumulation of Anthocyanins in Blood Oranges^{WUQ}

Eugenio Butelli,^a Concetta Licciardello,^b Yang Zhang,^a Jianjun Liu,^c Steve Mackay,^a Paul Bailey,^a Giuseppe Reforgiato-Recupero,^b and Cathie Martin^{a,1}

^aJohn Innes Centre, Norwich NR4 7UH, United Kingdom
^bCentro di Ricerca per l'Agrumicoltura e le Colture Mediterranee, 95024 Acireale, Italy
^cSichuan Academy of Agricultural Sciences, Chengdu City, Sichuan 610066, China

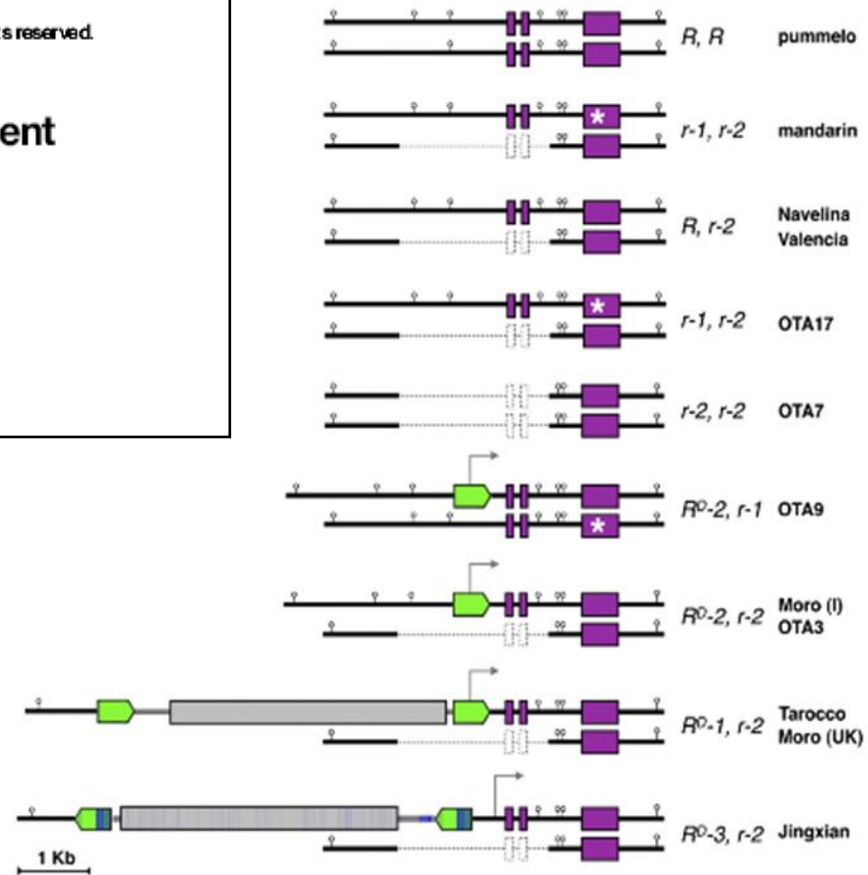
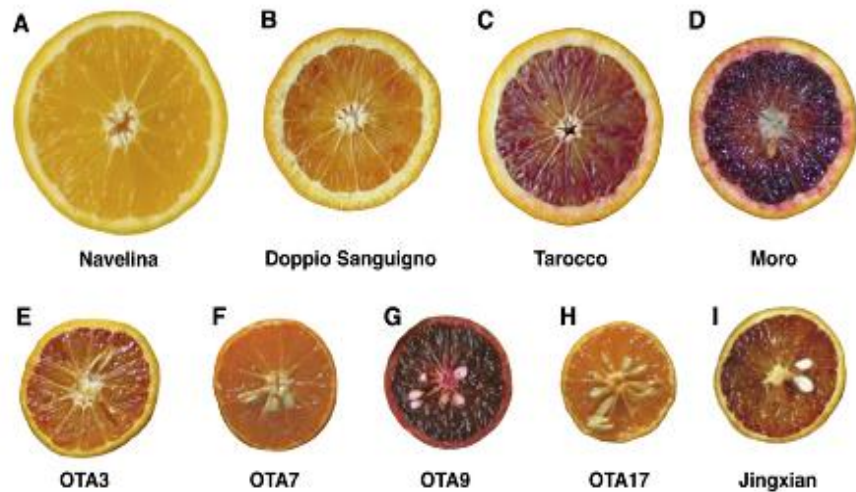
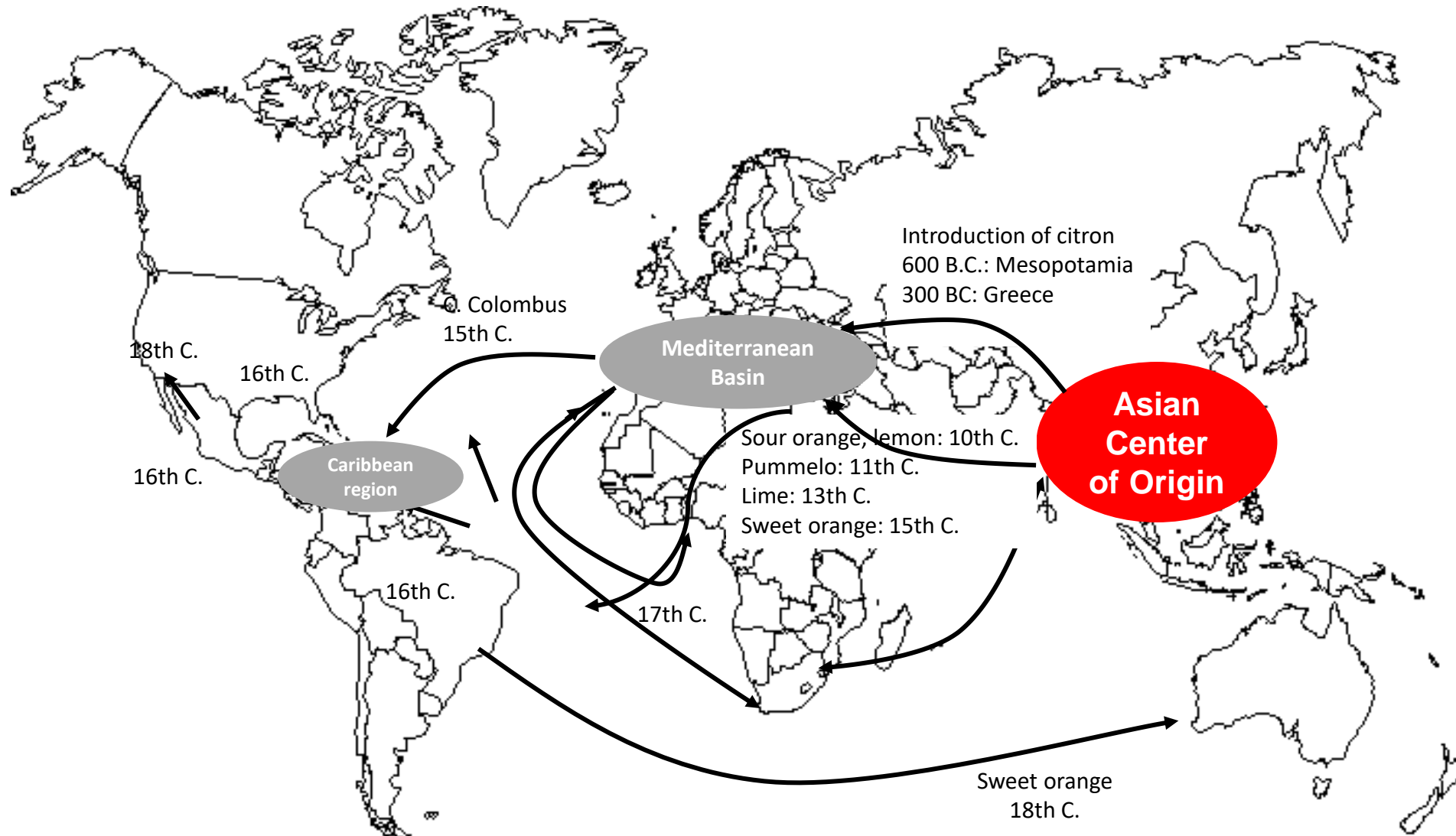


Figure 6. Maps of Structures of the *Ruby* Locus in the Different *Citrus* Species, Orange Accessions, and Hybrids.

Citrus origin and dispersion



Citrus diversity for genetic improvement



The objectives of breeding programs are:

- to diversify the panel of citrus cultivars,
- to extend the period of citrus production
- to produce new cultivars / rootstocks adapted to the environment and constraints

- How the citrus diversity can be mobilized ?
- Which main strategies for each kind of citrus ?
- How can breeding strategies be adapted to the characteristics of the species' biology ?

Constraints and advantages of conventional breeding

Reproduction

Self-incompatibility

backcross is hampered



Gametic sterility due to chromosomal aberrations
embryo sac or seed abortion, or gene (CMS)



Partial apomixis

Polyembryonic seeds give somatic embryos from maternal tissue (nucellus).
Unfit for progeny development (competition)



The delays for fruiting are long and limit the number of generations in a genealogical selection

Long juvenile phase (~6 years)



Overcrowding of adult progenies



~25 m²

Genetic control of gametic self-incompatibility

Mol Genet Genomics (2017) 292:325–341
DOI 10.1007/s00438-016-1279-8

Liang et al. 2020

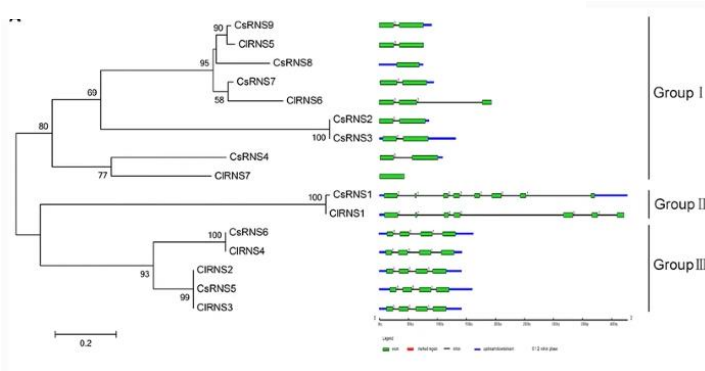


ORIGINAL ARTICLE

Genome-wide identification and functional analysis of S-RNase involved in the self-incompatibility of citrus

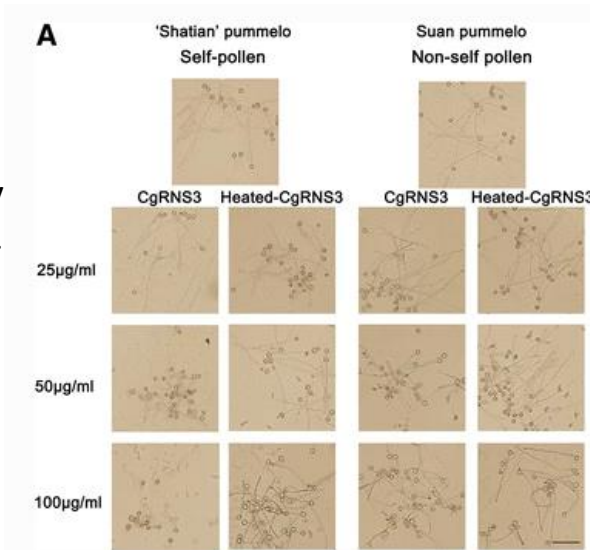
Mei Liang¹ · Wei Yang¹ · Shiyong Su¹ · Lili Fu¹ · Hualin Yi¹ · Chuanwu Chen² · Xiuxin Deng¹ · Lijun Chai¹

Homologous RNase genes from genome data base



CsRNS3 gene expressed only in pistil

CgRNS3 protein significantly inhibited the growth of self-pollen tubes from 'Shatian' pummelo



Ollitrault et al. 2021

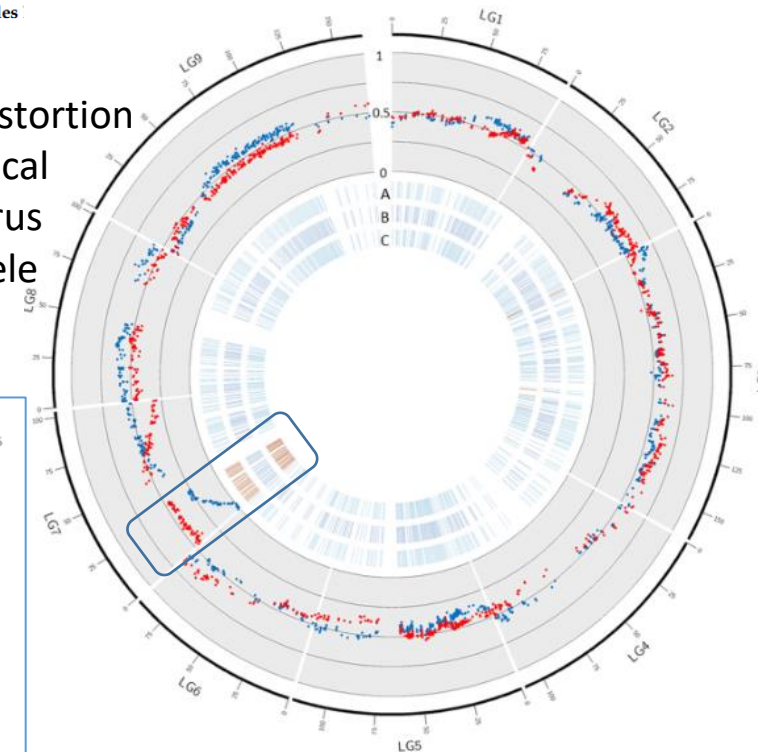
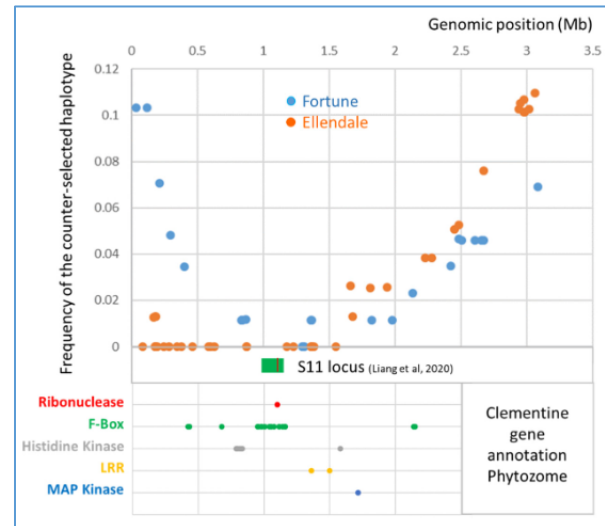


Article

Segregation Distortion for Male Parents in High Density Genetic Maps from Reciprocal Crosses between Two Self-Incompatible Cultivars Confirms a Gametophytic System for Self-Incompatibility in Citrus

Patrick Ollitrault^{1,2,*}, Dalel Ahmed³, Gilles Costantino³, Jean-Charles Pierre Mournet^{1,2}, Aude Perdereau⁴ and Yann Froelicher^{1,2}

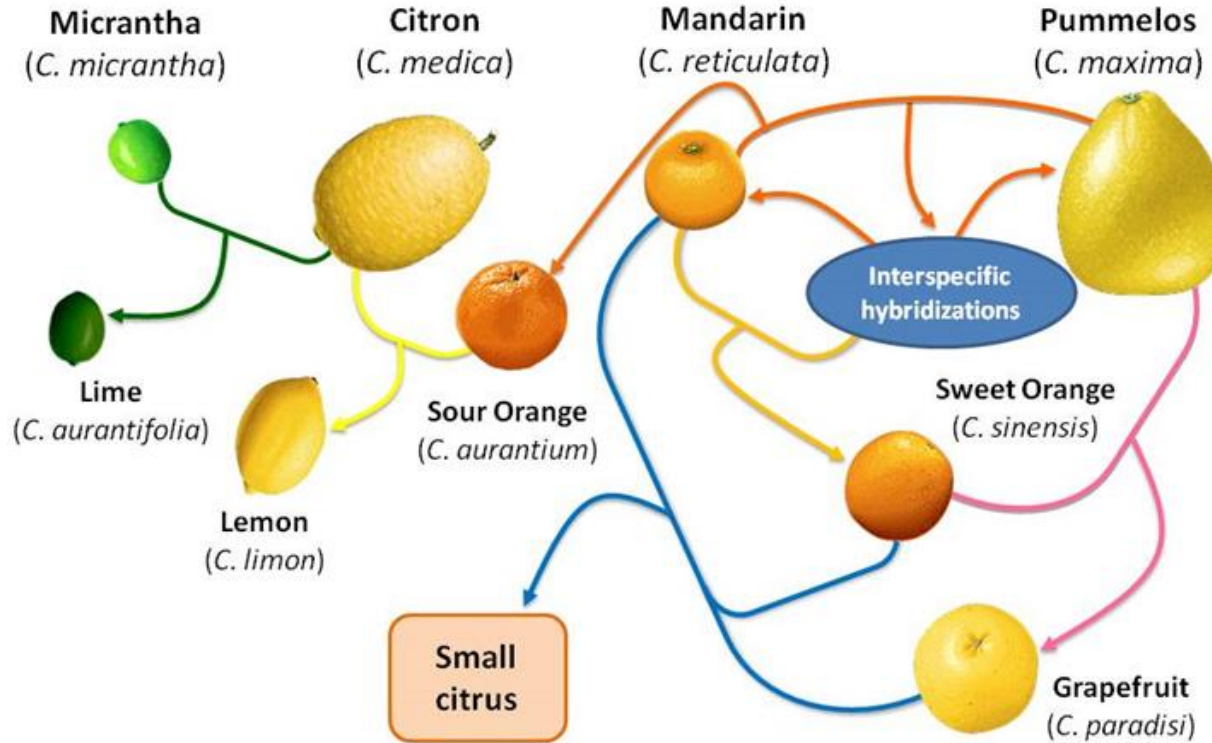
S locus location by segregation distortion (rejection of one allele) in reciprocal crosses of 2 self-incompatible citrus sharing one self-incompatible allele



gametophytic S-RNase system with the S locus located at the beginning of chromosome 7

Constraints and advantages of conventional breeding

Phylogeny of *Citrus* sp.

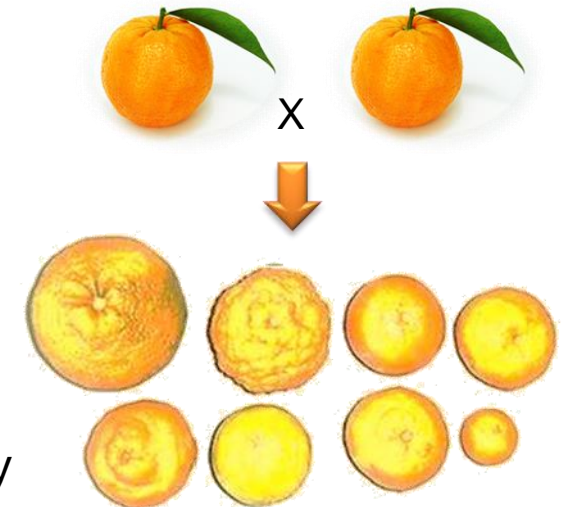


Complex genetic structures

Citrus crops are interspecific hybrids fixed by vegetative propagation



High Heterozygosity



Conventional breeding of the citrus crops is hampered by the complex genetic structures that determine their specific phenotypes.

The **highly heterozygous** interspecific mosaic structure of their genome is broken by sexual recombination => high heterogeneous phenotype with expression of deleterious alleles

Solutions for secondary species breeding



The majority of citrus cultivars are spontaneous mutants

Mutagenesis

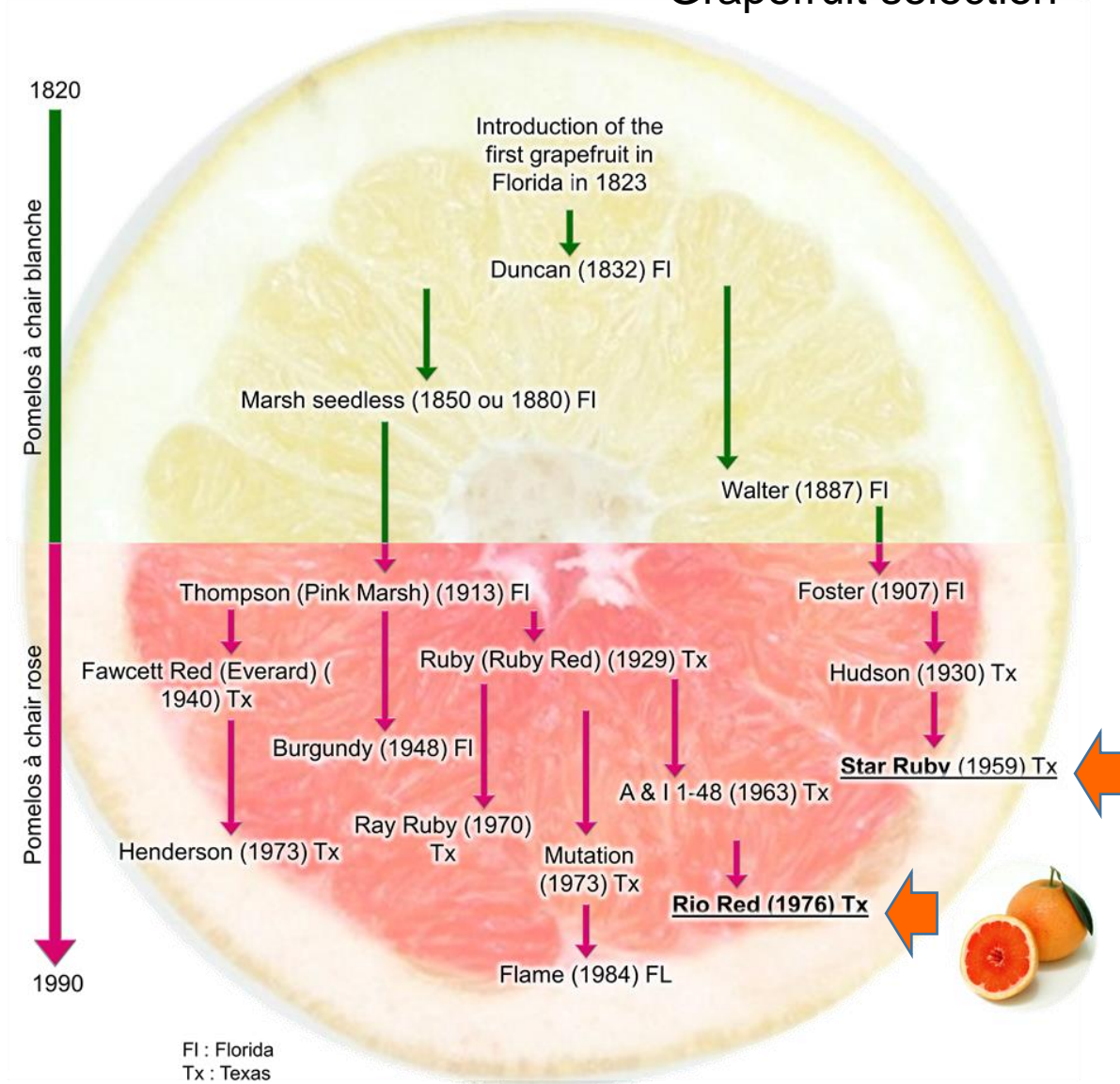
Mainly by gamma irradiation, very efficient for **sterility** / without modification on other traits

Grapefruits combined spontaneous and induced mutants: **Star Ruby** (irradiated bud woods to produce sterility), **Rio Red** (irradiated seeds to intensify the red flesh colour)

Used as the final step of selection process to produce sterility of improved hybrids

Mandarins: Afourer => **Tangold** , Murcott => **Mor**

Grapefruit selection



Citrus breeding: seedlessness



Seedless = major trait for citrus fresh fruit market

Breeding associates **parthenocarpy** and mechanisms that prevent ovule fertilization

Self-incompatibility was a very efficient way to select seedless cultivars :

Clementines and mandarins as 'Fortune', 'Ellendale', 'Nadorcott', 'Nova'

Present in pummelos (*C. maxima*) : allogamic reproduction

Seedless fruit production is obtained if the trees are grown in monovarietal orchards to avoid cross fertilization

The possibilities of using hybridization for citrus improvement



Available for **creation of new varieties** in mandarins

The breeding strategies are generally based in one reproductive cycle

Sterility associated with **parthenocarpy** is the best way to obtain seedless fruits and to avoid cross-pollination in self-incompatible varieties such as clementine



The **selection of triploid hybrids** is the most widespread used strategy in the world to obtain new sterile seedless varieties (dysfunction of meiosis => abnormal gametes)

Natural citrus polyploids

- ✓ 3x hybrids = fertilization between unreduced and normal haploid gametes
- ✓ The frequency of 3x hybrids in 2x X 2x progenies varies to 0 at 25%
- ✓ 4x = doubled diploid from chromosome doubling in somatic embryos available for 3x production by 2x X 4x fertilization



Tahiti (Key) lime
C. latifolia



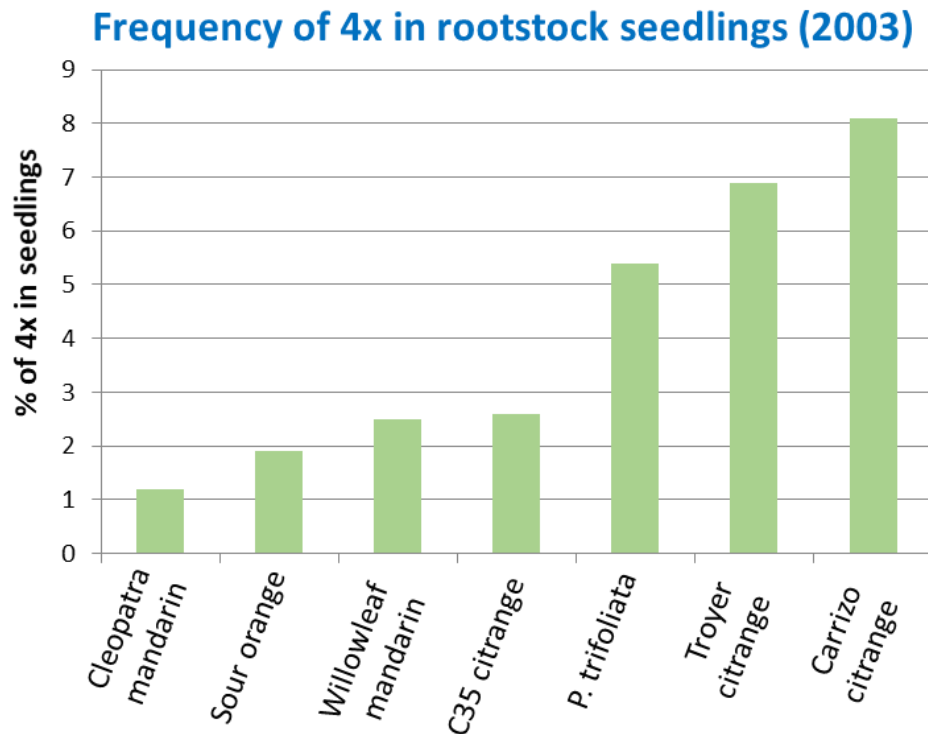
The frequency of polyploids in citrus fruits

Triploid hybrid frequency depends on variety and environment

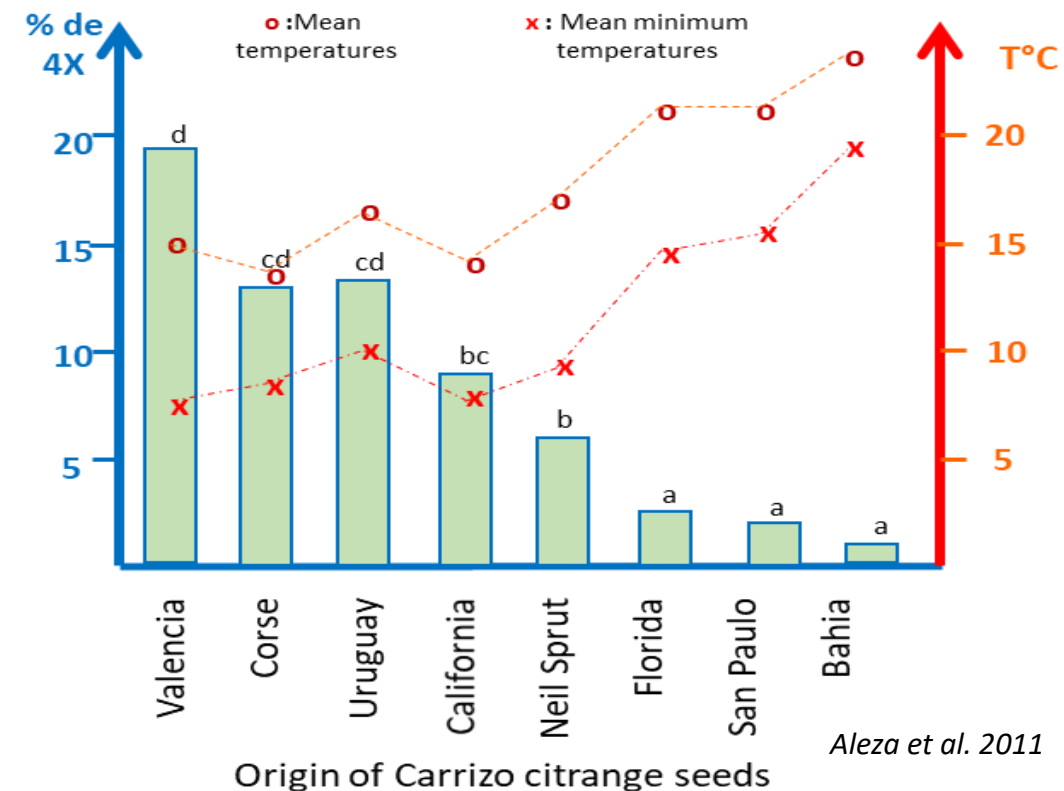
Clementine	1%	Wilking mandarin	14,6%
Citron	4%	Sweet Orange	8 to 33%
King mandarin	7%	Ortanique tangor	25%

Cold temperatures affect the meiosis process

The frequency of **doubled diploids** in seedlings depends on the variety... and the environment



Relation between temperature during blossom and frequency of 4X



Aleza et al. 2011

The INRAE-Cirad breeding program: exploitation of polyploidy

Mandarins : new cultivars

Lime (*C. latifolia*): reproduction of an elite genotype

Rootstocks : adaptation to environment constraints



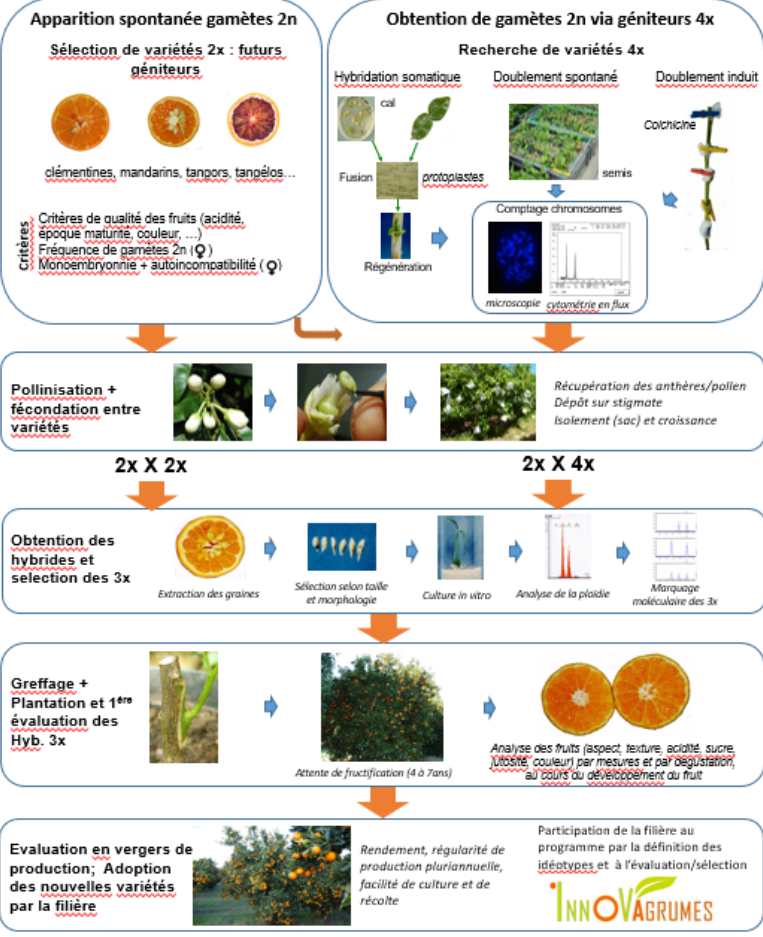
The INRAE-Cirad mandarin breeding program

Le programme Inra-Cirad de création variétale mandarine : exploitation de la polypléidie par croisements sexuels

Objectifs :

- Etendre la période de maturité au-delà de celle de la clémentine (Nov-Dec)
- Type (forme et qualité) proche de la clémentine ou alors très différent (arôme mandarine; couleur sanguine...)
- Variétés stériles (double) pour assurer une production de fruits sans pépin (pas de pollinisation des clémentiniers)

Stratégie: Créer des variétés hybrides, triploïdes (3 stocks de chromosomes => stérilité)



Objectives

- Extend the ripening period beyond that of the clementine (after December)
- PhenoType (shape and quality) close to the clementine or very different (mandarin aroma; blood color...)
- Sterility of cultivars (double M&F) to ensure seedless fruit production (no pollination of clementine trees)

Strategy

Create 3x hybrid varieties (3 stocks of chromosomes => sterility)



Selection of triploid hybrids in 2x X 2x and 2x X 4x

Utilization and mobilization of the mandarin biodiversity

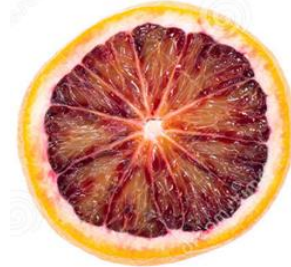
Selection of 2x genitors



Clementine



Mandarins



Tangors, tangelos

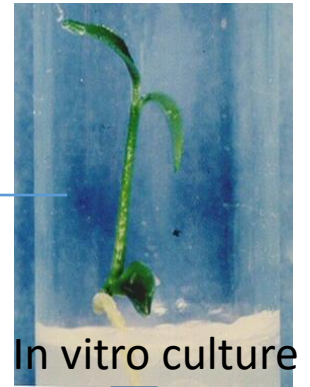
Selected Criteria

Fertilized parent :

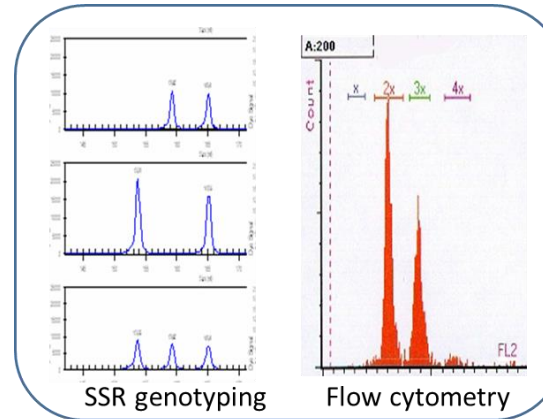
- ✓ Parthenocarpic + self-incompatible + monoembryonic
- ✓ 2n gamete frequency

Pollinators:

Quality traits (acidity, color, aroma, maturity period...)



In vitro culture



SSR genotyping

Flow cytometry

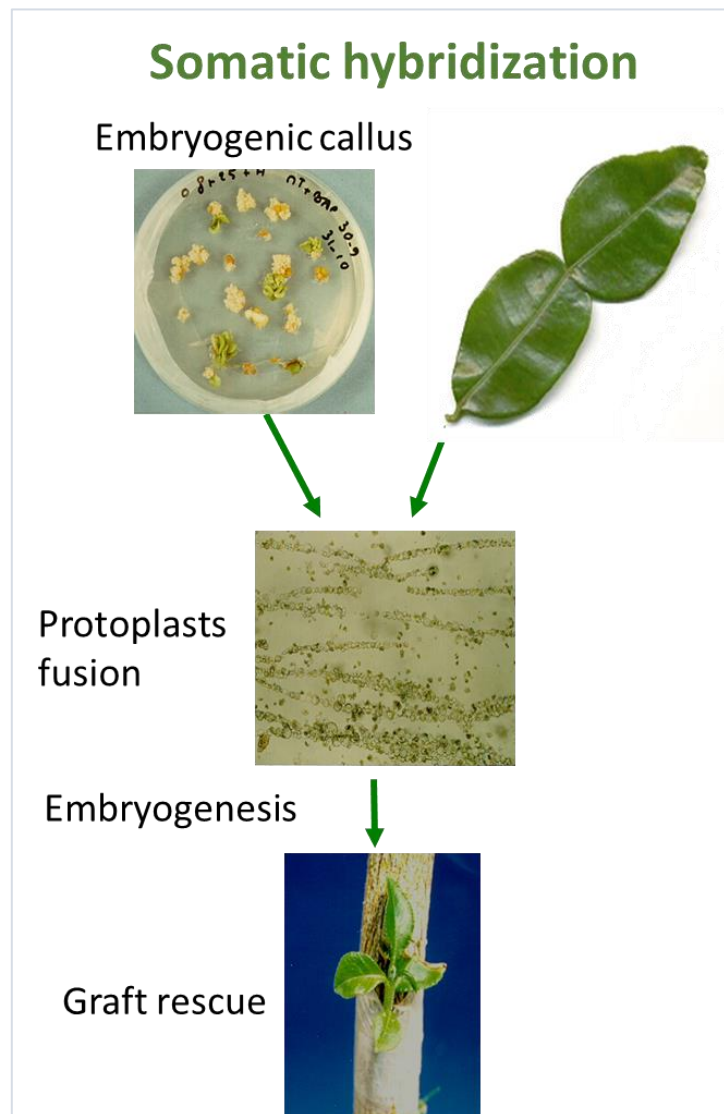
breeding program =
15 – 20 years



Safor : Spanish triploid mandarin is exploited commercially (*Aleza et al., 2010*)

Selection of triploid hybrids in 2x X 2x and 2x X 4x

How to select 4x varieties?



Spontaneous 4x (doubled diploids)

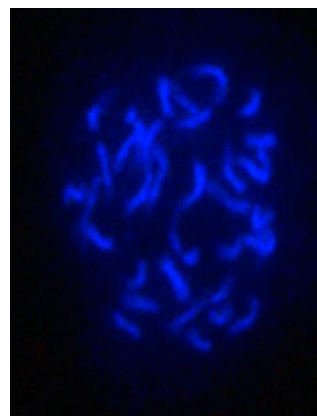


Seedlings of polyembryonic varieties

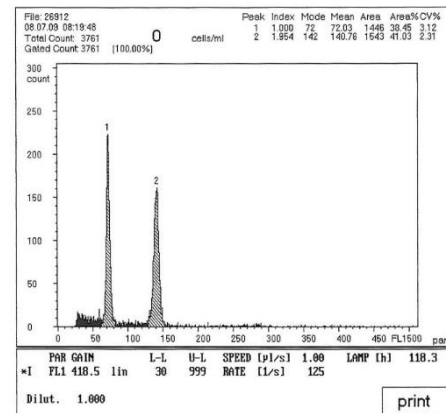
Chromosome doubling induction



Ploidy evaluation



chromosome counting



Flow cytometric evaluation

The INRAE-Cirad lime breeding program



How phylogenetic origin and genomic structures of modern cultivars can help the reconstruction of the complex genetic structure ?



Tahiti lime
C. latifolia

The case of Tahiti lime (*C. latifolia*)



Context:

- ✓ Limes (*C. latifolia*) not well adapted to the Mediterranean climate (early de-greening),
- ✓ Fruit skin contains toxic phenolic compounds (furocoumarins),
- ✓ Sensitive to Huanglongbing (very severe bacterial disease),
- ✓ Flowering is mostly grouped on one period while the needs are all of the year.

Objectives :

- Discover the genetic origin (parents & meiosis mechanisms),
- Reproduce the original parental cross,
- Select by DSNP the hybrid genotypes close to the Tahitian lime,
- Evaluate the phenotype

How genomic investigations help the reconstruction of a complex genotype

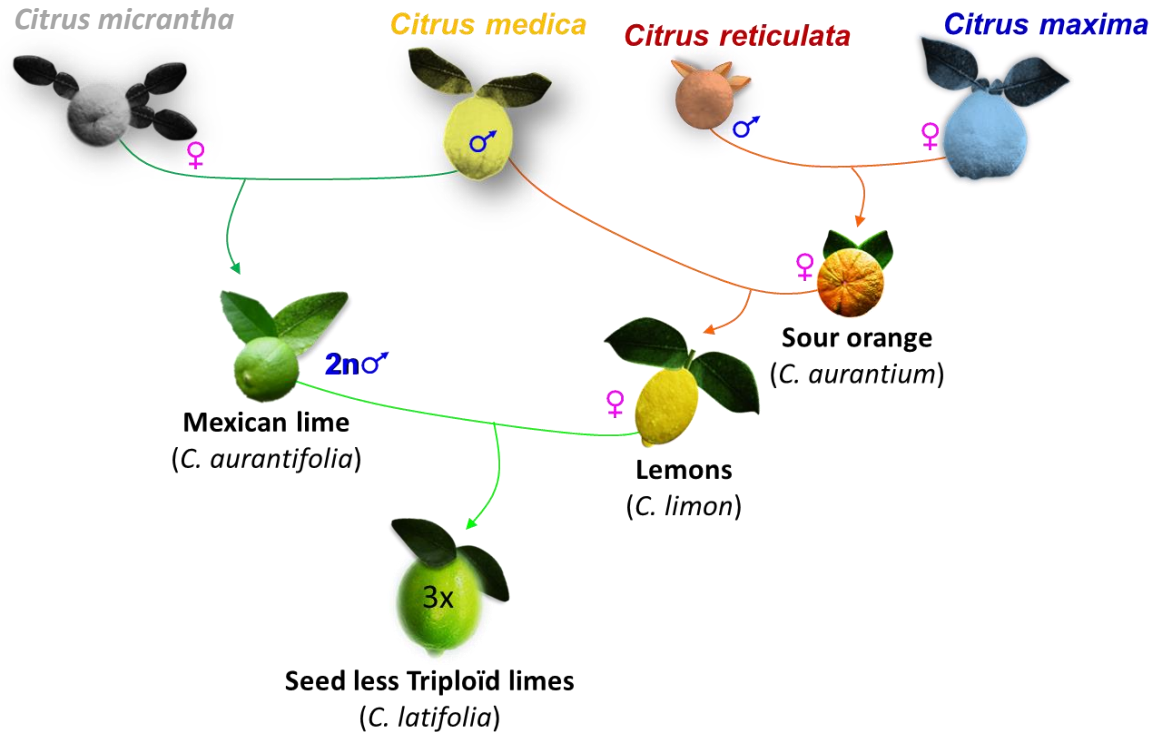


Tahiti lime
C. latifolia

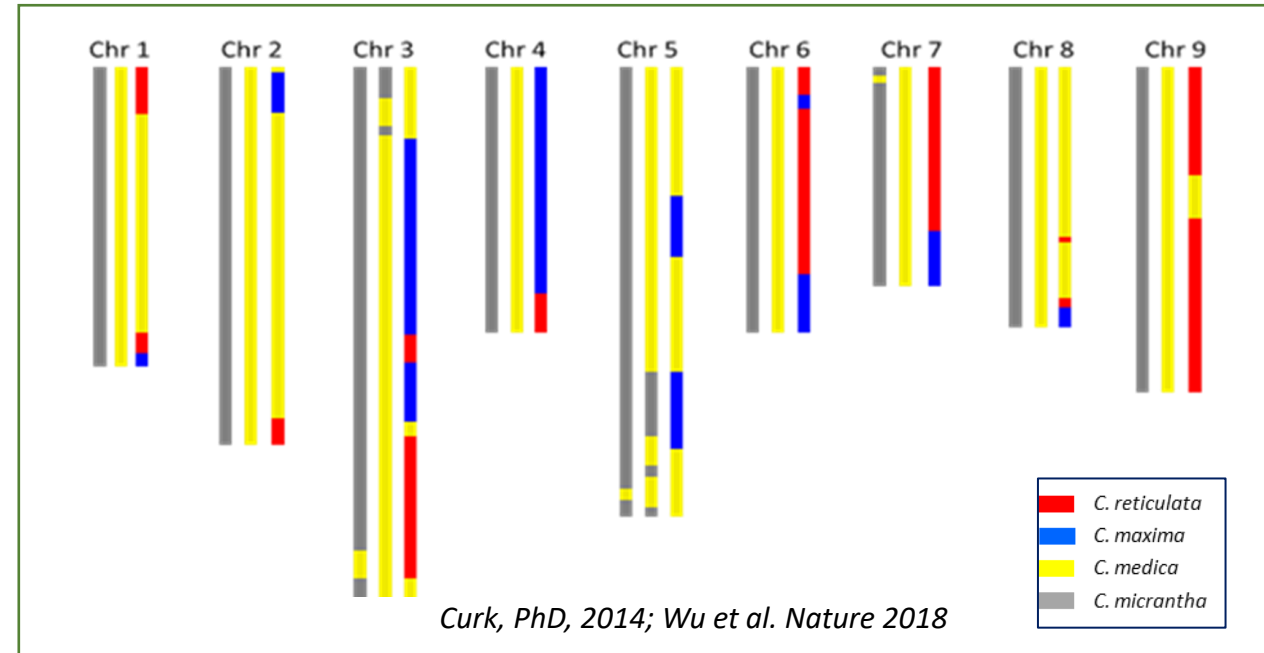


C. latifolia phylogeny deciphered by diagnostic SNPs

Deciphering interspecific mosaic structures from WGS



Structure of *C. latifolia* (3x) genome



C. latifolia (Tahiti lime) genome is an admixture of 4 ancestral species and is derived from fertilization of a lemon ovule (1n) by unreduced pollen (2n) of *C. aurantifolia* (Key lime)

Giant Key lime (*C. aurantifolia*, 4X) meiosis is predominantly disomic; this inheritance is consistent with the *C. limon* (2x) X *C. aurantifolia* (4x) cross that produced *C. latifolia* (3x)

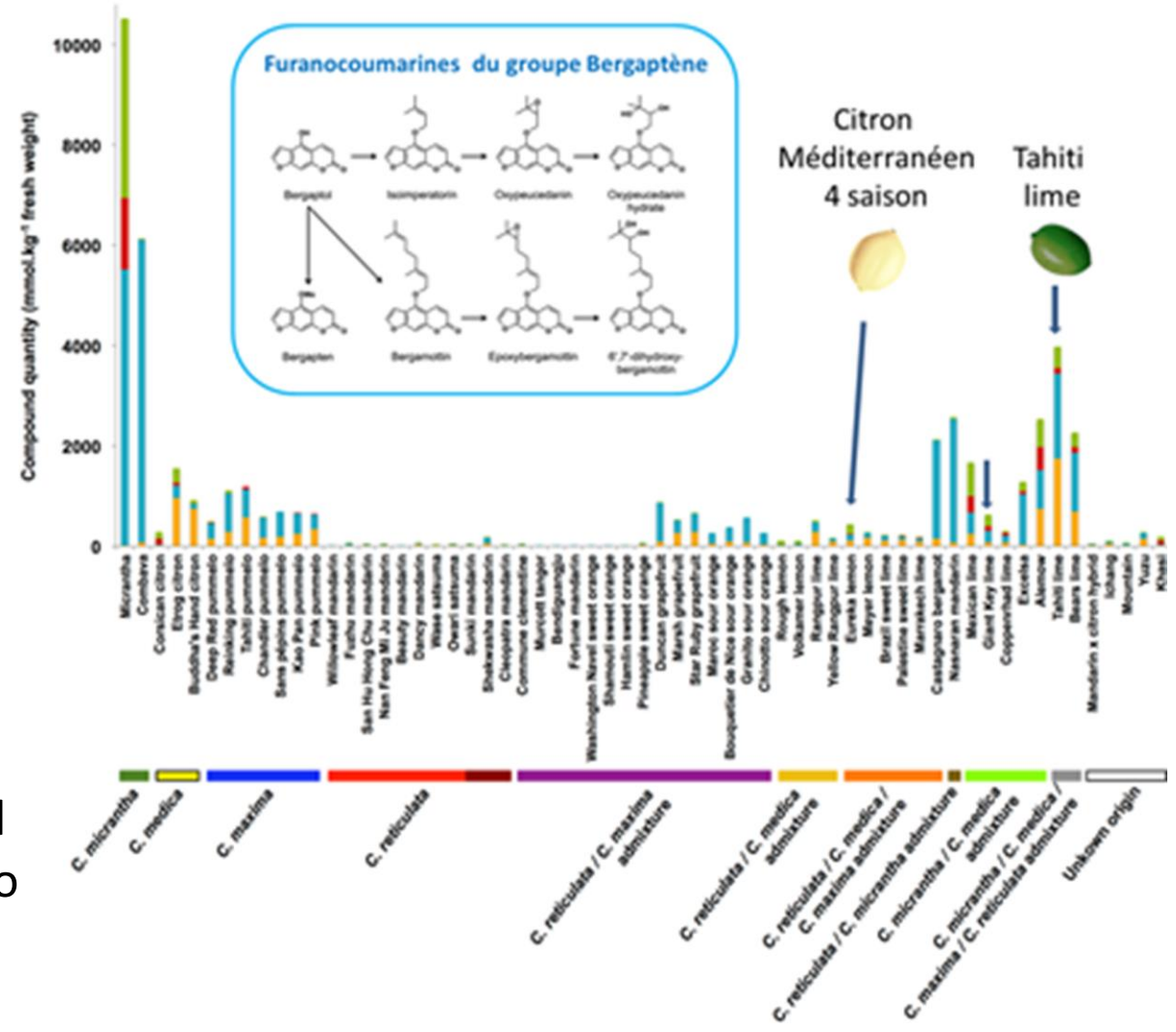
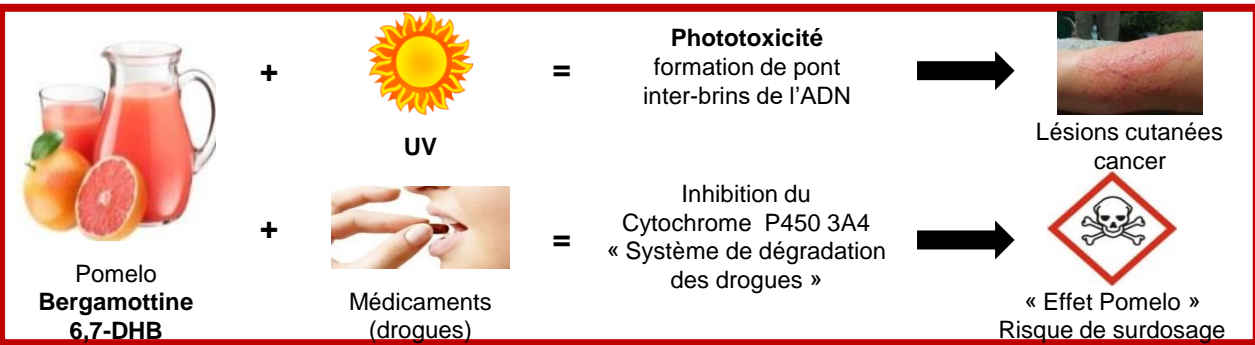
Curk et al., PlosOne, 2015; Annals of Botany 2016

Riouss et al., Annals Bot. 2018

Some inappropriate traits of lime *C. latifolia*



Furanocoumarins (Flavanoids) are toxic



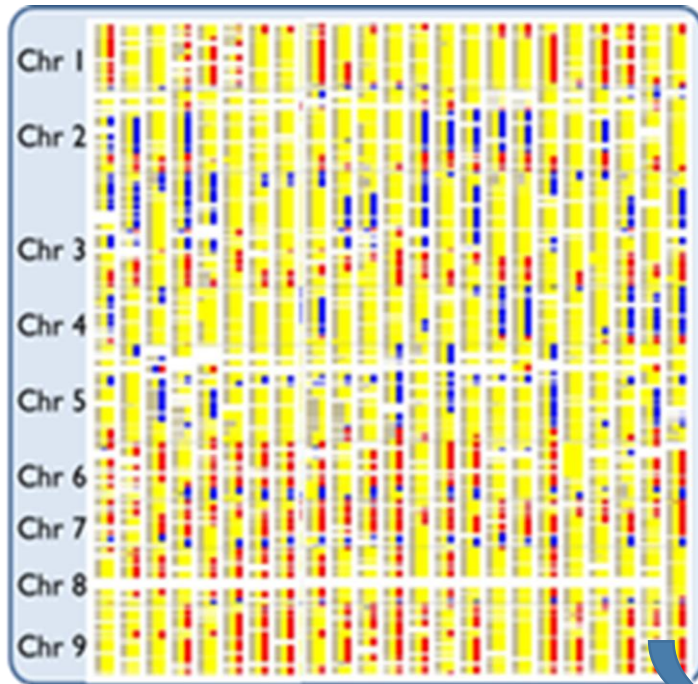
C. latifolia, like its ancestor *C. micrantha*, has very high levels of furanocoumarins; the lemon tree, its maternal parent, is almost devoid of them and Giant Key (4x) also

Dugrand-Judek et al. PlosOne 2015

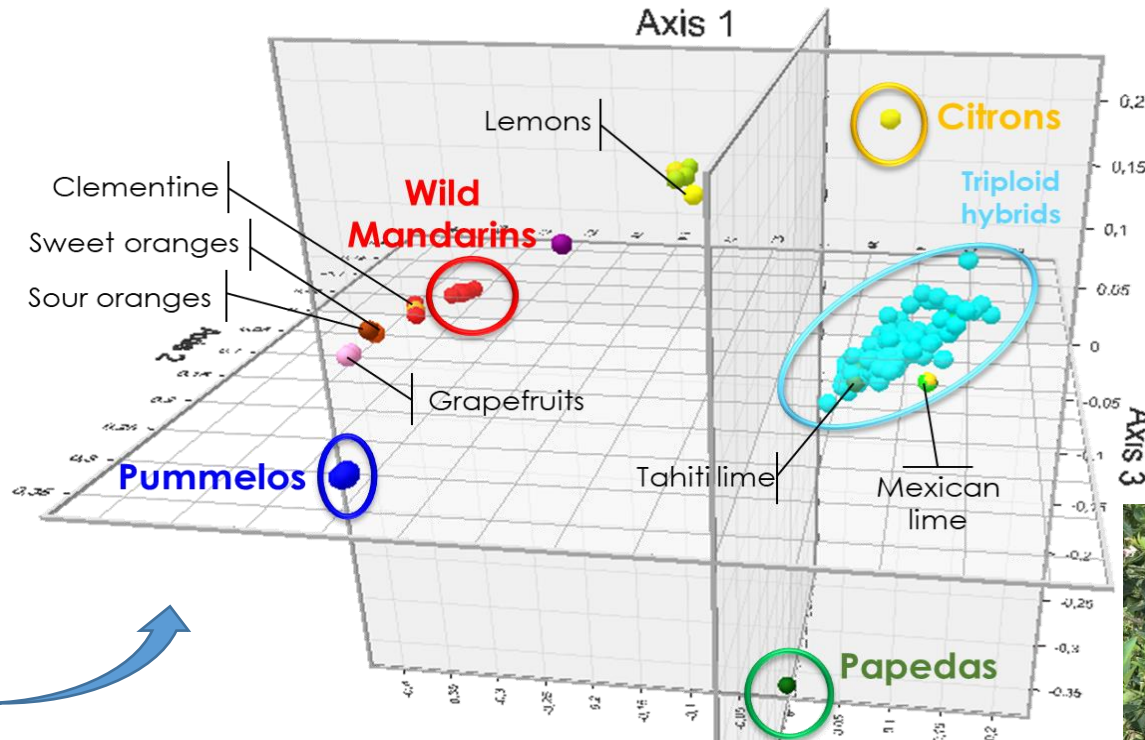
Reproduction of the cross at the origin of the Tahiti lime

C. limon (2x) x *C. aurantifolia* 4X (Giant Key) => 500 hybrids (3x)

Analyze of hybrid genomes by DSNP (GBS)



Diversity of hybrid population in relation with parental genotypes and ancestral species (DSNP)



Triploid hybrids are close to *C. latifolia*

Ahmed et al. Front. Plant Sci. 2020



Evaluation of hybrids in progress in Guadeloupe and in Corsica (HLB resistance, aroma...)



A promising 3x hybrid with permanent blooming

**Rootstock ploidy manipulation for a better
adaptation; new breeding strategies**

Adapting to climate change through rootstock selection

Constraints related to climate change

Abiotic stress:

- ★ Water deficit / drought
- ★ Salinity
- ★ Temperature (low)
- ★ Ferric chlorosis (Alkalinity)

- ★ Solution par sélection P-G
- C Non présent en Méditerr.
- C Non présent en Corse

Biotic stress:

- ★ Root rot Gommosis (*Phytophthora*)
- Brown spot (*Alternaria alternata*)
- ★ Mal secco (*Deuterophoma tracheiphila*)
- ★ Tristeza (CTV)
- Variegated Chlorosis (*Xylella fastidiosa*)
- Citrus canker (*Xanthomonas campestris*)
- ★ HLB (*Candidatus liberobacter*)

Fruit quality traits:

- ★ Acidity / early collapse
- ★ Peel coloration
- ★ Fruit size
- ★ Juiciness



Most constraints can be managed or modulated via the choice of rootstock

Adapting to climate change through rootstock selection

There are sources of tolerance to the various stresses in the different gene pools, which are useful for selecting tolerant rootstocks.



	<i>Alk.</i>	<i>Chl.</i>	<i>Bore</i>	<i>Dr.</i>	<i>Trist.</i>	<i>Nem.</i>	<i>Phyto</i>
<i>Citrus relatives</i>							
<i>Eremocitrus glauca</i>	T	TT	TT	TT	T	S	T
<i>Severinia buxifolia</i>	pT	TT	TT	pT	T	T	T
<i>Cleopatra mandarin</i>	T	T	pT	pT	T	S	S
<i>Sour orange</i>	T	T	pT	T	S	S	T
<i>Sweet Orange</i>	pT	S	pT	T	T	S	S
<i>C. junos</i>	S	pT	?	T	T	S	T
<i>C. taiwanica</i>	T	T	pT	pT	pT	S	T
<i>Lemon group</i>							
<i>Rangpur lime</i>	T	TT	T	TT	T	S	S
<i>Rough lemon</i>	pT	S	pT	T	T	S	pT
<i>C. macrophylla</i>	T	pT	pT	T	S	S	T
<i>C. volkameriana</i>	T	pT	pT	T	T	S	T
<i>P. trifoliata</i>	S	S	S	pT	TT	TT	TT

Rootstock ploidy manipulation for a better adaptation; physiological insights

Doubled diploid AB → AABB

Obtention: spontaneous chromosome stock doubling in nucellar cells (Aleza et al., 2011)
Most traditional citrus rootstock are actually available as doubled diploid

Advantages:

- global conservation of tolerance to biotic stress,
- enhancement of tolerance to some abiotic stresses,
- frequent vigor reduction in doubled diploid

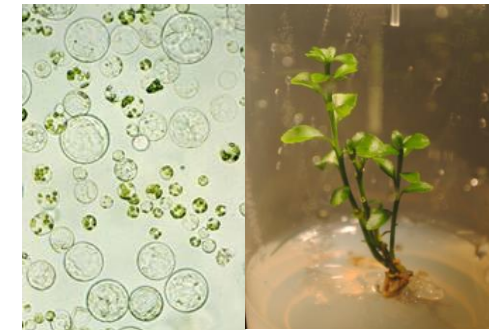


Somatic hybrids AB + CD → ABCD

Obtention: protoplast fusion (Grosser et al., 2010; Dambier et al., 2011)

Advantages:

- addition of dominant traits of complementary parents (Poncirus/Citrus)
- frequent vigor reduction



Tetrazyg AABB x CCDD; ABCD x EFGH

Obtention: sexual hybridization at tetraploid level (Grosser et al. 2010)

Advantages:

- very large possibility of allelic combinations
- tetraploid behavior facing abiotic stresses

Limits: totally new genotypes and numerous selection objectives

Needs: efficient early selection ; marker trait association



Sweet orange grafted on FLHORAG1 somatic hybrid (CIRAD)
Calcareous soils Morocco

Valencia delta orange grafted on tetraploid (DD) Rangpur lime

11 days without watering

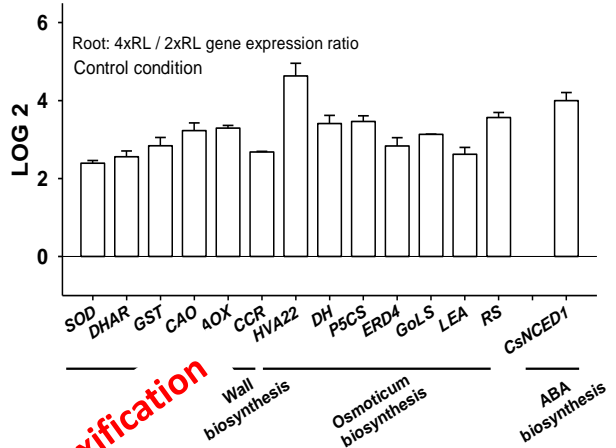


Rangpur L. 2x/V

Rangpur L. 4x/V

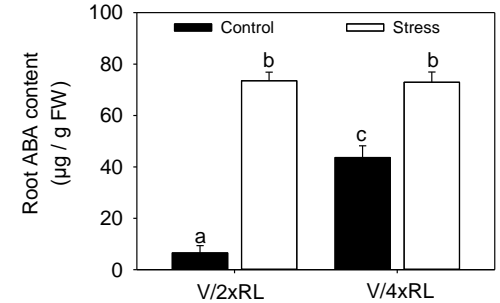
Rootstock ploidy manipulation for a better adaptation; physiological insights

Candidate gene for adaptation expression at the root level:
4x > 2x



Detoxification

[ABA] in root:
2x < 4x

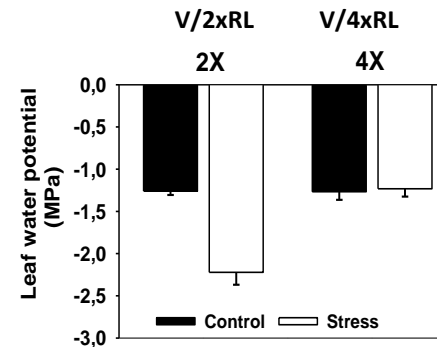
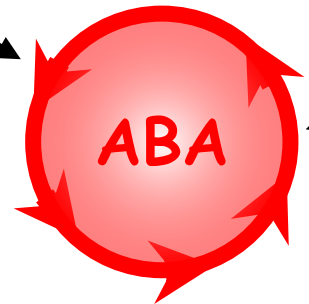


Leaf transpiration in control condition: V/2xRL > V/4xRL

Plant growth:
2x > 4x

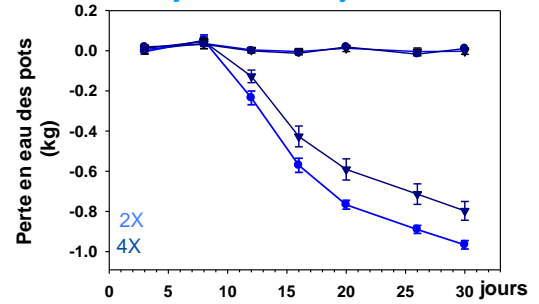
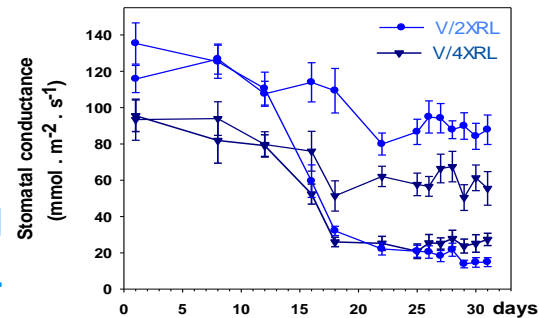


Tree water consumption:
V / 2xRL > V / 4xRL



Leaf water potential in stress:
V/2xRL < V/4xRL

Pot water loss along water deficit:
V/2xRL > V/4xRL



Huanglongbing: does exist varietal solution for HLB ?

- No strict resistance to *Liberobacter asiaticus* in the genetic resources
- However, the impact is different depending of the variety



- Comportment that is different in infected orchards: grapefruit (Jackson et Triumph)
- Minor sensitivity in some small citrus Temple, KanSan mandarins)
- Better comportment of 3x (Tahiti) compared to 2x (Mexican lime, etc)
- Appetence of psyllids for the variety is importance regarding the sensibility



Orange trees infected using sensitive and « tolerant » rootstocks

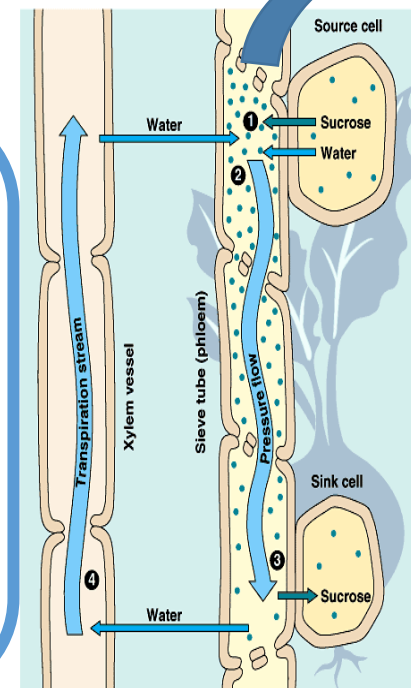
- Some 4x rootstocks present a reduced sensitivity (or a partial tolerance?)

Huanglongbing: what do we know ?

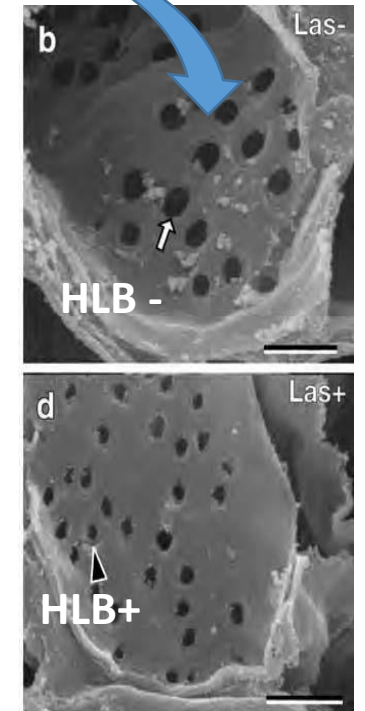
- High concentration of starch and sugars in leaves
- Callosis synthesis in the cells of phloem
- Low presence of assimilates at root level
- Deficiency in leaf lead to accumulation of starch and sugars

Symptoms seem to be associated to physiological and metabolomic perturbations:

- Leaf symptoms are similar to leaf removing => limitation of the transport of phloem
- Limitation of the conversion of starch to sugars
=> perturbation of the photosynthesis
- Callosis in the cells of the phloem: obstruction of connections between cells
=> Limitation of the symplast transport



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Xylem Phloem



Conclusion

Citrus diversity is the result of a speciation phase during the allopatric evolution of ancestral populations and subsequent interspecific crosses.

Citrus reproduction is complex and diverse depending on the species, and apomixis is responsible for the clonal multiplication of interspecific hybrids elevated to the rank of secondary species.

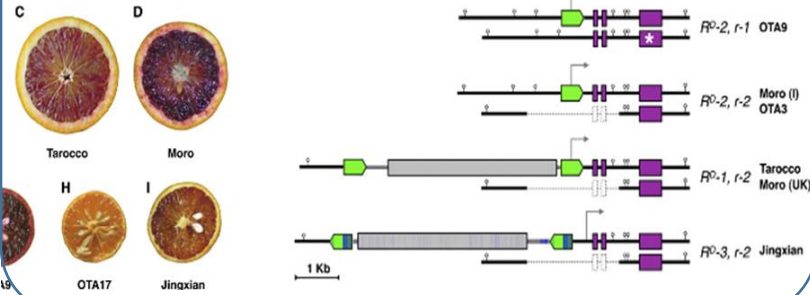
Breeding strategies are adapted to biological constraints: mutagenesis for secondary species, hybridization for primary species mixing triploidy (sterility) and tetraploidy (increased tolerance to biotic and abiotic stress)...

Genome investigations (DSNP, GBS and NGS) are useful for reproduction of a complex interspecific hybrids and for MAS



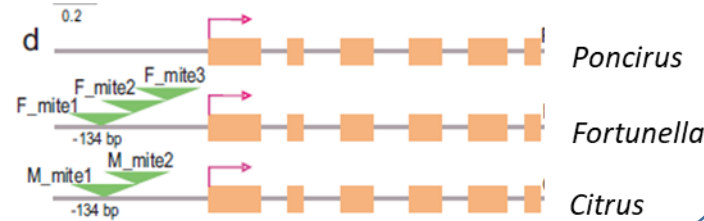
Toward marker assisted selection (MAS) or genome editing

Ruby gene



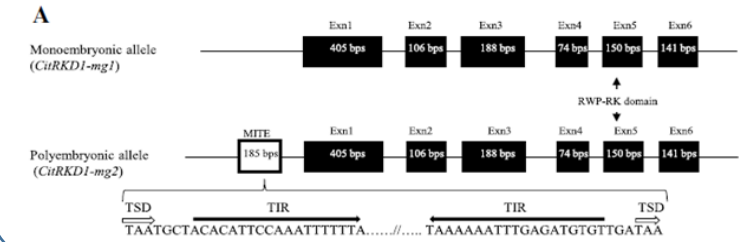
pulp acidity

(Wang et al. 2022)



Apomixis (Polyembryony)

(Shimada et al. 2018)

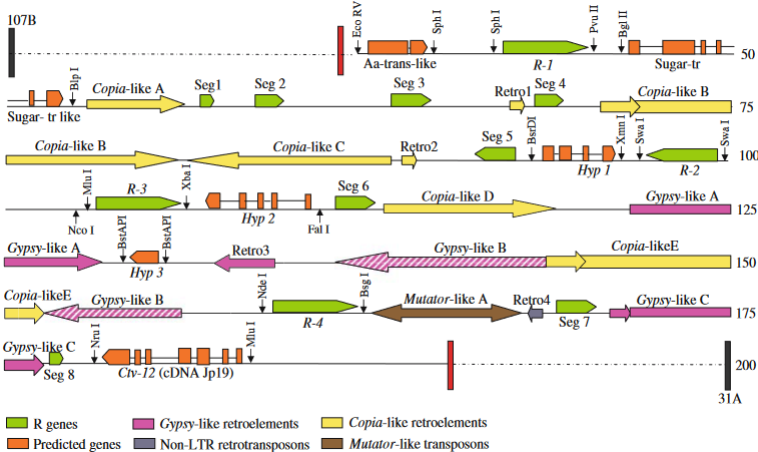


Plant Molecular Biology (2006) 61:399-414
DOI 10.1007/s1103-006-0018-7

CTV resistance gene

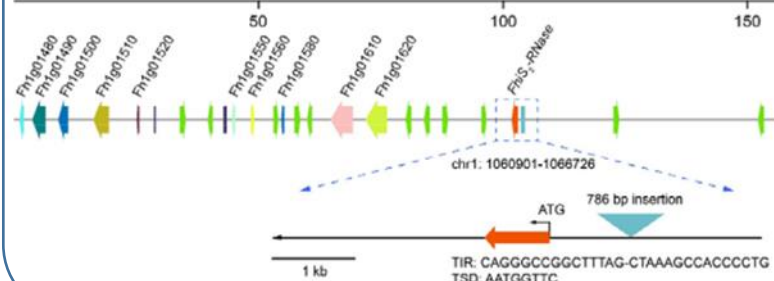
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Refinement of the Citrus tristeza virus resistance gene (*Ctv*) positional map in *Poncirus trifoliata* and generation of transgenic grapefruit (*Citrus paradisi*) plant lines with candidate resistance genes in this region



self-compatibility

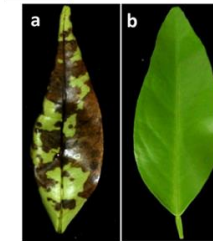
(Hu et al. 2023)



SNP and InDel markers can be used for selection of different traits in breeding programs

frontiers
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Fine Mapping for Identification of Citrus Alternaria Brown Spot Candidate Resistance Genes and Development of New SNP Markers for Marker-Assisted Selection

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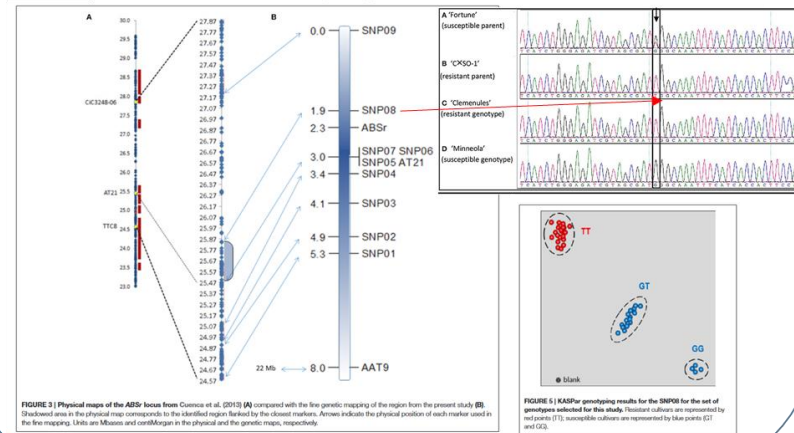


FIGURE 3 | Physical maps of the *ABSr* locus from Cuenca et al. (2013) (A) compared with the fine genetic mapping of the region from the present study (B). Shaded areas in the physical map correspond to the identified region flanked by the closest markers. Arrows indicate the physical position of each marker used in the fine mapping. Units are Mbases and centiMorgan in the physical and the genetic maps, respectively.

FIGURE 5 | MAS-Par genotyping results for the SNPs for the set of genotypes selected for this study. Resistant cultivars are represented by red points (TT), susceptible cultivars are represented by blue points (GG) and GG.



Thank
You...