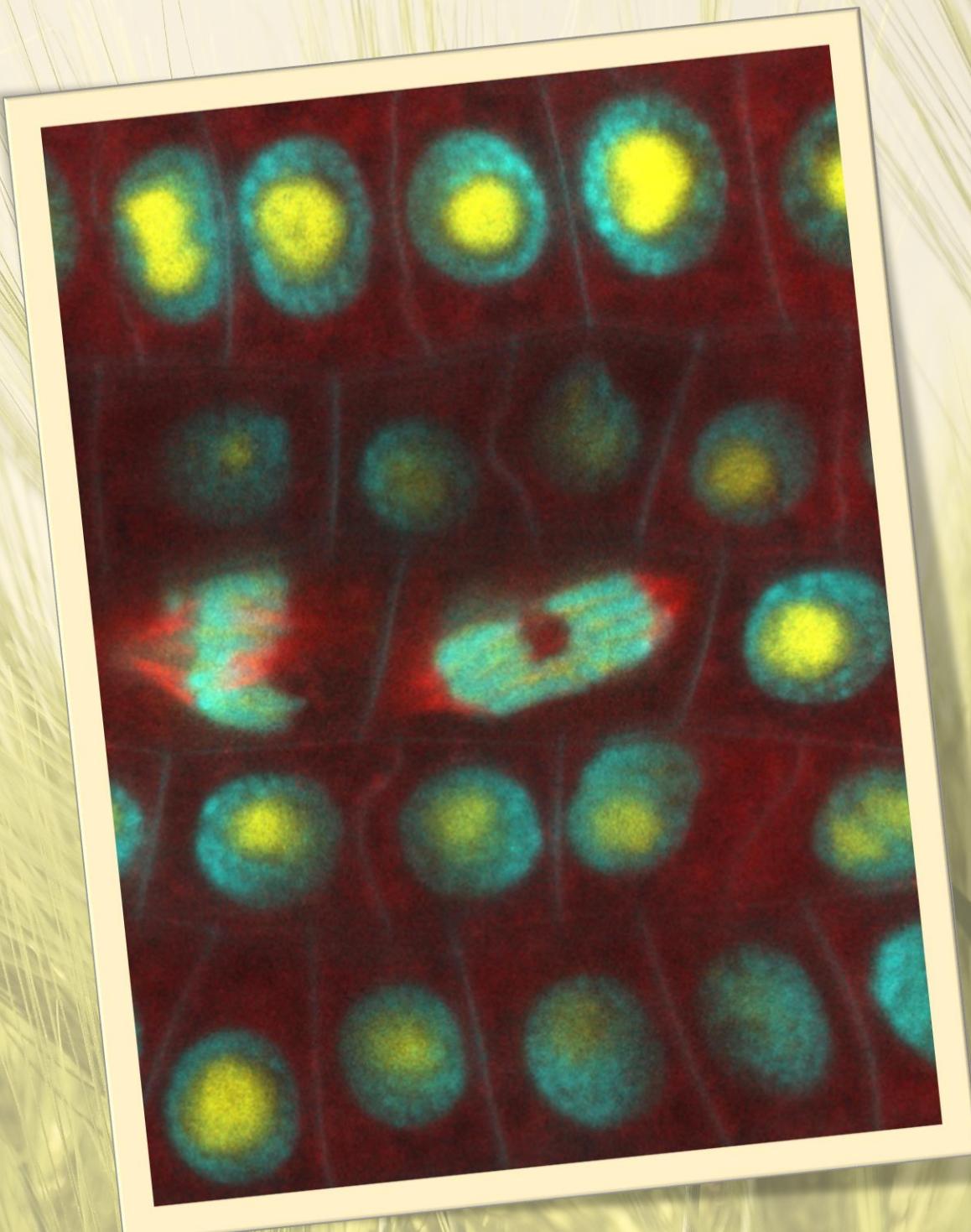


CHROMOSOME BIOLOGY

2024



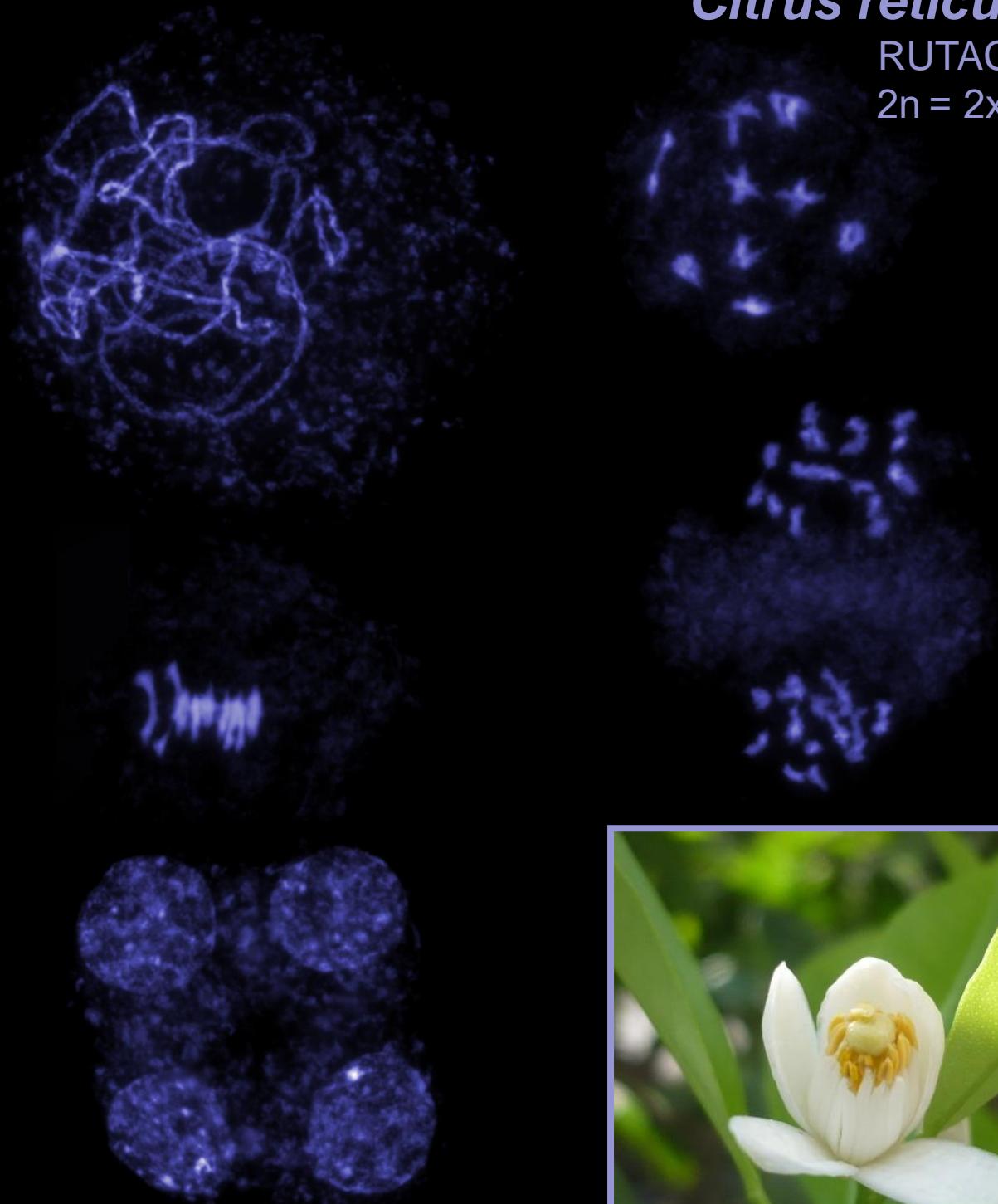
7th Edition

JANUARY

Citrus reticulata

RUTACEAE

$2n = 2x = 18$



Composition of different meiotic stages from diploid *Citrus reticulata* by spreading technique with DAPI staining. The production of triploid citrus hybrids is relevant to get new seed-less varieties.



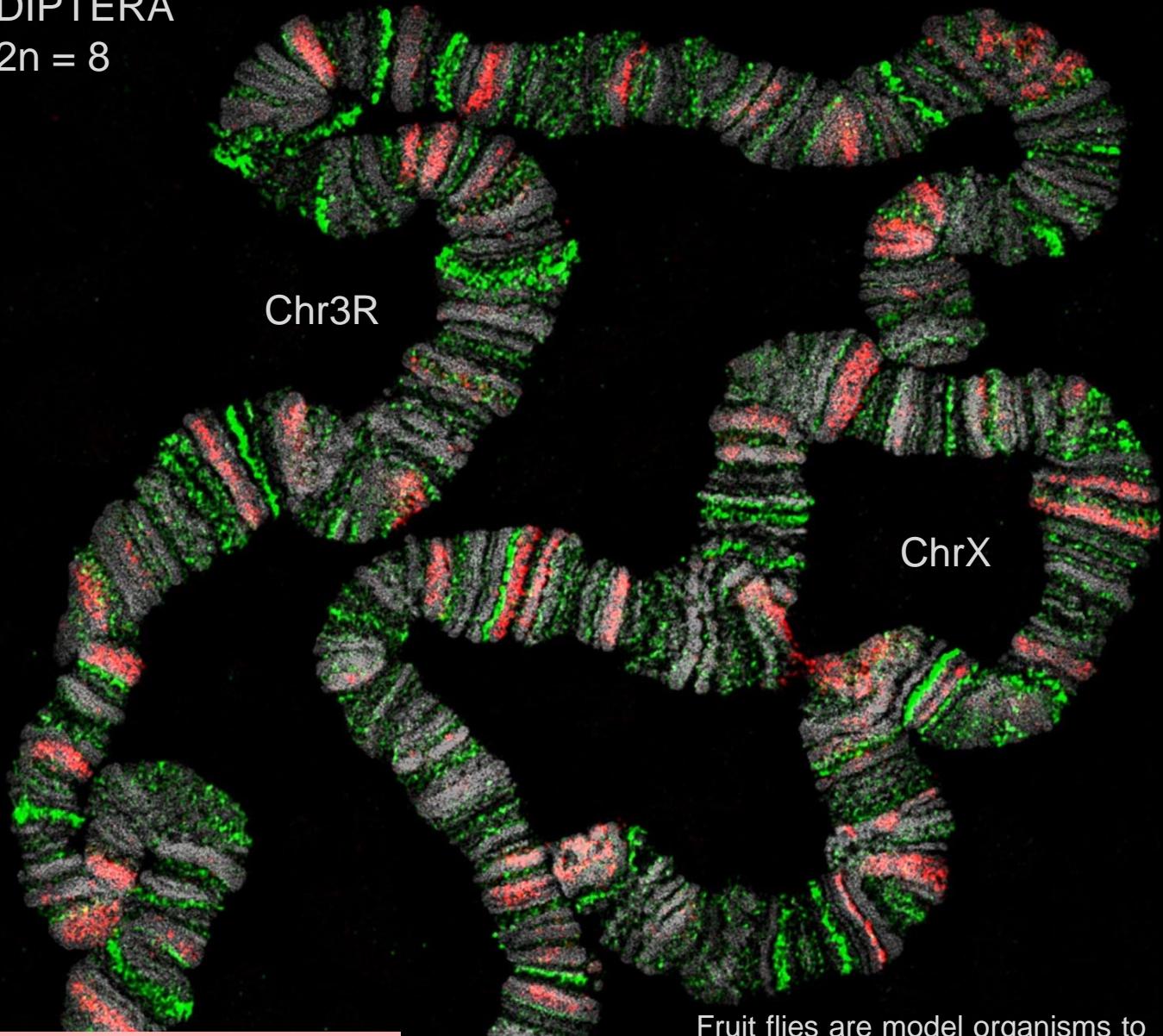
Nadia Fernández-Jiménez and Mónica Pradillo (Universidad Complutense de Madrid, Spain)
Andrés García-Lor and Pablo Aleza (Instituto Valenciano de Investigaciones Agrarias, Valencia, Spain)

FEBRUARY

Drosophila melanogaster

DIPTERA

2n = 8



Fruit flies are model organisms to analyze transcription (green) and replication (red) in polytene chromosomes comprising altogether 2048 chromatids aligned in both paired homologues.

5 μ m

Tatyana Kolesnikova, Viktoria Dovgan and Veit Schubert (Institute of Molecular and Cellular Biology, Novosibirsk, Russia; IPK Gatersleben, Germany)

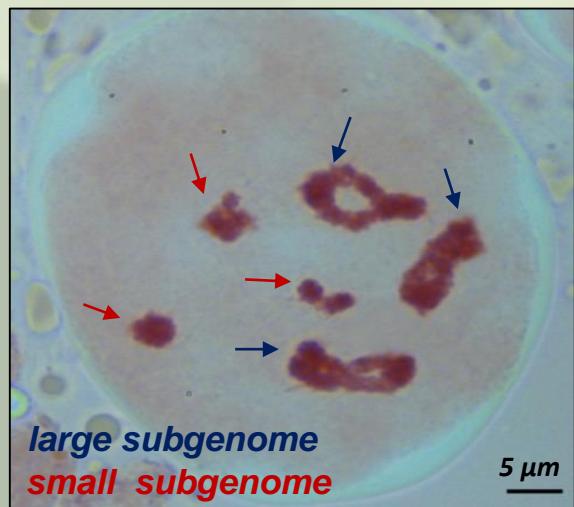
123 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29

MARCH

Dipcadi goaense

ASPARAGACEAE

$2n = 2x = 12$



Top: Mitotic metaphase of *D. goaense* showing $2n = 12$ chromosomes and a bimodal karyotype.

Bottom: Pollen mother cell at diakinesis showing $n = 6$ bivalents



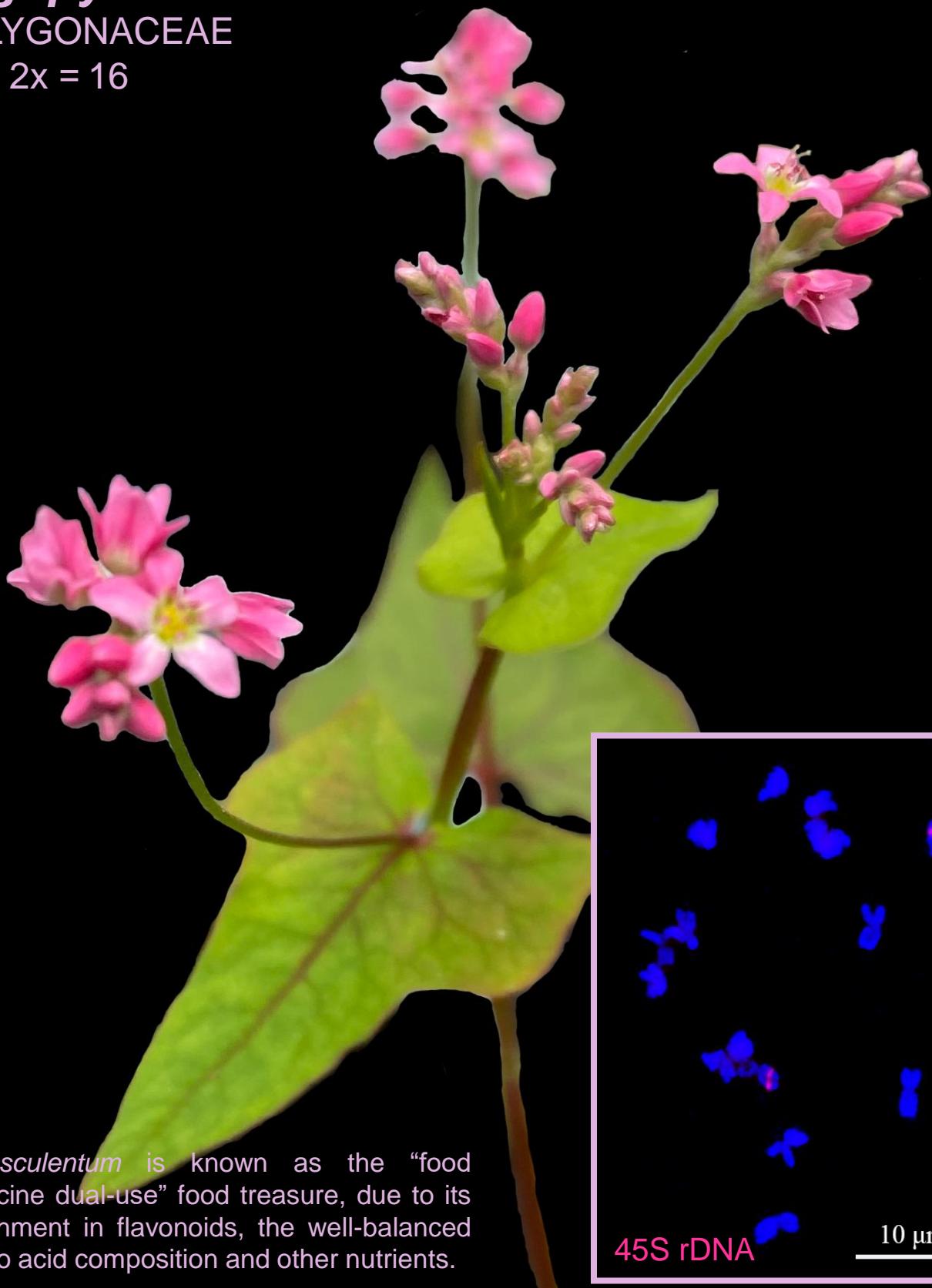
Priya E. Shelke, Shrirang R. Yadav and Manoj M. Lekhak (Shivaji University, Kolhapur, India)

APRIL

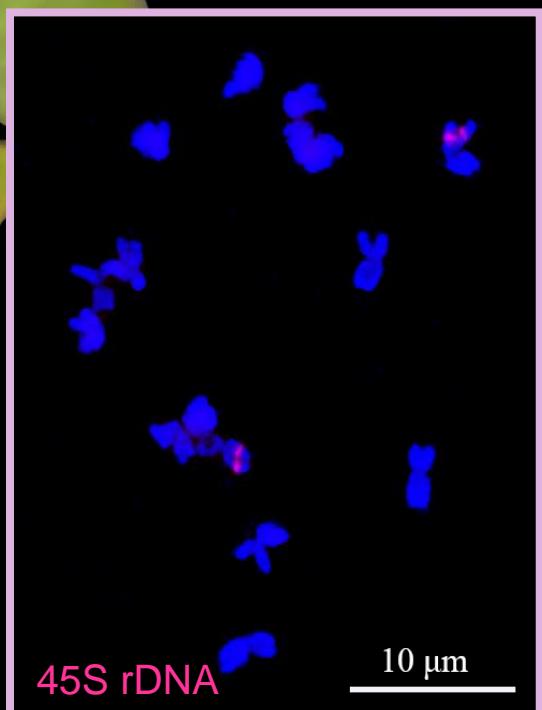
Fagopyrum esculentum

POLYGONACEAE

$2n = 2x = 16$



F. esculentum is known as the “food medicine dual-use” food treasure, due to its enrichment in flavonoids, the well-balanced amino acid composition and other nutrients.



Wen Zheng and Dandan Wu (Sichuan Agricultural University, China)

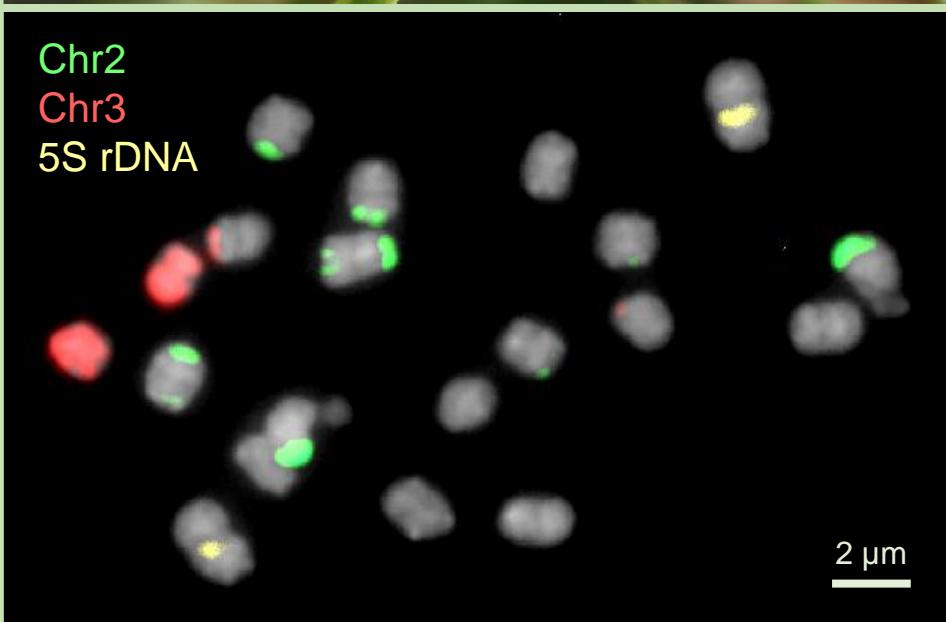
1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30

MAY

Phaseolus angustissimus

FABACEAE

$2n = 2x = 20$



Oligo painting probes designed from the common bean genome for the orthologous Chr2 (in green) and Chr3 (in red) show translocations for both chromosomes in the genome of *P. angustissimus*.

Thiago Nascimento, André Marques and Andrea Pedrosa-Harand (Federal University of Pernambuco, Brazil and Max-Planck Institute for Plant Breading Research, Cologne, Germany)

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31

JUNE

Arabidopsis thaliana

BRASSICACEAE

2n = 10



Maria Cuacos and Stefan Heckmann (IPK Gatersleben, Germany)

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30

Meiosis occurs inside the female gametophyte embedded within multiple cell layers, making its investigation a challenge. Here, female meiotic cells are visualized by expression of the meiotic axis protein ASY1 fused to mRuby2 (red). This line also expresses CENH3 fused to mTurquoise2 (blue).

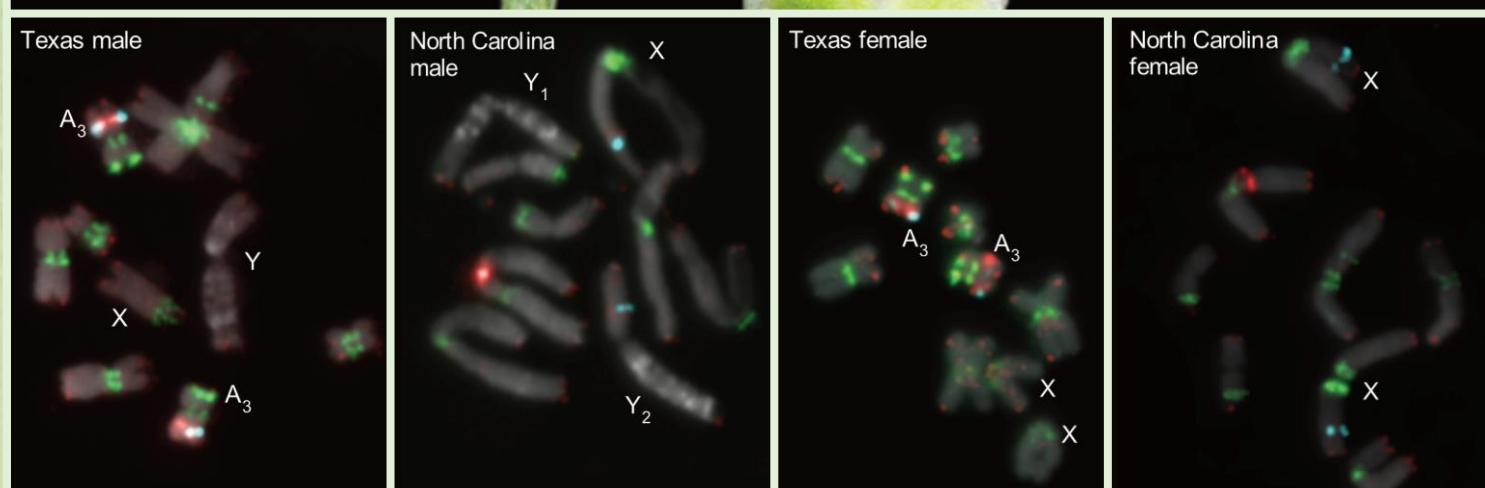
50 μ m

JULY

Rumex hastatulus POLYGONACEAE

R. hastatulus is a dioecious weed native to North America. This species has two cytologically distinct populations that vary in their heteromorphic sex chromosomes.

FISH with 5S rDNA (cyan), pericentromeric repeat (green) and telomere-specific (red) probes. ▼



Jana Kružlicová, Roman Hobza (Institute of Biophysics CAS, Brno, Czech Republic) and Stephen Wright (University of Toronto, Canada)

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31

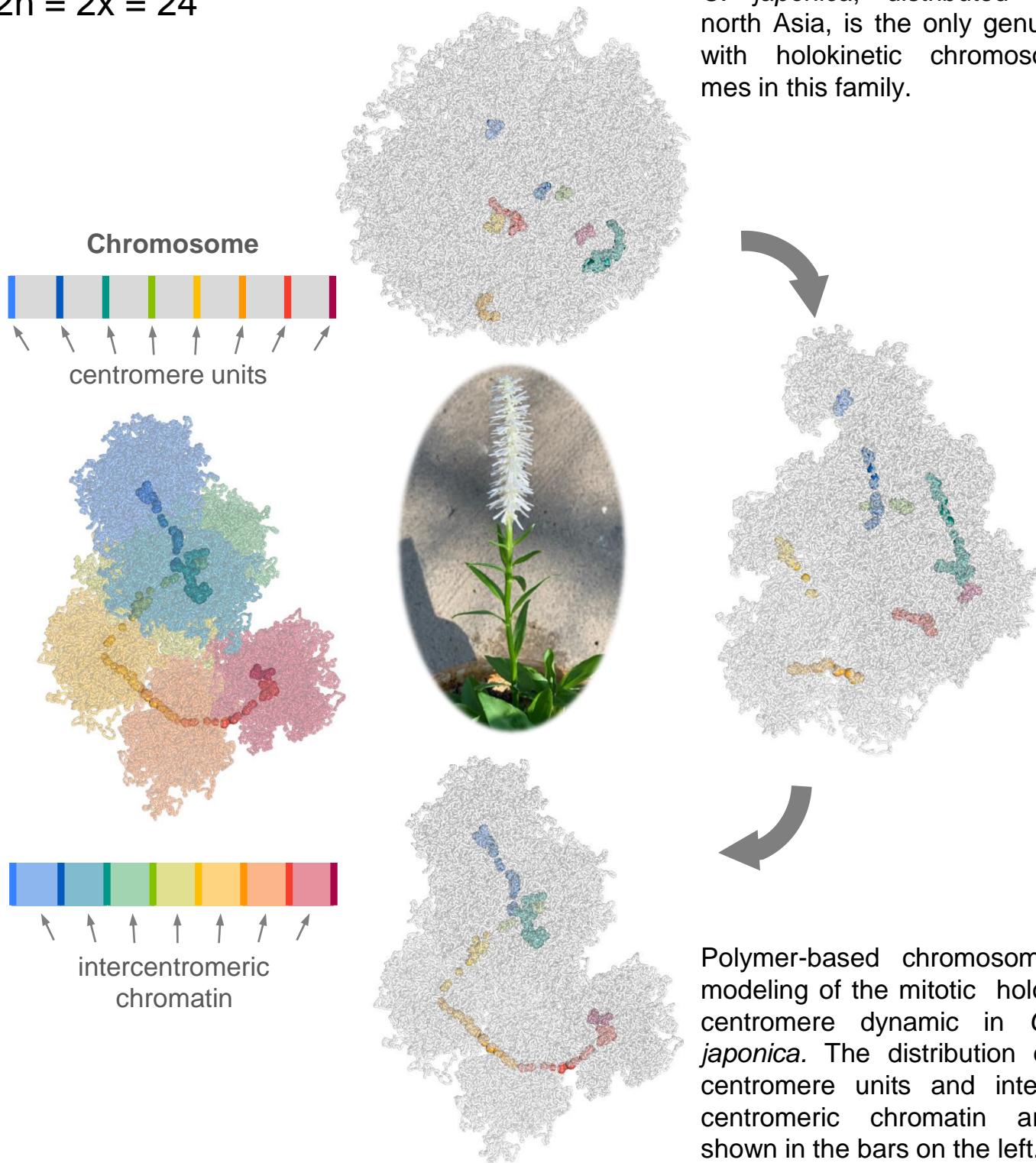
AUGUST

Chionographis japonica

MELANTHIACEA

$2n = 2x = 24$

C. japonica, distributed in north Asia, is the only genus with holokinetic chromosomes in this family.



Polymer-based chromosome modeling of the mitotic holocentromere dynamic in *C. japonica*. The distribution of centromere units and intercentromeric chromatin are shown in the bars on the left.

Amanda Câmara, Yi-Tzu Kuo and Andreas Houben (IPK Gatersleben, Germany)

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31

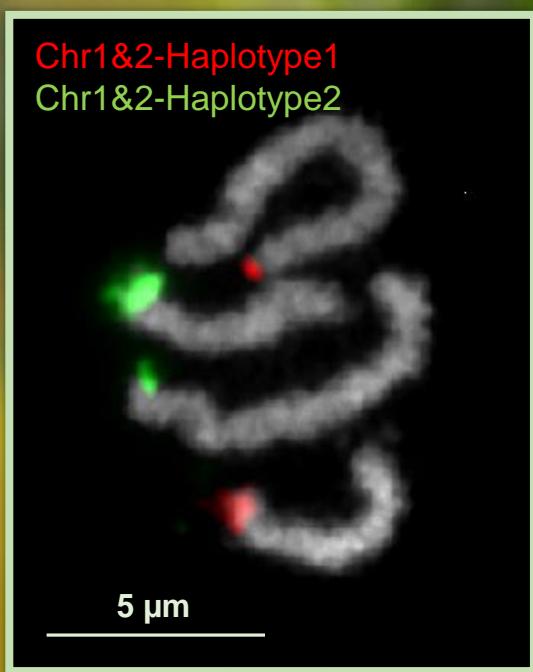
SEPTEMBER

Rhynchospora tenuis

CYPERACEAE

$2n = 2x = 4$

R. tenuis is a holocentric beaksedge showing achiasmatic meiosis. It presents the lowest chromosome number reported for flowering plants ($2n = 4$), which makes it a model candidate in the study of meiotic adaptations and karyotype evolution in holocentric species.



◀ Haplotype-specific oligo probes showing a complex translocation between the non-homologous chromosomes of *R. tenuis*. Probes in red are specific for the haplotype 1 and probes in green specific to haplotype 2.

Thiago Nascimento and André Marques (Max-Planck Institute for Plant Breeding Research, Cologne, Germany)

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30

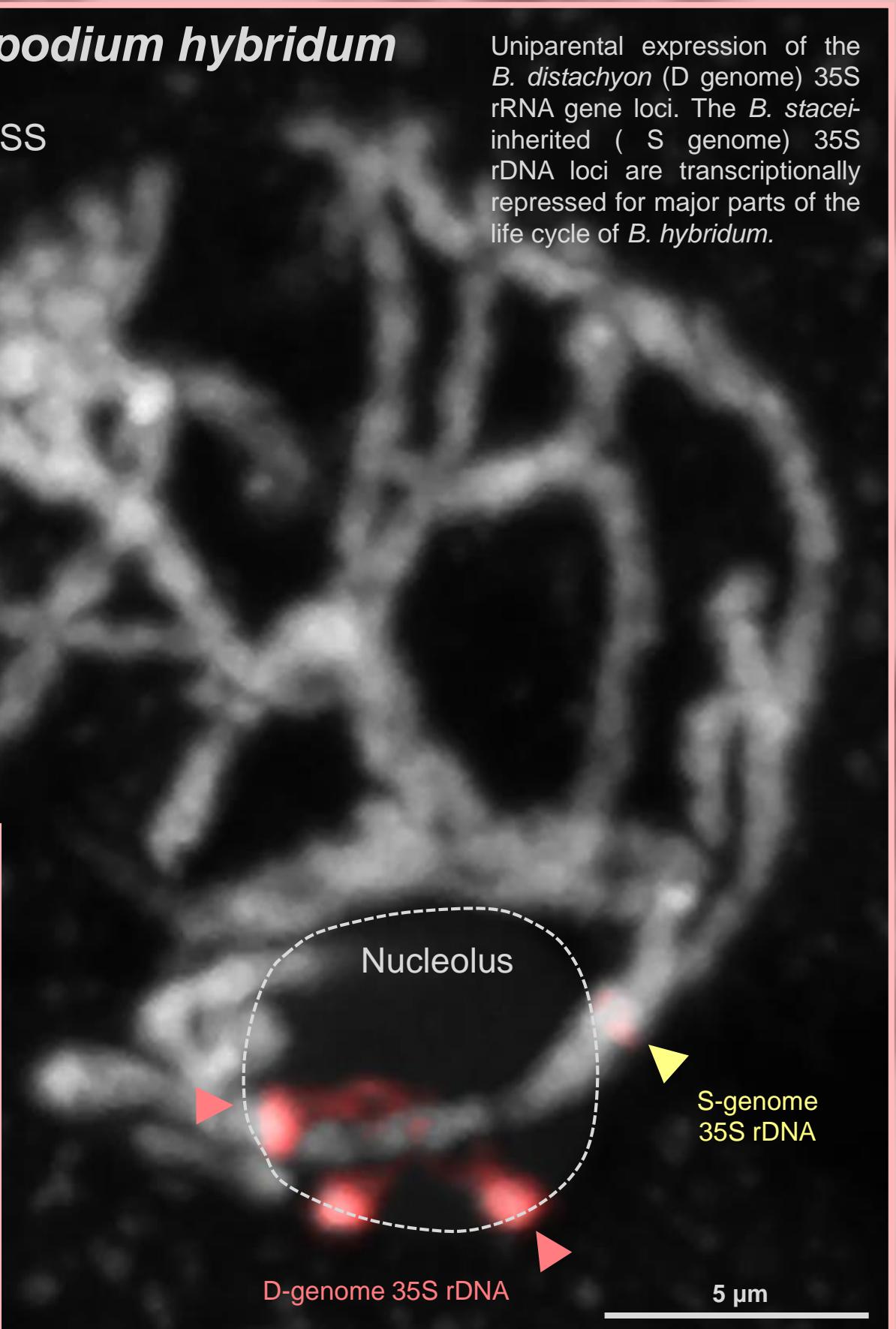
OCTOBER

Brachypodium hybridum

POACEAE

2n=30, DDSS

Uniparental expression of the *B. distachyon* (D genome) 35S rRNA gene loci. The *B. stacei*-inherited (S genome) 35S rDNA loci are transcriptionally repressed for major parts of the life cycle of *B. hybridum*.



Natalia Borowska-Żuchowska, Ewa Robaszkiewicz and Robert Hasterok (University of Silesia in Katowice, Poland)

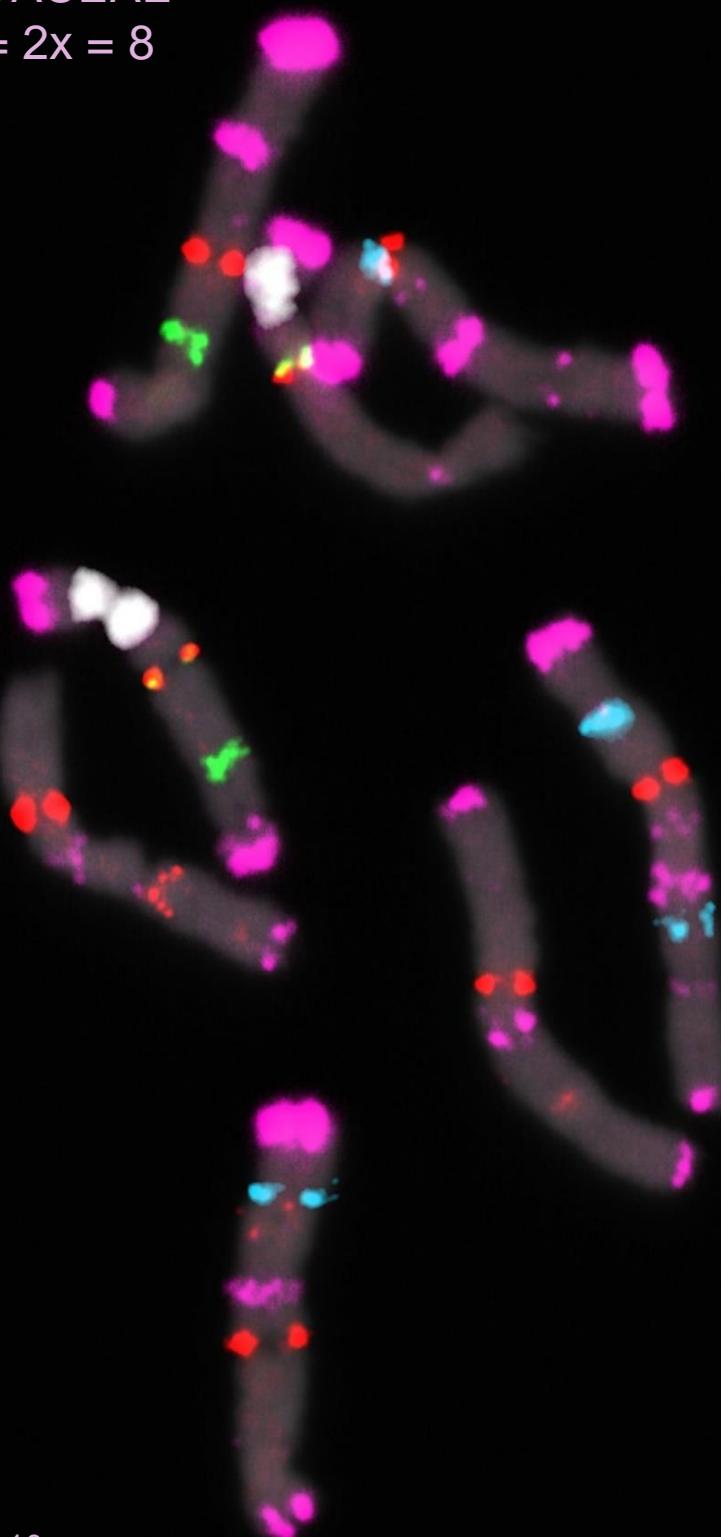
1 2 3 4 5 **6** 7 8 9 10 11 12 **13** 14 15 16 17 18 19 **20** 21 22 23 24 25 26 **27** 28 29 30 31

NOVEMBER

Crocus vernus

IRIDACEAE

$2n = 2x = 8$



10 μm



The spring crocus (*C. vernus*) is growing in alpine meadows of the northern Pyrenees, the Alps and the Dinaric Alps. An interesting cytogenetic feature of *C. vernus* is its high heterozygous karyotypes.

5S rDNA

45S rDNA

CI60

CI154

CI188 (centromeric repeat)

Nomar Espinosa Waminal, Frank R. Blattner and Dörte Harpke (IPK, Gatersleben, Germany)

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30

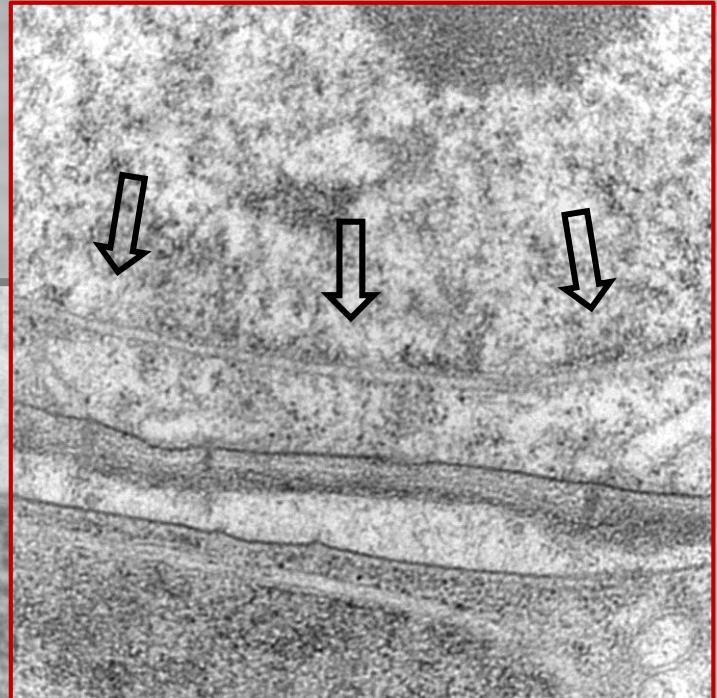
DECEMBER

Myristica fragrans

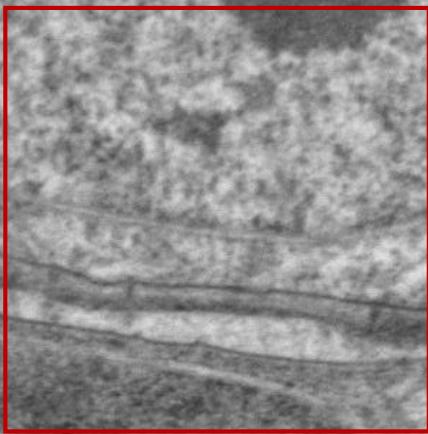
MYRISTICACEAE

$2n = 2x = 44$

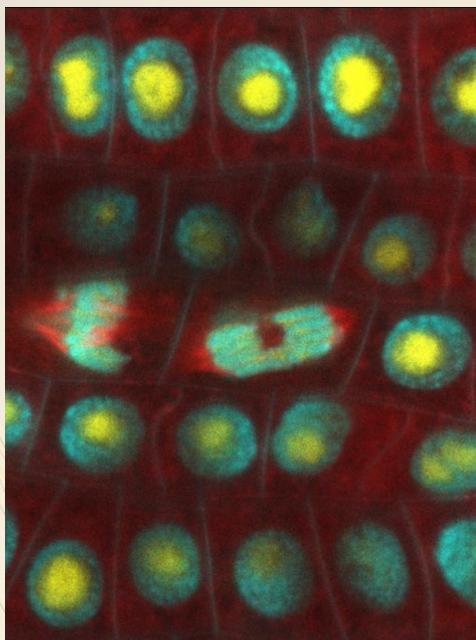
Nutmeg seed and red aril within fruit.



Interphase nucleus of *M. fragrans* analysed by transmission electron microscopy. Arrows indicate the nuclear membrane in the further enlarged picture.



Michael Melzer, Yi-Tzu Kuo, Jacob G. Kurian and Andreas Houben (IISER Thiruvananthapuram, India; IPK, Gatersleben, Germany)



Cover picture

Mitosis in living barley plant visualized with a multi-marker fluorescent line for chromatin (CFP-H2B), nucleolus (EYFP-FIB1) and microtubules (mCHERRY-TUA3).

Kateřina Kaduchová and Aleš Pečinka (IEB, Olomouc, Czech Republic)

Acknowledgement

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Jörg Fuchs and Andreas Houben (IPK, Gatersleben, Germany)



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