

# Origen, consecuencias y estudio de la poliploidía en plantas

Genética, Genómica y Mejora Vegetal

Febrero de 2023

**Máster en Genética y Evolución**  
(Especialidad Agroalimentaria)



**UNIVERSIDAD  
DE GRANADA**

**Rafael Navajas-Pérez**

# Una visión naif de la Genética Vegetal

## Un gen, una función...

### Mendel's Genes: Toward a Full Molecular Characterization

James B. Reid and John J. Ross<sup>1</sup>

School of Plant Science, University of Tasmania, Hobart, Tasmania 7001, Australia

Genetics, Vol. 189, 3–10 September 2011

Table 1 Seven characters of *P. sativum* examined by Mendel and a summary of the genes, phenotypes, and presumed mutations involved

Trait	Dominant phenotype	Recessive phenotype	Symbol group	Linkage group	Cloned	Gene function	Molecular nature of mutation
Seed shape	Round	Wrinkled	<i>R</i>	V	Yes	Starch branching enzyme 1	0.8-kb insertion
Stem length	Tall	Dwarf	<i>LE</i>	III	Yes	GA 3-oxidase1	G-to-A substitution
Cotyledon color	Yellow	Green	<i>I</i>	I	Yes	Stay-green gene	6-bp insertion
Seed coat/flower color	Purple	White	<i>A</i>	II	Yes	bHLH transcription factor	G-to-A at splice site
Pod color	Green	Yellow	<i>GP</i>	V	No	Chloroplast structure in pod wall	Unknown
Pod form	Inflated	Constricted	<i>V?</i>	III	No	Sclerenchyma formation in pods	Unknown
Position of flowers	Axial	Terminal	<i>FA</i>	IV	No	Meristem function	Unknown

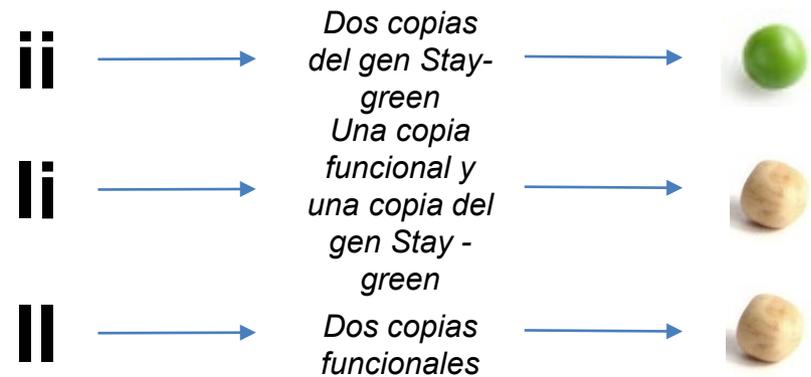
References are given in the text.



**Dominancia/Recesividad**

**I**

**i**



Fuente: Fotolia

# Una visión naif de la Genética Vegetal

## Un gen, una función... (II)

*VvmybA1*

Inserción Gret1

10422 bp

Excisión Gret1

44 bp

829 bp

Modificado de This et al., 2007 (Imágenes de unifeed.club)



Dos copias intactas del gen *VvmybA1*

$V^+V^+$

Producción normal de antocianinas



Dos copias mutadas del gen *VvmybA1*

$VV$

Sin producción de antocianinas



Una copia mutada y otra funcional del gen *VvmybA1*

$V^+V$

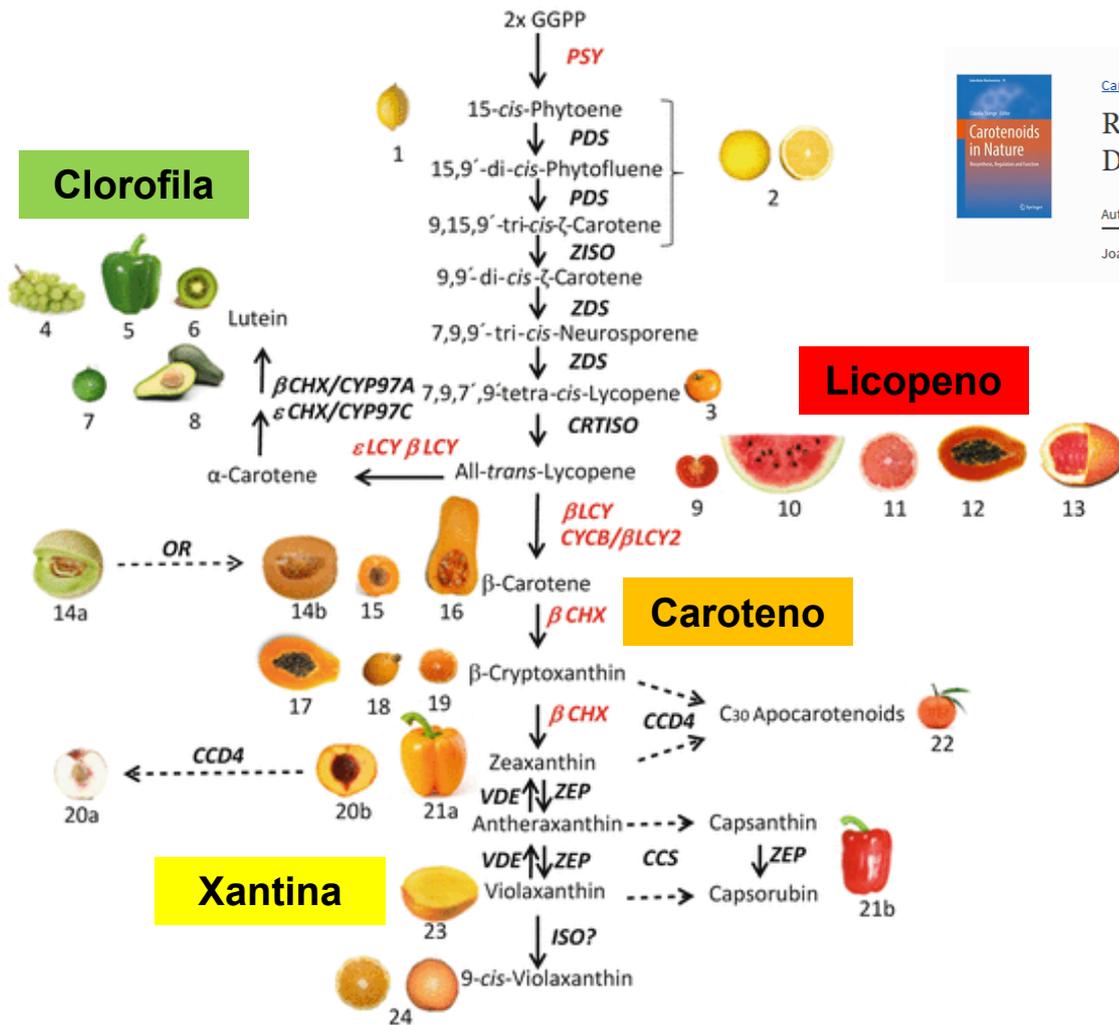
Se produce la mitad de la cantidad normal de antocianinas

### Herencia Intermedia

**Wine grape (*Vitis vinifera* L.) color associates with allelic variation in the domestication gene *VvmybA1***

# Una visión naif de la Genética Vegetal

Varios genes, una función... (Interacción génica)



Carotenoids in Nature pp 161-198 | [Cite as](#)

Regulation of Carotenoid Biosynthesis During Fruit Development

Authors [Authors and affiliations](#)

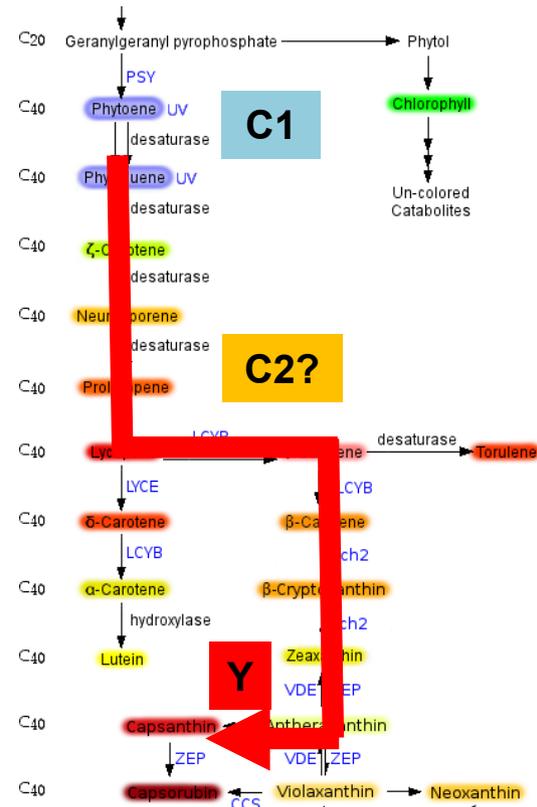
Joanna Lado , Lorenzo Zacarías, María Jesús Rodrigo



Foto: R. Navajas

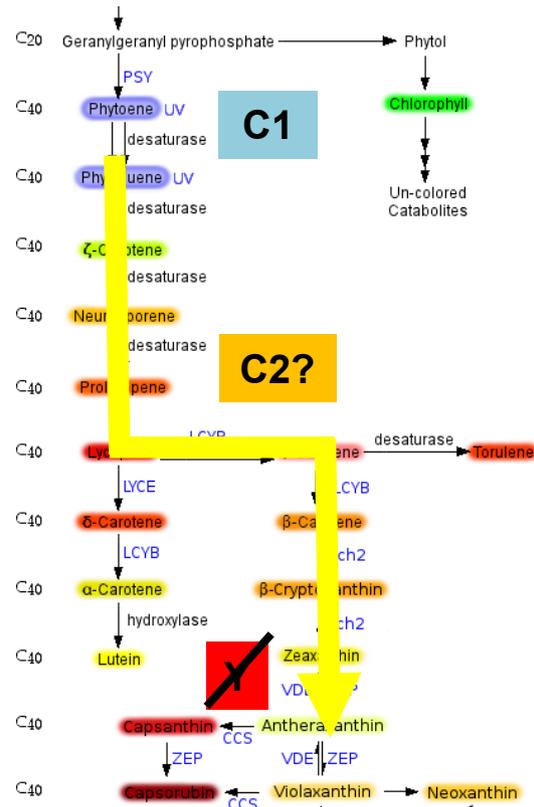
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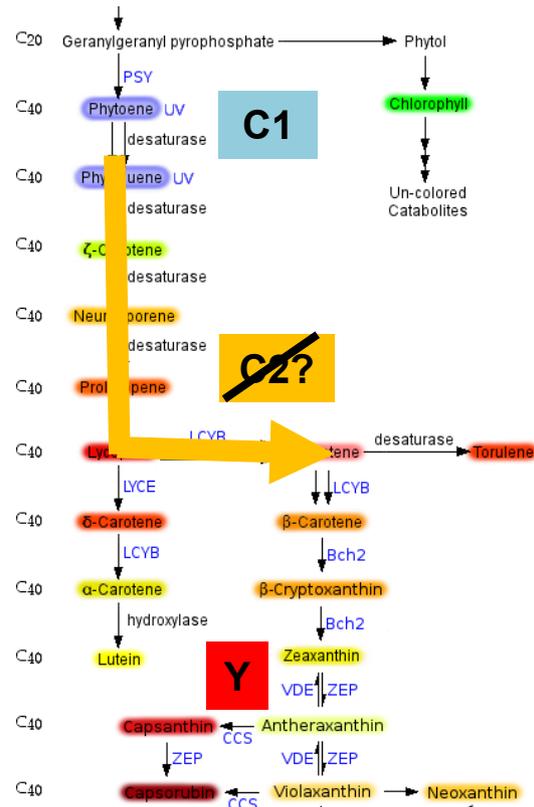
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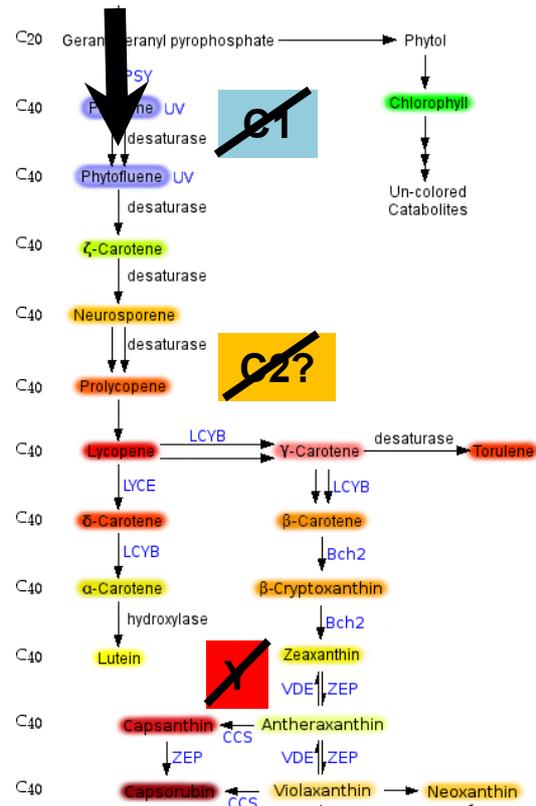
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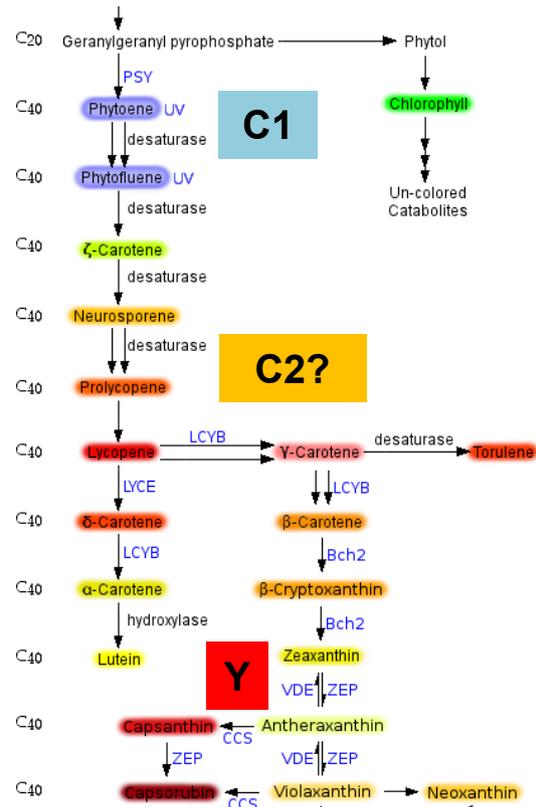
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Varios genes, una función... (Interacción génica)

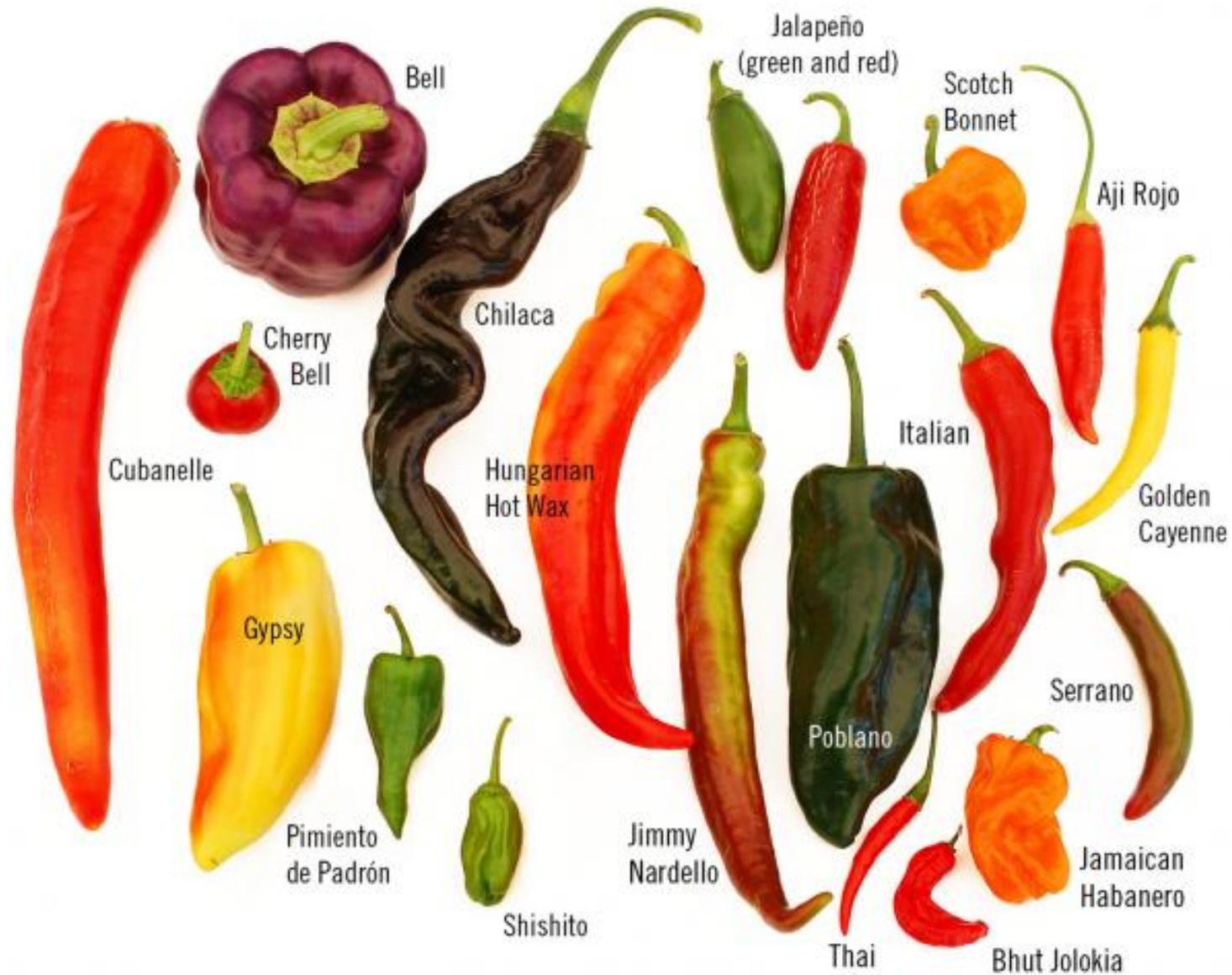


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Varios genes, una función... (Interacción génica)

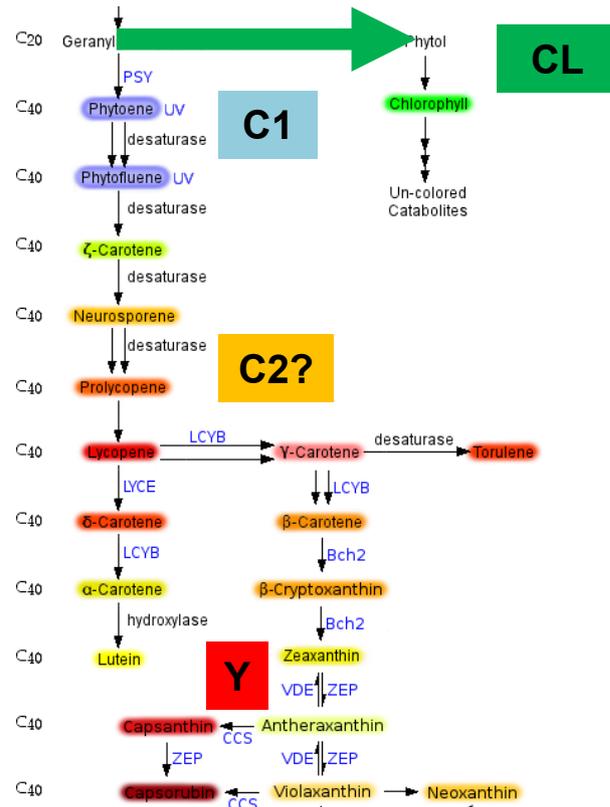


# CUESA's GUIDE to SWEET AND HOT PEPPERS

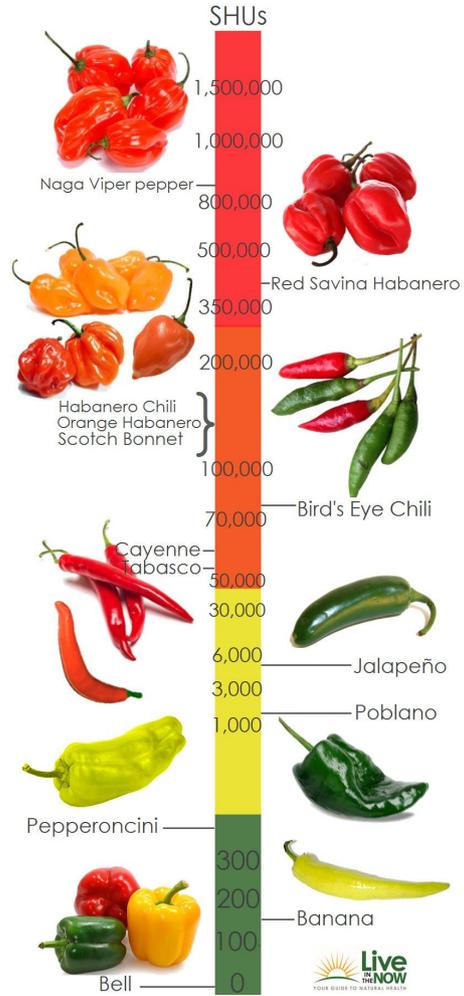


# Una visión naif de la Genética Vegetal

Varios genes, una función... (Interacción génica)

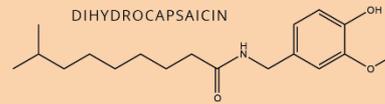
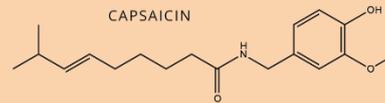


Pssst... Capsaicin benefits heart health! So,  
**HOW SPICY IS THAT PEPPER?**



# THE CHEMISTRY OF A CHILLI

## CAPSAICINOIDS

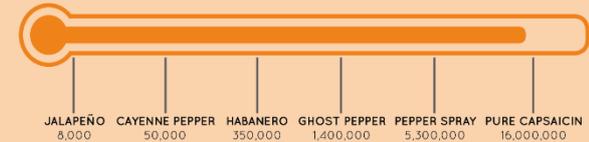


The spiciness of chillis is due to the presence of compounds called capsaicinoids. The two compounds above are the main capsaicinoids in chilli peppers. They cause a burning sensation when they come into contact with mucous membranes, due to their interaction with pain and heat sensing neurons.

Capsaicin is also used in some brands of pepper spray, and studies have shown it may be capable of killing prostate and lung cancer cells. It is toxic in large quantities.



## THE SCOVILLE HEAT INDEX



The Scoville scale is a taste detection based method for rating the heat of chilli peppers. A measured amount of pepper extract has sugar added to it incrementally until the heat is undetectable through taste. Though it is an imprecise method, it has been estimated that 1 unit corresponds to 18µM.

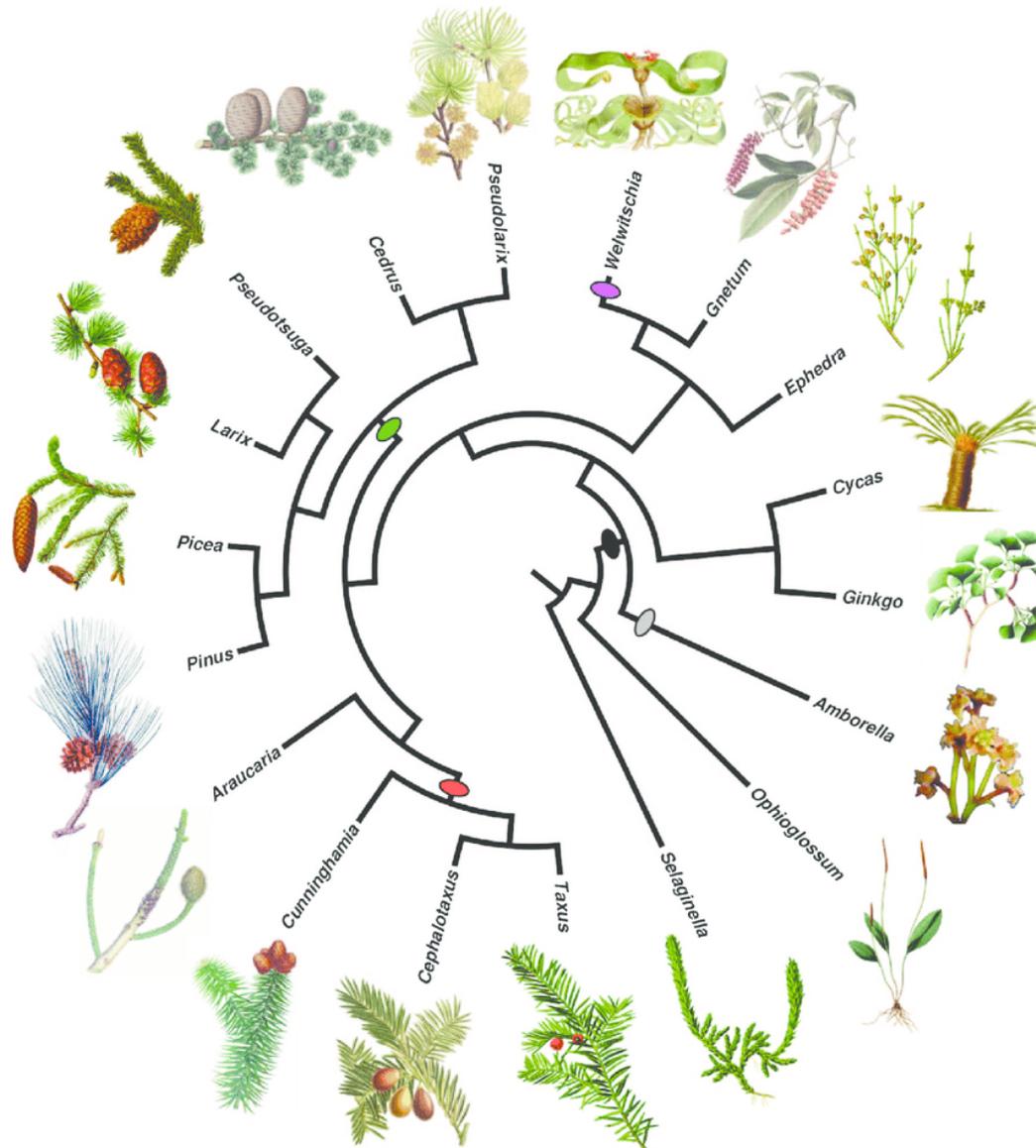
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 PHOTO: <https://www.flickr.com/photos/mainaphotonut/6978313358/>



# Early genome duplications in conifers and other seed plants

Zheng Li,<sup>1</sup> Anthony E. Baniaga,<sup>1</sup> Emily B. Sessa,<sup>2</sup> Moira Scascitelli,<sup>3</sup> Sean W. Graham,<sup>3</sup> Loren H. Rieseberg,<sup>3,4</sup> Michael S. Barker<sup>1\*</sup>

Li et al. *Sci. Adv.* 2015;1:e1501084 20 November 2015



# Large-Scale Gene Relocations following an Ancient Genome Triplication Associated with the Diversification of Core Eudicots

Citation: Wang Y, Ficklin SP, Wang X, Feltus FA, Paterson AH (2016) Large-Scale Gene Relocations following an Ancient Genome Triplication Associated with the Diversification of Core Eudicots. PLoS ONE 11(5): e0155637. doi:10.1371/journal.pone.0155637

Yupeng Wang<sup>1</sup>, Stephen P. Ficklin<sup>2</sup>, Xiyin Wang<sup>1</sup>, F. Alex Feltus<sup>3</sup>, Andrew H. Paterson<sup>1\*</sup>

- **Eudicotas basales:**
  - [Buxales](#)
  - [Proteales](#)
  - [Ranunculales](#)
  - [Trochodendrales](#)
- **Eudicotas centrales:**
  - Clados
    - [Superasterids](#)
    - [Superrosids](#)
  - Órdenes
    - [Berberidopsidales](#)
    - [Caryophyllales](#)
    - [Dilleniales](#)
    - [Gunnerales](#)
    - [Santalales](#)
    - [Saxifragales](#)



La mayoría de los paleopoliploides son consecuencia de duplicaciones, excepto algunos que conservan **triplicaciones e incluso quintuplicaciones**:

## A genome triplication associated with early diversification of the core eudicots

Yuannian Jiao<sup>1,2</sup>, Jim Leebens-Mack<sup>3</sup>, Saravanaraj Ayyampalayam<sup>3</sup>, John E Bowers<sup>3</sup>, Michael R McKain<sup>3</sup>, Joel McNeal<sup>3,4</sup>, Megan Rolf<sup>5</sup>, Daniel R Ruzicka<sup>5</sup>, Eric Wafula<sup>2</sup>, Norman J Wickett<sup>2,6</sup>, Xiaolei Wu<sup>7</sup>, Yong Zhang<sup>7</sup>, Jun Wang<sup>7,8</sup>, Yeting Zhang<sup>2,9</sup>, Eric J Carpenter<sup>10</sup>, Michael K Deyholos<sup>10</sup>, Toni M Kutchan<sup>5</sup>, Andre S Chanderbali<sup>11,12</sup>, Pamela S Soltis<sup>11</sup>, Dennis W Stevenson<sup>13</sup>, Richard McCombie<sup>14</sup>, J Chris Pires<sup>15</sup>, Gane Ka-Shu Wong<sup>7,16</sup>, Douglas E Soltis<sup>12</sup> and Claude W dePamphilis<sup>1,2\*</sup>

Jiao *et al. Genome Biology* 2012, **13**:R3  
<http://genomebiology.com/2012/13/1/R3>

Received 23 Oct 2013 | Accepted 22 Apr 2014 | Published 23 May 2014

DOI: 10.1038/ncomms4930

OPEN

## The *Brassica oleracea* genome reveals the asymmetrical evolution of polyploid genomes



doi:10.1038/nature11119

## The tomato genome sequence provides insights into fleshy fruit evolution

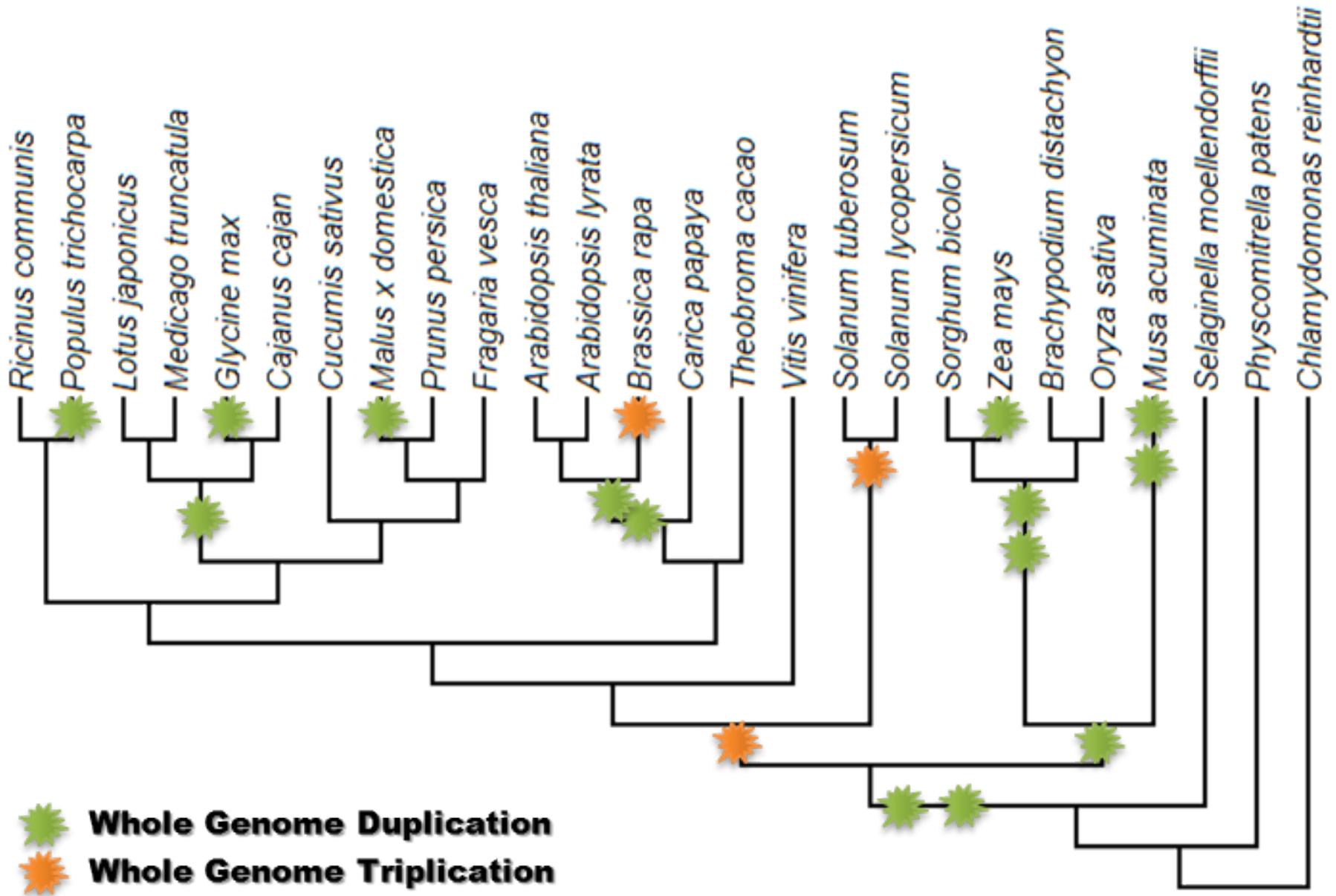
31 MAY 2012 | VOL 485 | NATURE | 635

The Tomato Genome Consortium\*

## Comparative genomic de-convolution of the cotton genome revealed a decaploid ancestor and widespread chromosomal fractionation

*New Phytologist* (2016) **209**: 1252–1263  
doi: 10.1111/nph.13689

Xiyin Wang<sup>1,2,3</sup>, Hui Guo<sup>1,4</sup>, Jinpeng Wang<sup>2,3</sup>, Tianyu Lei<sup>2,5</sup>, Tao Liu<sup>2,5</sup>, Zhenyi Wang<sup>2,3</sup>, Yuxian Li<sup>2,3</sup>, Tae-Ho Lee<sup>1</sup>, Jingping Li<sup>1</sup>, Haibao Tang<sup>6,7,8</sup>, Dianchuan Jin<sup>2,5</sup> and Andrew H. Paterson<sup>1,4</sup>



# POLIPLOIDÍA EN MEJORA GENÉTICA

- Gran parte de las **plantas cultivadas son poliploides** (alfalfa, algodón, patata, café, fresas, trigo, etc...).
- En algunos casos, se observa una correlación entre la poliploidía y el **proceso de domesticación**.
- Hay poliploides **naturales e inducidos** (colchicina).

# **POLIPLOIDÍA EN MEJORA GENÉTICA**

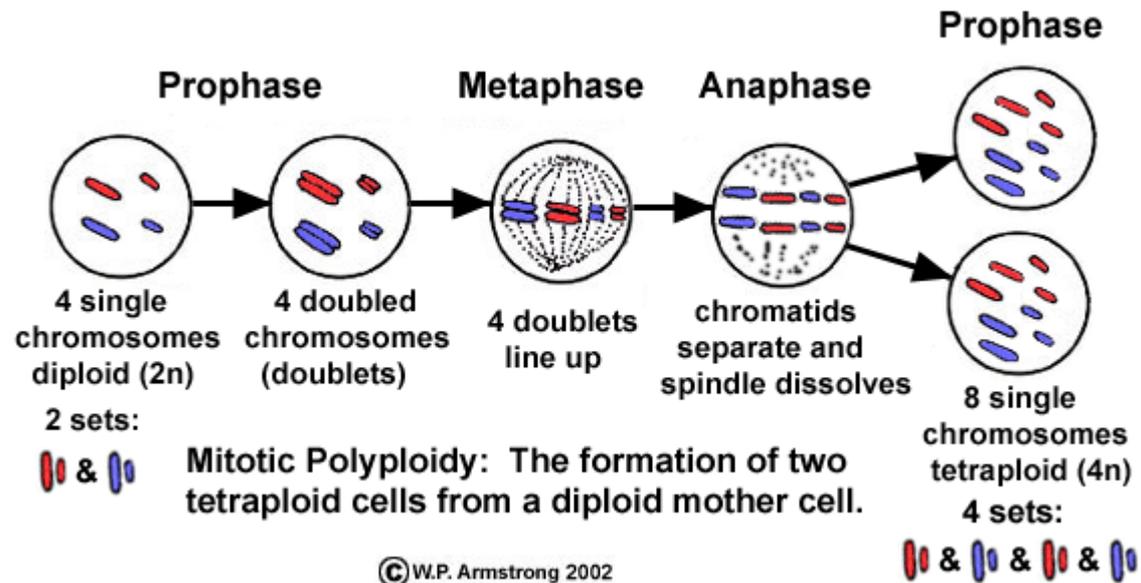
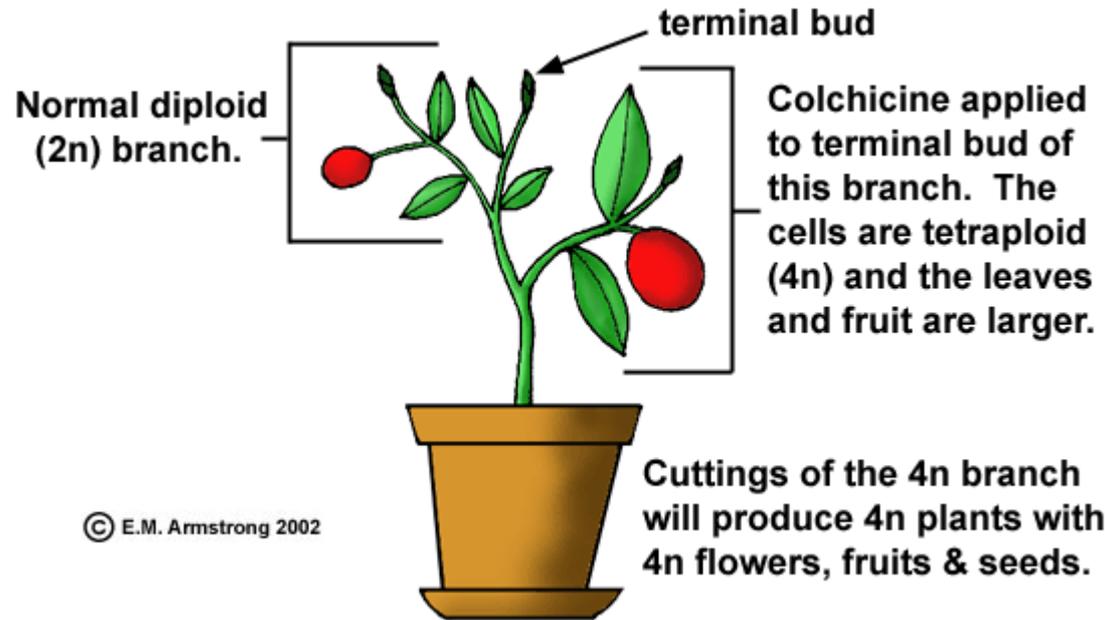
Mejora de plantas **autógamas**  
[Selección de **caracteres de interés**]

Mejora de plantas **alógamas**  
[Obtención de **híbridos**]

# POLIPLOIDÍA EN MEJORA GENÉTICA

- La poliploidía podría dar lugar a nuevos **rasgos con interés productivo** (fenotipo gigas).
- Se puede usar (o no) para recuperar la **fertilidad de híbridos**.
- Como vector para la **transferencia de genes** de unas especies/variedades a otras.

# Heterosis o vigor híbrido



# POLIPLOIDÍA EN MEJORA GENÉTICA



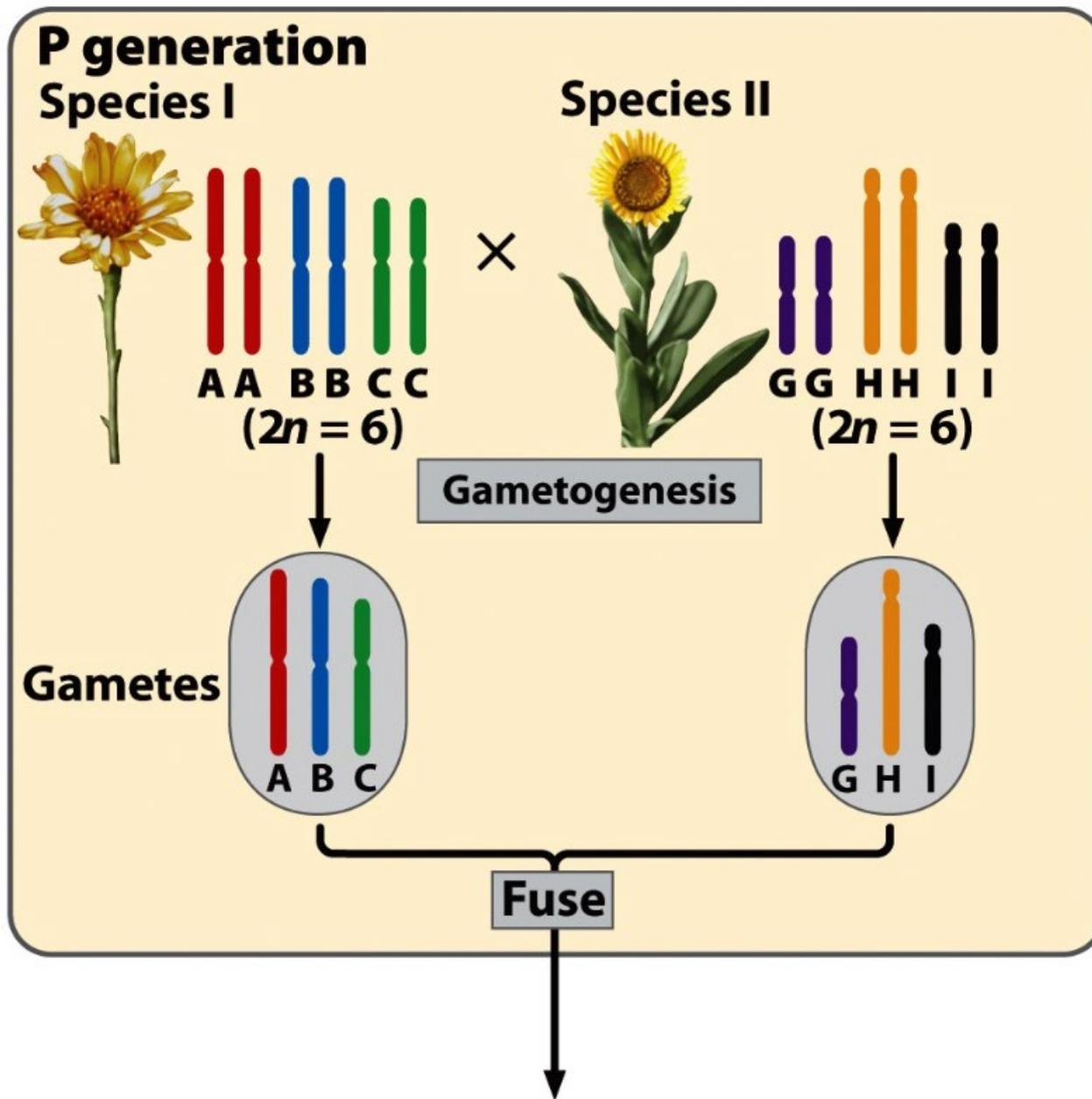


Figure 9-28 part 1  
*Genetics: A Conceptual Approach, Third Edition*  
© 2009 W.H. Freeman and Company

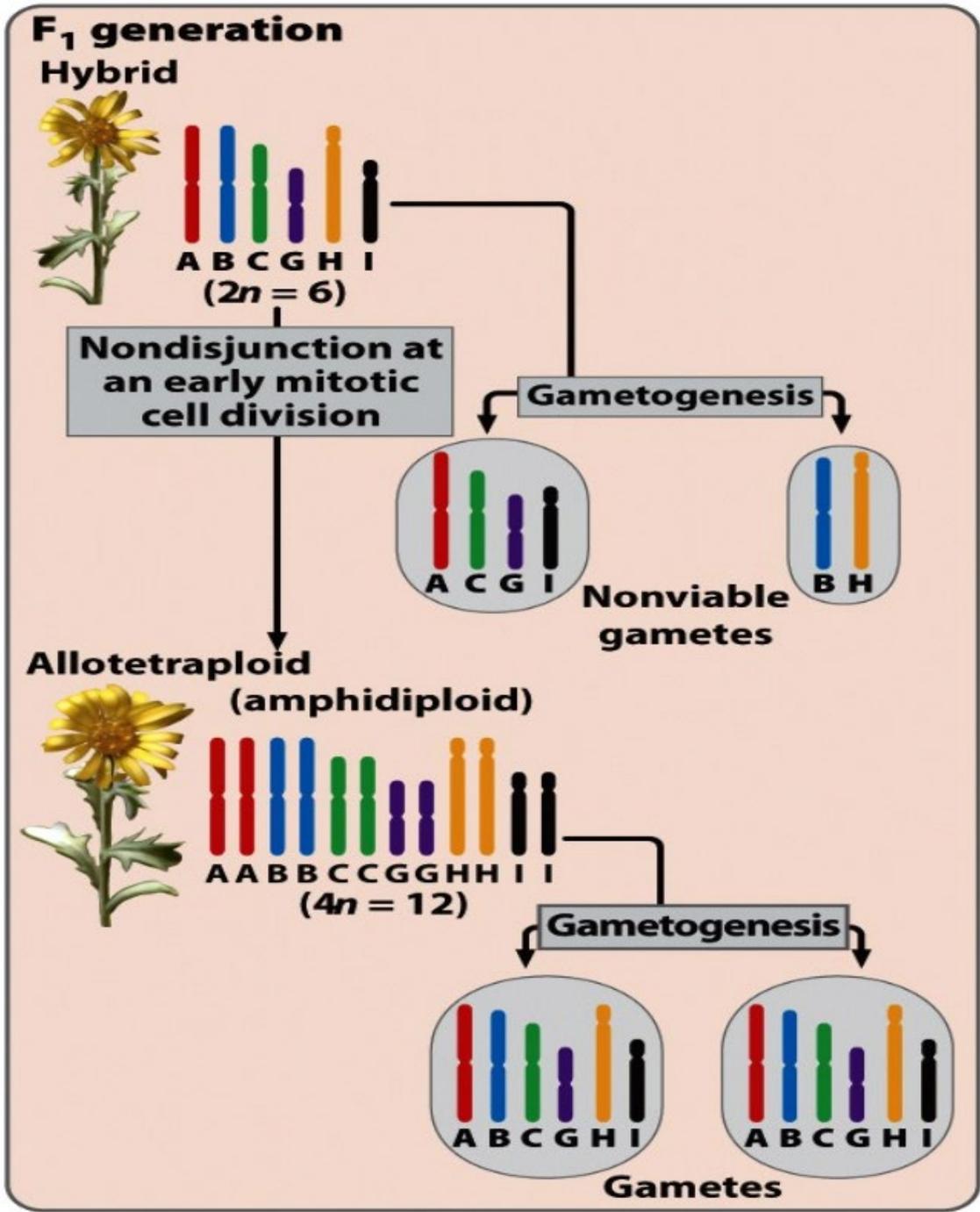
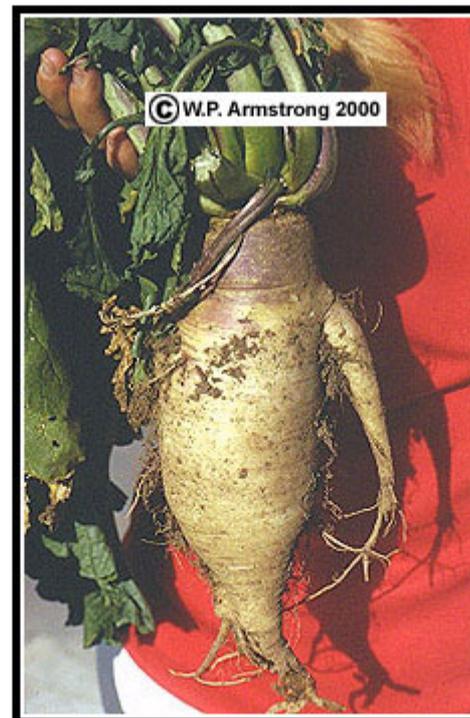
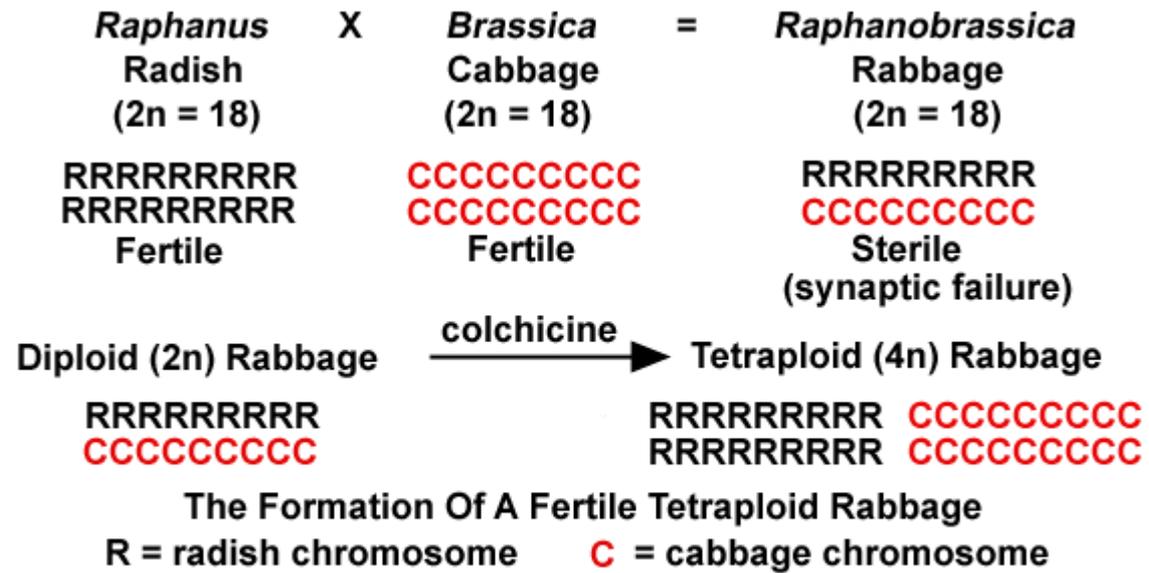


Figure 9-28 part 2  
*Genetics: A Conceptual Approach, Third Edition*  
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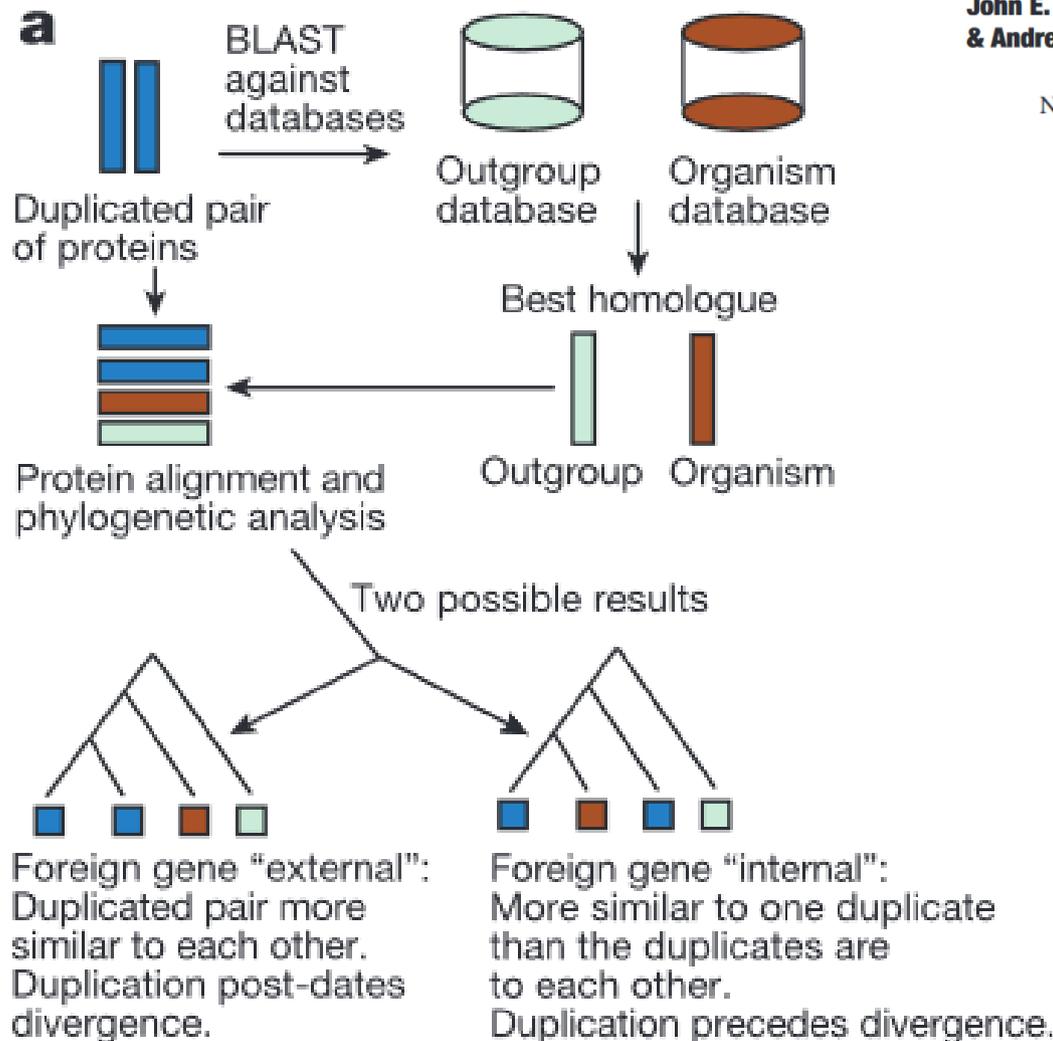
*Raphanus sativus*

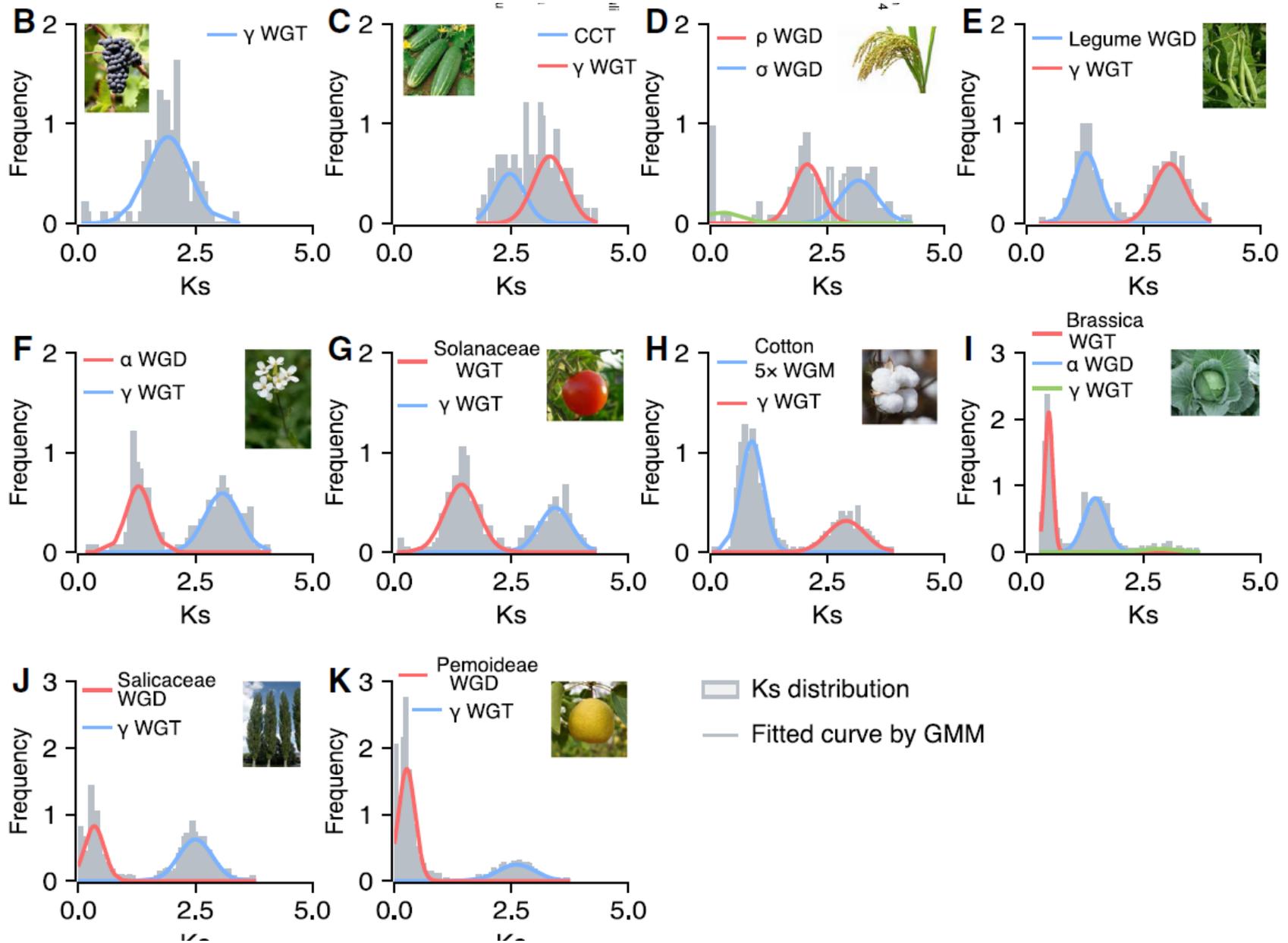
(Rábano x Col) + Poliploidización

# Unravelling angiosperm genome evolution by phylogenetic analysis of chromosomal duplication events

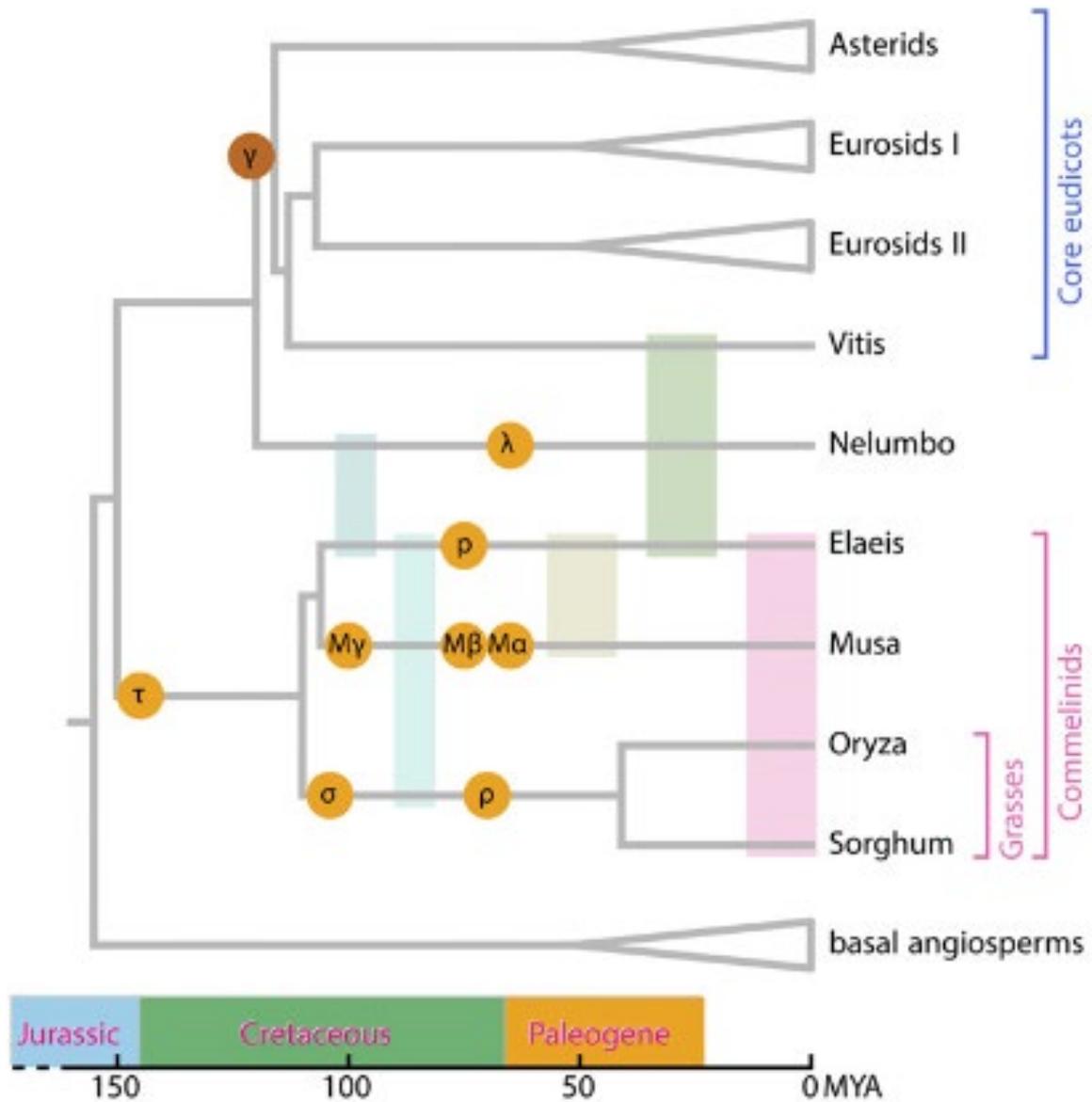
John E. Bowers\*, Brad A. Chapman\*, Junkang Rong  
& Andrew H. Paterson

NATURE | VOL 422 | 27 MARCH 2003 | [www.nature.com/nature](http://www.nature.com/nature)









The tragic ripples of  
an epic fraud p. 535

Insect pest profits from  
maize defenses pp. 642 & 654

Photoredox activation  
of methane pp. 647 & 658

# Science

\$15  
17 AUGUST 2018  
science.org

AAAS



## ROAD MAP FOR **WHEAT**

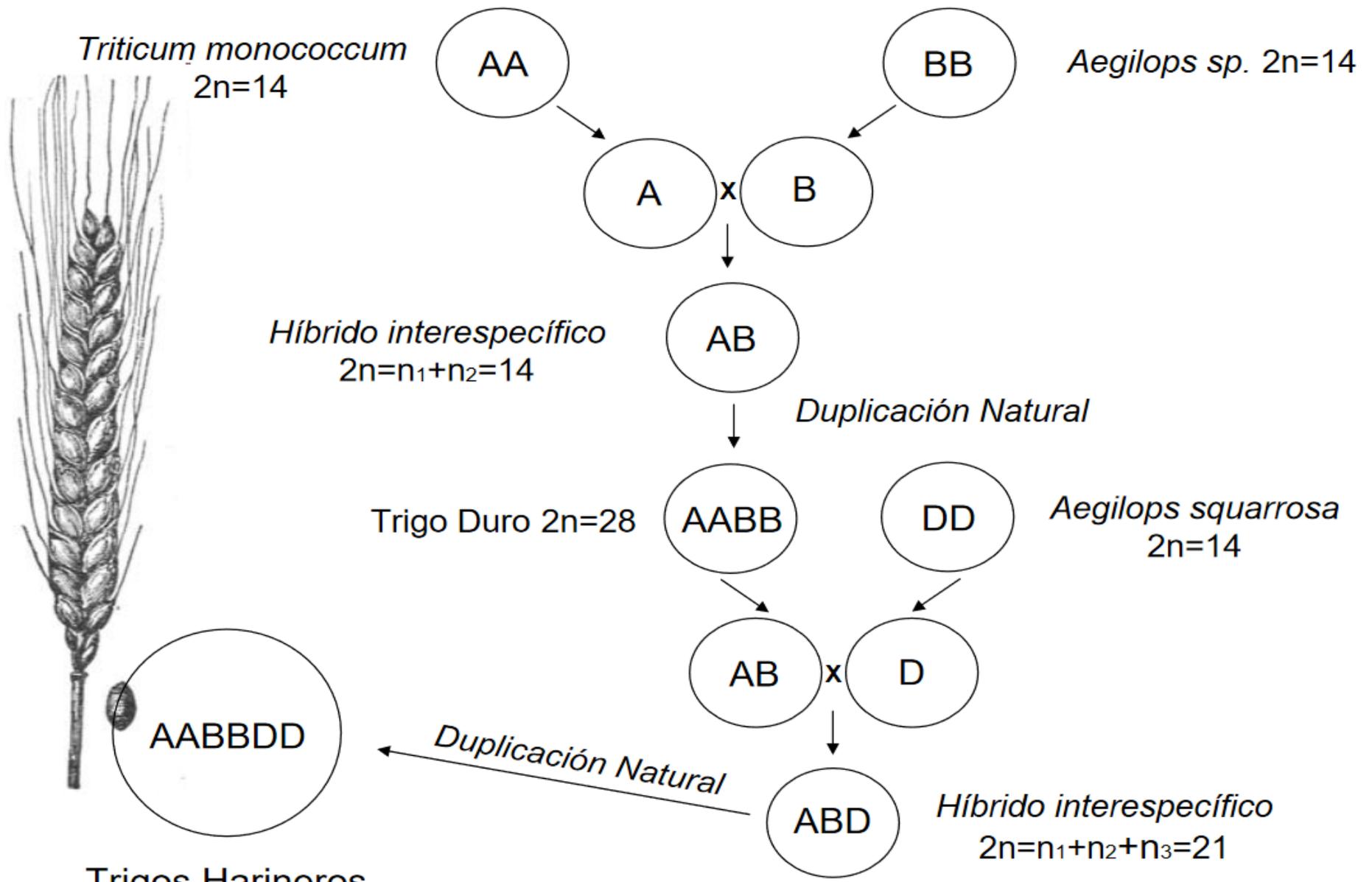
Ordered sequence will  
speed research pp. 635, 661, & 662

## The transcriptional landscape of polyploid wheat

R. H. Ramírez-González<sup>1,\*</sup>, P. Borrill<sup>1,\*†</sup>, D. Lang<sup>2</sup>, S. A. Harrington<sup>1</sup>, J. Brinton<sup>1</sup>, L. Venturini<sup>3</sup>, M. Davey<sup>4</sup>, J. Jacobs<sup>4</sup>, F. van...

\* See all authors and affiliations

Science 17 Aug 2018:  
Vol. 361, Issue 6403, eaar6089  
DOI: 10.1126/science.aar6089



**Trigos Harineros (hexaploides)**

- Triticum aestivum ssp. spelta*
- Triticum aestivum ssp. aestivum*
- Triticum aestivum ssp. vavilovii*

**Table 5.1** Economically important polyploids

Taxa	Natural/synthetic	Ploidy auto/allo
Wheat ( <i>Triticum aestivum</i> )	Synthetic	6X: Allopolyploid
Maize ( <i>Zea mays</i> )	Synthetic	4X: Autopolyploid
Oats ( <i>Avena sativa</i> )	Natural	6X: Allopolyploid
Alfalfa ( <i>Medicago sativa</i> )	Natural	4X: Autopolyploid
Sugarcane ( <i>Saccharum officinale</i> )	Natural	8X: Autopolyploid
Potato ( <i>Solanum tuberosum</i> )	Natural	4X: Autopolyploid
Sweet potato ( <i>Ipomoea batatas</i> )	Natural	6X: Autopolyploid
→ Banana ( <i>Musa sapientum</i> )	Natural	2X, 3X: Autopolyploid
Cotton ( <i>Gossypium hirsutum</i> )	Natural	4X: Allopolyploid
Tobacco ( <i>Nicotiana tabacum</i> )	Natural	4X: Allopolyploid
Coffee ( <i>Coffea arabica</i> )	Natural	4X: Allopolyploid
Apple ( <i>Malus domestica</i> )	Natural	2X, 3X: Allopolyploid
Pear ( <i>Pyrus communis</i> )	Natural	2X, 3X: Allopolyploid
Strawberry ( <i>Fragaria cascadiensis</i> )	Synthetic	10X: Allopolyploid
Peanut ( <i>Arachis hypogaea</i> )	Natural	4X: Allopolyploid
Watermelon ( <i>Citrullus lanatus</i> )	Synthetic	3X: Autopolyploid
Plum ( <i>Prunus domestica</i> )	Natural	6X: Allotetraploid
Rapeseed ( <i>Brassica napus</i> )	Synthetic	4X: Allotetraploid
Soybean ( <i>Glycine max</i> )	Natural	4X: Allopolyploid
Lime ( <i>Citrus aurantifolia</i> )	Natural	3X: Allopolyploid

# Effect of paleopolyploidy and allopolyploidy on gene expression in banana

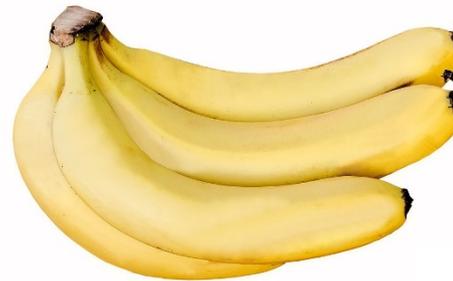


Alberto Cenci<sup>1\*</sup> , Yann Hueber<sup>1</sup>, Yasmin Zorrilla-Fontanesi<sup>2</sup>, Jelle van Wesemael<sup>2</sup>, Ewaut Kissel<sup>2</sup>, Marie Gislard<sup>3</sup>, Julie Sardos<sup>1</sup>, Rony Swennen<sup>2,4,5</sup>, Nicolas Roux<sup>1</sup>, Sebastien Christian Carpentier<sup>2,4</sup> and Mathieu Rouard<sup>1\*</sup>

Cenci et al. *BMC Genomics* (2019) 20:244  
<https://doi.org/10.1186/s12864-019-5618-0>



AAA: *Musa acuminata*. Cavendish



AAB: *Musa acuminata* and one by *Musa balbisiana*. Silk



ABB: *Musa acuminata* and two by *Musa balbisiana*. Blugoe subgroup



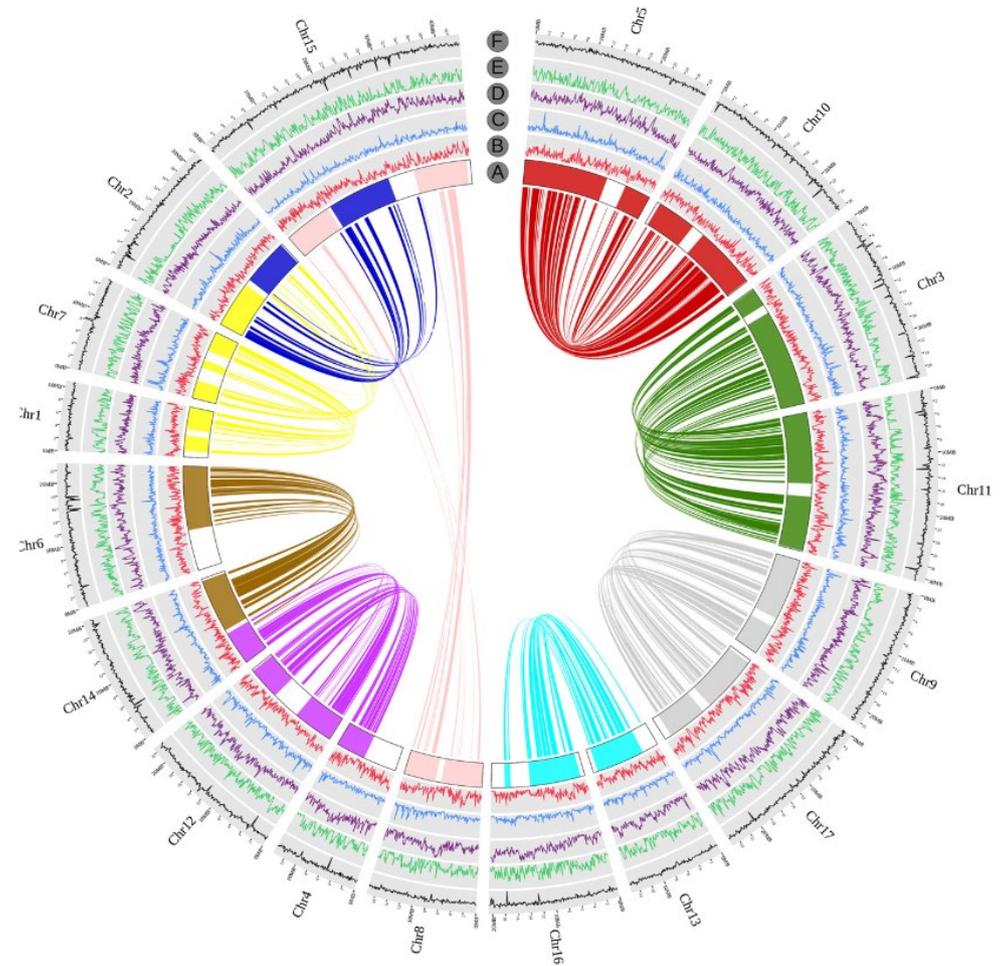
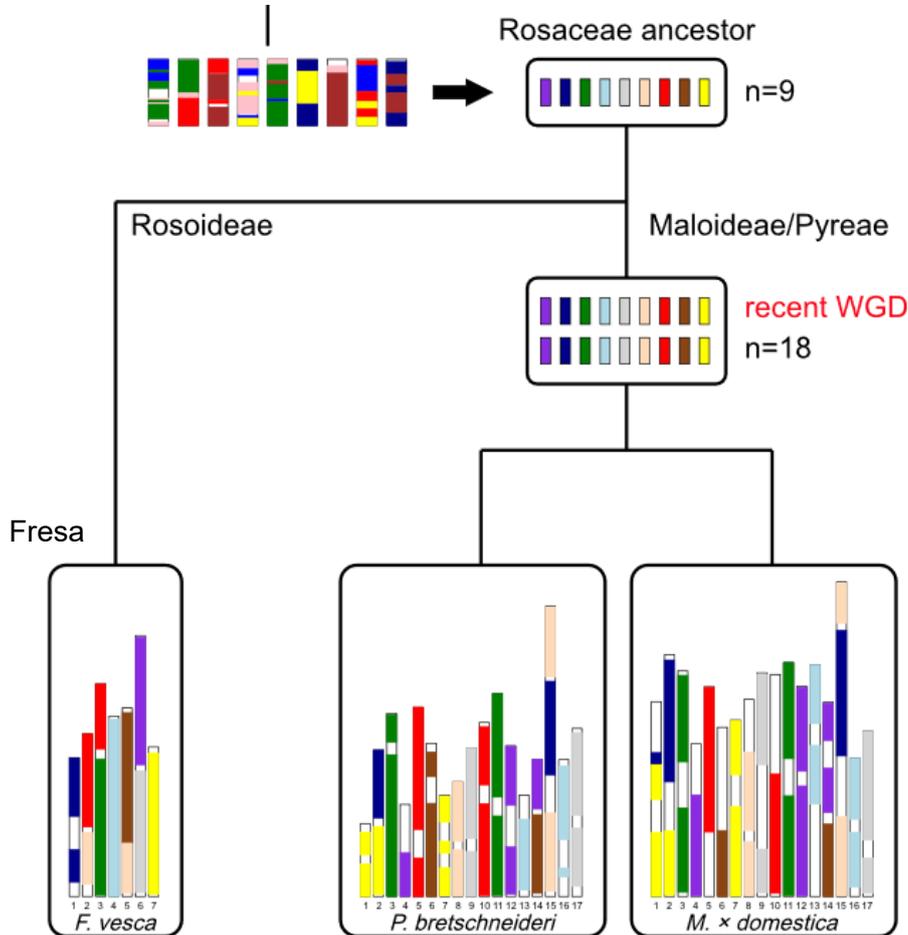
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Sugarcane ( <i>Saccharum officinale</i> )	Natural	8X: Autopolyploid
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# The genome of the pear (*Pyrus bretschneideri* Rehd.)

Jun Wu,<sup>1,11</sup> Zhiwen Wang,<sup>2,11</sup> Zebin Shi,<sup>3,11</sup> Shu Zhang,<sup>2,11</sup> Ray Ming,<sup>4,11</sup> Shilin Zhu,<sup>2,11</sup> M. Awais Khan,<sup>5</sup> Shutian Tao,<sup>1</sup> Schuyler S. Korban,<sup>5</sup> Hao Wang,<sup>6</sup> Nancy J. Chen,<sup>7</sup> Takeshi Nishio,<sup>8</sup> Xun Xu,<sup>2</sup> Lin Cong,<sup>2</sup> Kaijie Qi,<sup>1</sup> Xiaosan Huang,<sup>1</sup> Yingtao Wang,<sup>1</sup> Xiang Zhao,<sup>2</sup> Juyou Wu,<sup>1</sup> Cao Deng,<sup>2</sup> Caiyun Gou,<sup>2</sup> Weili Zhou,<sup>2</sup> Hao Yin,<sup>1</sup> Gaihua Qin,<sup>1</sup> Yuhui Sha,<sup>2</sup> Ye Tao,<sup>2</sup> Hui Chen,<sup>1</sup> Yanan Yang,<sup>1</sup> Yue Song,<sup>1</sup> Dongliang Zhan,<sup>2</sup> Juan Wang,<sup>2</sup> Leiting Li,<sup>1,4</sup> Meisong Dai,<sup>3</sup> Chao Gu,<sup>1</sup> Yuezhi Wang,<sup>3</sup> Daihu Shi,<sup>2</sup> Xiaowei Wang,<sup>2</sup> Huping Zhang,<sup>1</sup> Liang Zeng,<sup>2</sup> Danman Zheng,<sup>5</sup> Chunlei Wang,<sup>8</sup> Maoshan Chen,<sup>2</sup> Guangbiao Wang,<sup>2</sup> Lin Xie,<sup>2</sup> Valpuri Sovero,<sup>9</sup> Shoufeng Sha,<sup>1</sup> Wenjiang Huang,<sup>1</sup> Shujun Zhang,<sup>3</sup> Mingyue Zhang,<sup>1</sup> Jiangmei Sun,<sup>1</sup> Linlin Xu,<sup>1</sup> Yuan Li,<sup>1</sup> Xing Liu,<sup>1</sup> Qingsong Li,<sup>1</sup> Jiahui Shen,<sup>1</sup> Junyi Wang,<sup>2</sup> Robert E. Paull,<sup>7</sup> Jeffrey L. Bennetzen,<sup>6</sup> Jun Wang,<sup>2,10,12</sup> and Shaoling Zhang<sup>1,12</sup>

Genome Research 23:396-408 © 2013  
www.genome.org



**Pera y manzana**  
comparten una WGD  
ocurrida hace unos 30-45  
m.a.

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Alfalfa ( <i>Medicago sativa</i> )	Natural	4X: Autopolyploid
Sugarcane ( <i>Saccharum officinale</i> )	Natural	8X: Autopolyploid
Potato ( <i>Solanum tuberosum</i> )	Natural	4X: Autopolyploid
Sweet potato ( <i>Ipomoea batatas</i> )	Natural	6X: Autopolyploid
Banana ( <i>Musa sapientum</i> )	Natural	2X, 3X: Autopolyploid
Cotton ( <i>Gossypium hirsutum</i> )	Natural	4X: Allopolyploid
Tobacco ( <i>Nicotiana tabacum</i> )	Natural	4X: Allopolyploid
Coffee ( <i>Coffea arabica</i> )	Natural	4X: Allopolyploid
Apple ( <i>Malus domestica</i> )	Natural	2X, 3X: Allopolyploid
Pear ( <i>Pyrus communis</i> )	Natural	2X, 3X: Allopolyploid
Strawberry ( <i>Fragaria cascadiensis</i> )	Synthetic	10X: Allopolyploid
Peanut ( <i>Arachis hypogaea</i> )	Natural	4X: Allopolyploid
Watermelon ( <i>Citrullus lanatus</i> )	Synthetic	3X: Autopolyploid
Plum ( <i>Prunus domestica</i> )	Natural	6X: Allotetraploid
→ Rapeseed ( <i>Brassica napus</i> )	Synthetic	4X: Allotetraploid
Soybean ( <i>Glycine max</i> )	Natural	4X: Allopolyploid
Lime ( <i>Citrus aurantifolia</i> )	Natural	3X: Allopolyploid

$2n=18$

Yemas  
terminales

Inflorescencias

Yemas  
laterales

Tallos y  
flores

Tallo

Hojas

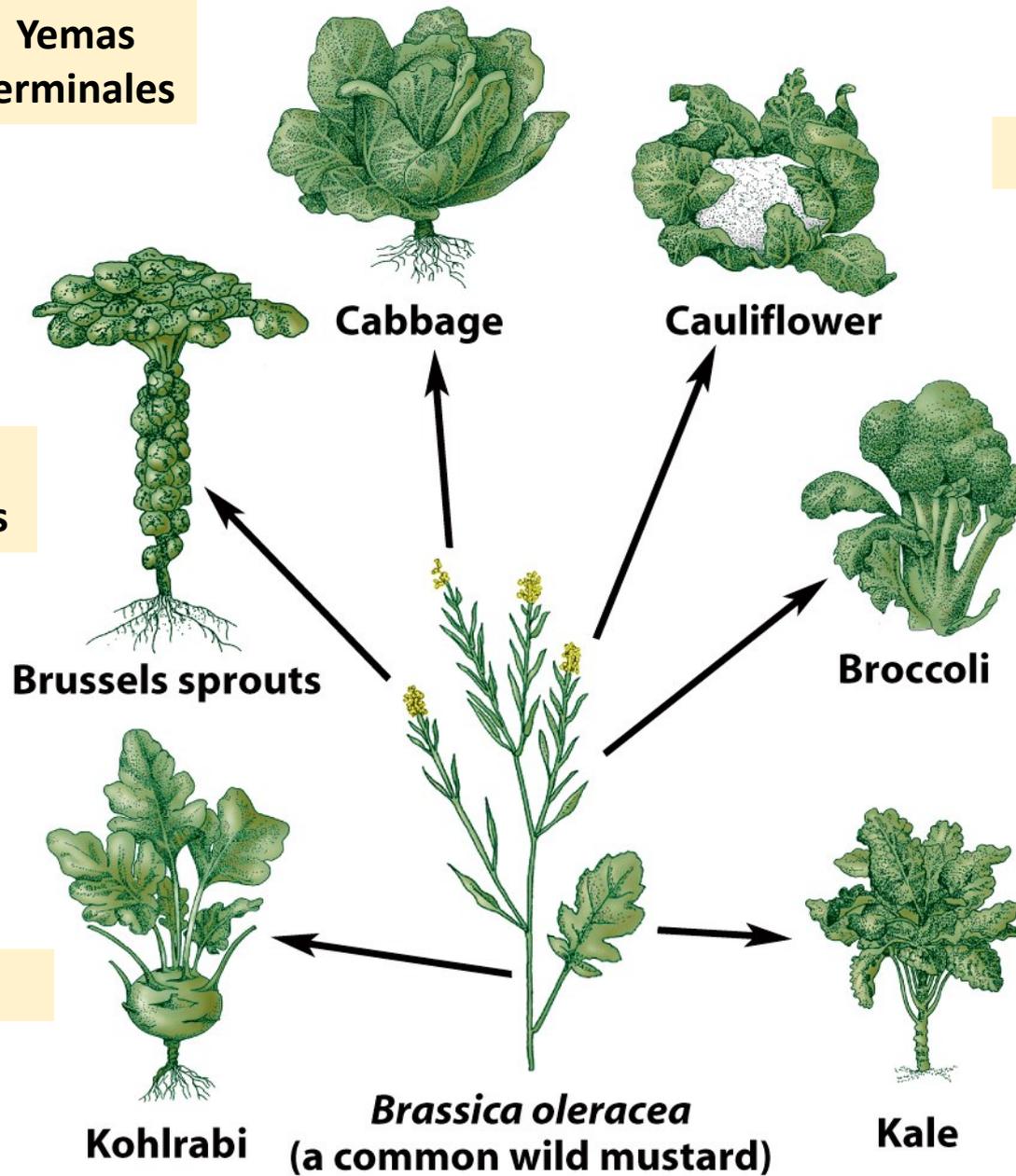
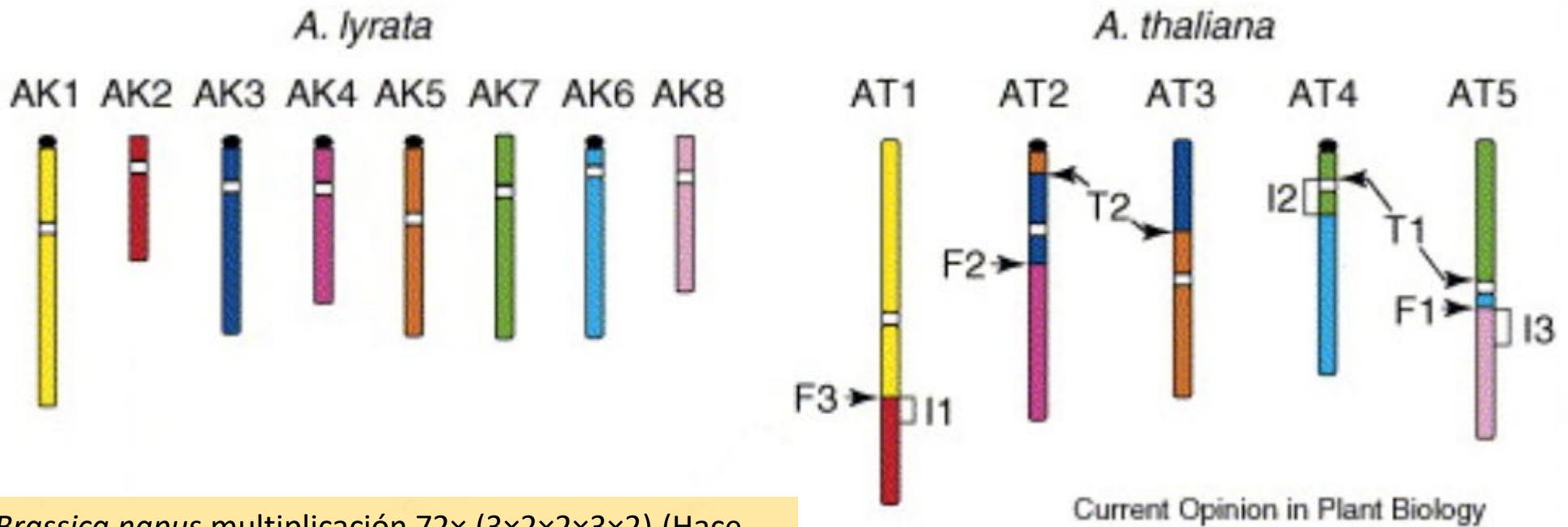
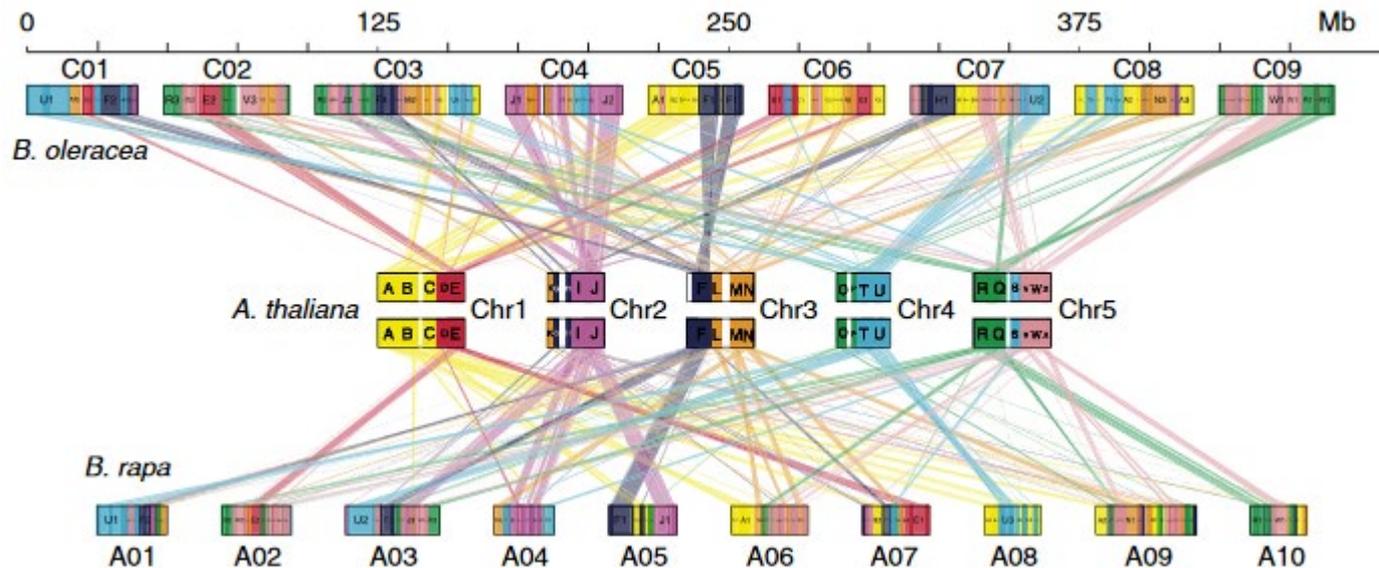


Figure 16-9 Discover Biology 3/e  
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*Brassica napus* multiplicación 72× (3×2×2×3×2) (Hace entre >100 m.a. a ~10,000 a.)



# Alopoliploidía

Berza



Mostaza negra



BB  
n=8

*Brassica nigra*

Mostaza parda



AABB  
n=10+8

*Brassica juncea*

BBCC  
n=9+8

*Brassica carinata*

CC  
n=9

*Brassica oleracea*

AACC  
n=10+9

*Brassica napus*

AA  
n=10

*Brassica rapa*

Colza



Col china, nabo

Coliflor, brocoli, col verde



**Table 5.1** Economically important polyploids

Taxa	Natural/synthetic	Ploidy auto/allo
Wheat ( <i>Triticum aestivum</i> )	Synthetic	6X: Allopolyploid
Maize ( <i>Zea mays</i> )	Synthetic	4X: Autopolyploid
Oats ( <i>Avena sativa</i> )	Natural	6X: Allopolyploid
Alfalfa ( <i>Medicago sativa</i> )	Natural	4X: Autopolyploid
Sugarcane ( <i>Saccharum officinale</i> )	Natural	8X: Autopolyploid
→ Potato ( <i>Solanum tuberosum</i> )	Natural	4X: Autopolyploid
Sweet potato ( <i>Ipomoea batatas</i> )	Natural	6X: Autopolyploid
Banana ( <i>Musa sapientum</i> )	Natural	2X, 3X: Autopolyploid
Cotton ( <i>Gossypium hirsutum</i> )	Natural	4X: Allopolyploid
Tobacco ( <i>Nicotiana tabacum</i> )	Natural	4X: Allopolyploid
Coffee ( <i>Coffea arabica</i> )	Natural	4X: Allopolyploid
Apple ( <i>Malus domestica</i> )	Natural	2X, 3X: Allopolyploid
Pear ( <i>Pyrus communis</i> )	Natural	2X, 3X: Allopolyploid
Strawberry ( <i>Fragaria cascadiensis</i> )	Synthetic	10X: Allopolyploid
Peanut ( <i>Arachis hypogaea</i> )	Natural	4X: Allopolyploid
Watermelon ( <i>Citrullus lanatus</i> )	Synthetic	3X: Autopolyploid
Plum ( <i>Prunus domestica</i> )	Natural	6X: Allotetraploid
Rapeseed ( <i>Brassica napus</i> )	Synthetic	4X: Allotetraploid
Soybean ( <i>Glycine max</i> )	Natural	4X: Allopolyploid
Lime ( <i>Citrus aurantifolia</i> )	Natural	3X: Allopolyploid

