Plant of the Day



Cyperus esculentus - Cyperaceae Chufa (tigernut) Top Crop for calory production per square meter One of the world's worst weeds

Big Questions

Is polyploidy an evolutionary dead-end?

If so, why are all land plants the products of multiple polyploidization events?

How do polyploid genomes diploidize (i.e., what are the rules)?

Paleopolyploidy

Ancient whole genome duplication

No different from neopolyploidy – except that it happened a long time ago

Track the historical contribution of polyploid speciation to evolution

Conventional views about the evolutionary role of polyploidy

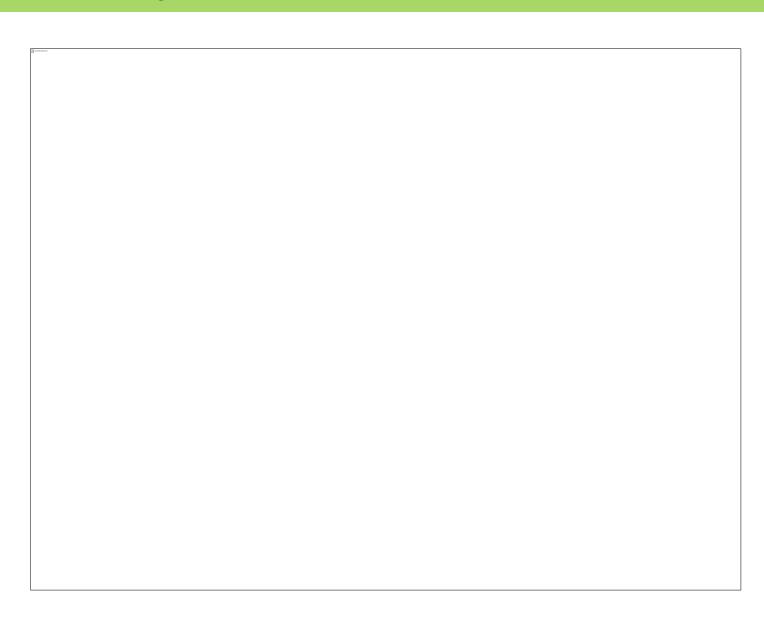
Polyploidy = Evolutionary noise (1970)

W. H. Wagner, Jr.

"...polyploidy has contributed little to progressive evolution" (1971)

G. L. Stebbins

Diploidization – the process of converting a polyploid genome into a diploid one.



Diploidization

Obscures evidence of paleopolyploidy

Return to a diploid genetic system

- Restoration of full bivalent pairing
- Gene and chromosome loss
- Chromosomal rearrangements

Proceeds at different rates in different lineages

Methods for Identifying Paleopolyploidy

Fossils

Synteny relationships of duplicated genes

conserved gene order

Age estimates of duplicate genes

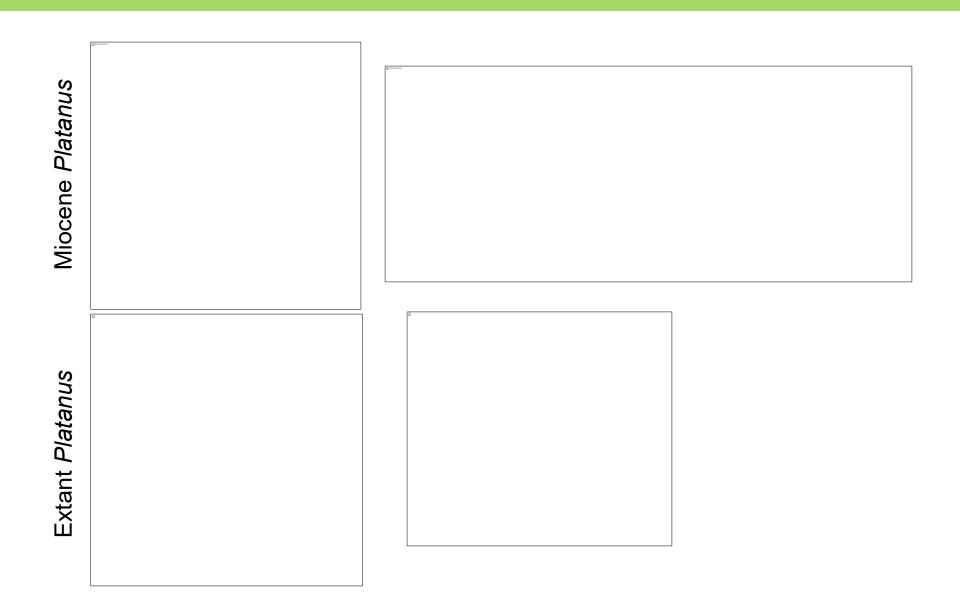
Cell Size Increase

Consequence of genome size increase

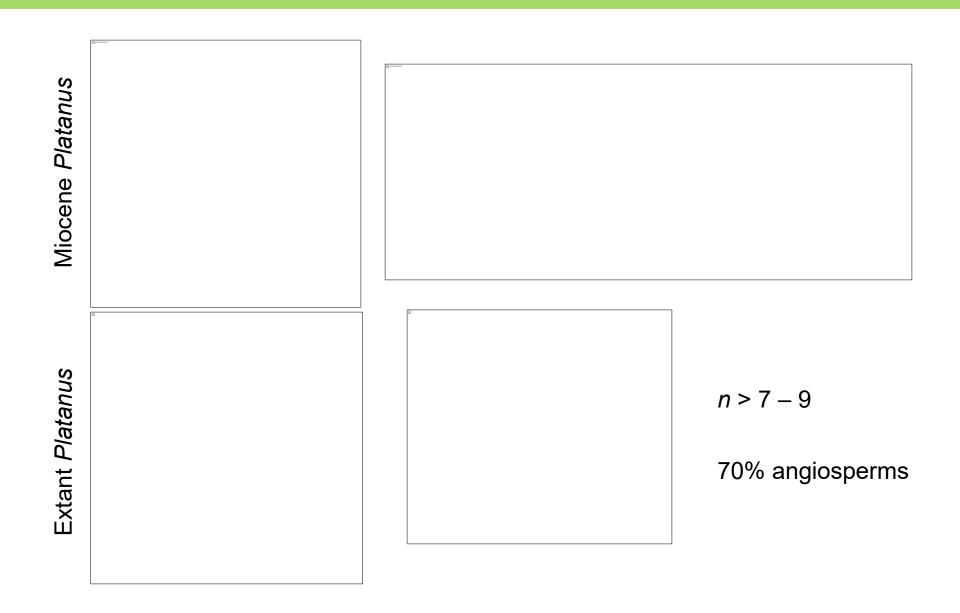
2 X increase in cell volume

 1.58 X increase in cell surface area 2n

Fossil Estimates



Fossil Estimates



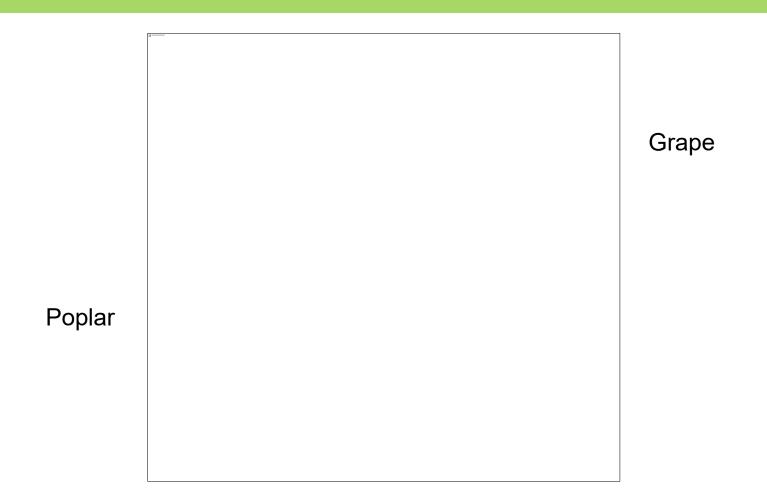






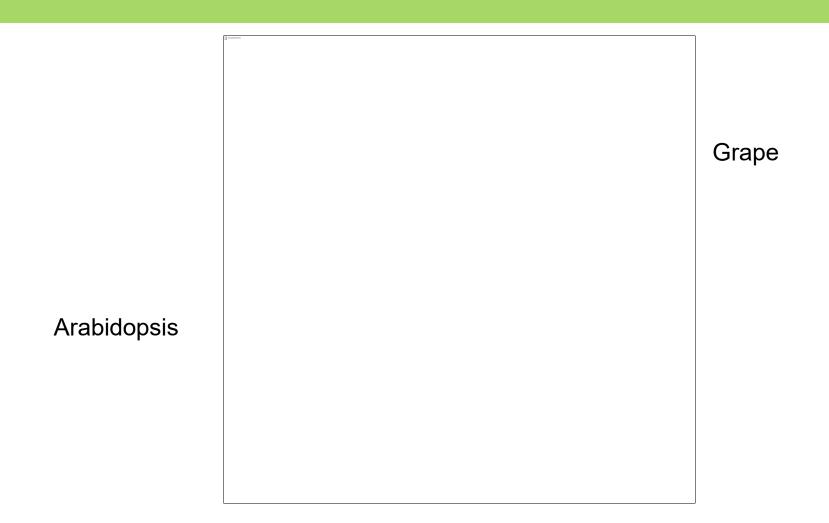


Synteny Analyses



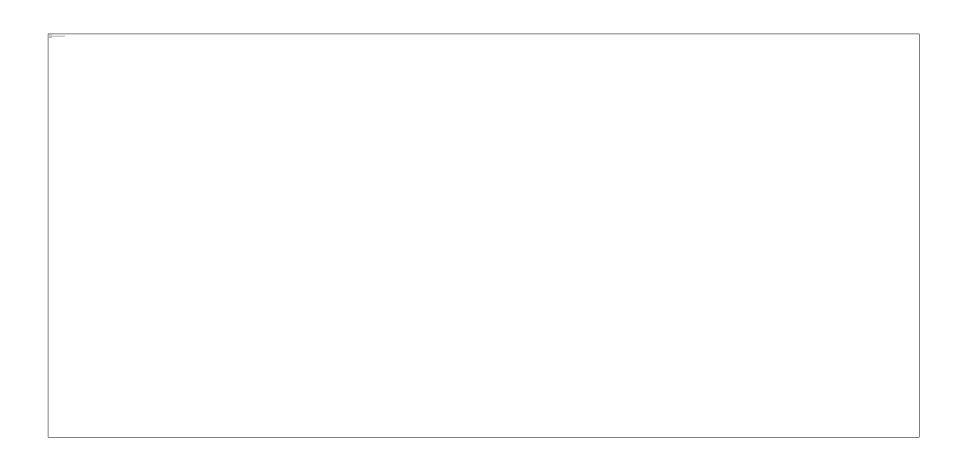
What is the history of paleopolyploidy?

Synteny Analyses



What is the history of paleopolyploidy?

Synteny Analyses



Whole Genome Sequences



Duplicate Gene Age Distributions

Carthamus tinctorius % of Duplicate Pairs

Ks (~ Time)

Inferring Paleopolyploidy

Histograms of gene age distributions for six species of Compositae and related families.

The y-axis is the number of gene duplications; the x-axis is synonymous divergence (Ks) of the duplication.

Peaks of gene duplication in panels A, B, and C correspond to whole-genome duplications.

American J of Botany, Volume: 103, Issue: 7, Pages: 1203-1211, First published: 16 June 2016, DOI: (10.3732/ajb.1600113)

Compositae Compositae

Calyceraceae

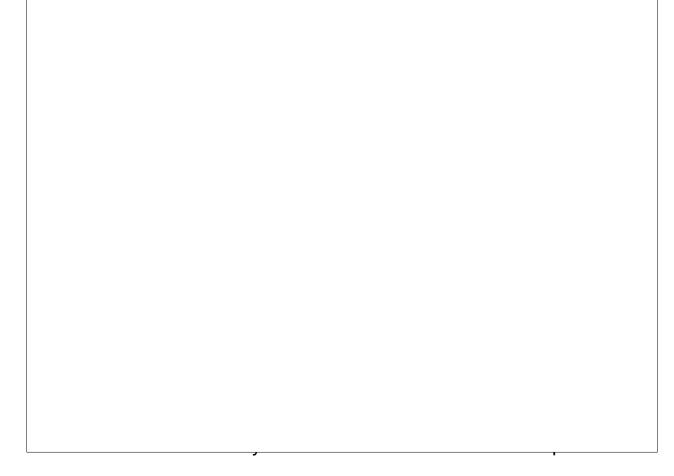
Goodeniaceae

Menyanthaceae

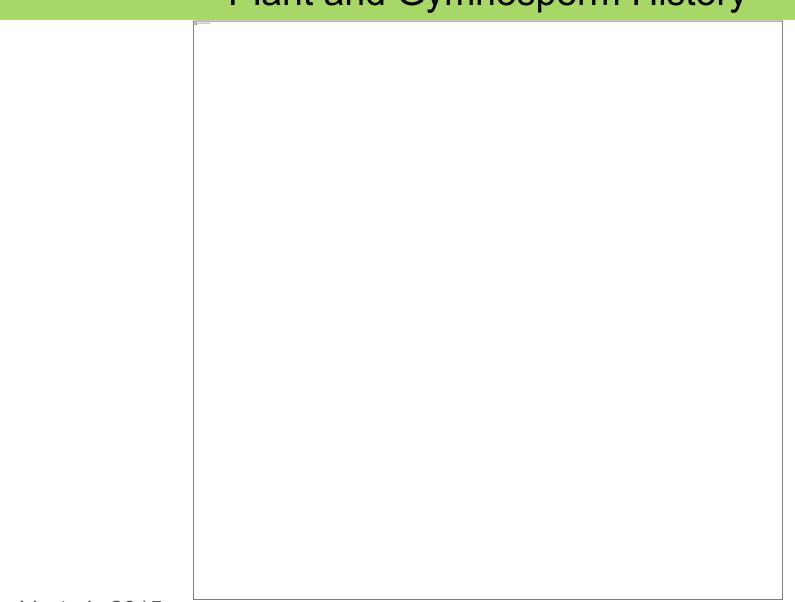
Campanulaceae

Inferring Paleopolyploidy

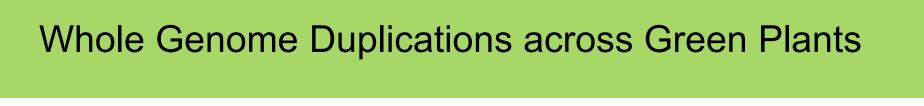
Species tree of Compositae and related families with percentage of gene duplications at each node.

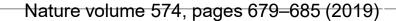


Whole Genome Duplications WGDs in Seed Plant and Gymnosperm History



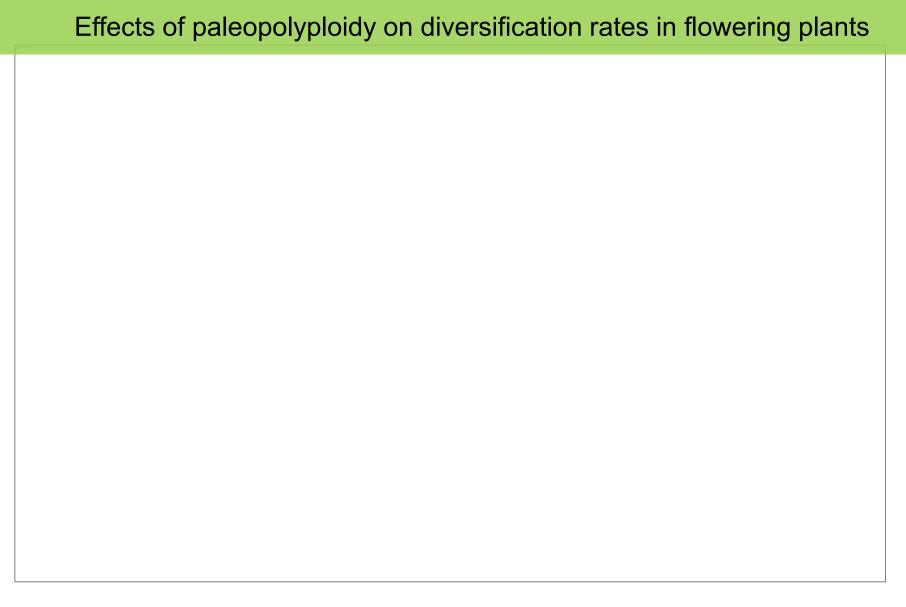
Li et al., 2015





Whole Genome Duplications across Green plants

- 244 Independent Paleopolyploidizations
- Most algal lineages show no evidence of WGDs
- All land plants except Selaginetta & some liverworts are ancient polyploids
- All seed plants are ancient polyploids
- All flowering plants characterized by at least two WGDs





Significant increases in diversification rates in flowering plants

•Half are associated with paleopolyploidy (*p* = 0.005)

Resolution

Polyploidy is most often an evolutionary dead end, but the expanded genomic potential of those polyploids that do persist drives longer term evolutionary success.

Unanswered questions

Do auto- and allopolyploids differ in their evolutionary success?

How long must a polyploid lineage persist before it transitions from a trajectory that favors extinction to one that favors diversification?

What evolutionary genetic changes/processes underlie this transition?