The ultrasmall green alga Ostreococcus unveils a unique regulation of the Calvin cycle

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Glyceraldehyde-3-phosphate dehydrogenase (GAPDH)

Calvin cvcle



- Iand plant: nuclear encoded
 - GapA: cyanobacterial (plastid) ancestor
 - GapC: proteobacterial (mitochondrial) origin
- interacts with CP12

Plays a role in light/dark regulation of the Calvin Cycle Contains characteristic carboxyterminal extension (CTE), to which it binds GAPDH

- transition from green algae to land plants
 - Origin of GapB (after GapA/B duplication)
 - land plant specific
 - also contains the characteristic carboxyterminal extension (CTE) as CP12
 - GapB is autonomously capable of forming high molecular GADPH complexes

Ostreococcus tauri is a unicellular green alga, belonging to the Prasinophyceae. With a size less than 1_m, comparable to that of a bacterium, it is the smallest eukaryotic organism described until now. The nuclear genome is 12.56 Mb, fragmented into 20 chromosomes making it the smallest of all known free-living eukaryotes. 8166 proteincoding genes were predicted, making it the most gene dense of all free-living eukaryotes sequenced to date.



Ultrastructure of Ostreococcus tauri

A. TEM section of Ostreococcus tauri ${\bf B}.$ Diagram illustrating the main features of the TEM section of the cell (G, Golgi body; Mt, Mitochondrion; N, nucleus; Np, nuclear pore; P, chloroplast; Sg, starch arain)

The GapA/B Gene Duplication Marks the Origin of Streptophyta

Petersen et al. MBE (23:1109-1118, (2006))

sequencing of GapA, GapB and CP12 genes of different plants

GapA: present in both Streptophyta and Chlorophyta and Mesostigma

CP12: universally present in green plants

°C

GapB:

only present in Streptophyta and Mesostigma



the GapB specific C-terminal domain allows an unambiguous identification of the respective gene

Phylogenetic analyses

the GapA/B gene duplication occurred in the common ancestor of Mesostiama. charophytes and land plants and coincides with the emergence of Streptophytes

Mesostigma viride belongs to the early Streptophyta

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GapA: present in both Ostreococcus strains

Situation in two Ostreococcus strains

CP12: absent

GapB: present

The GapB specific C-terminal domain is present



Both exhibit the GapB specific sequence pattern including two characteristic insertions, ruling out an independent CTE recruitment via gene fusion between GapA and CP12 duplicates



or This finding antedates the duplication



Systems

Biology

Conclusion: The prasinophyte Ostreococcus would be the first example where, due to the absence of CP12, the Calvin exclusively inactivated by the formation of GapAB complexes. cycle is

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