

INSIGHT

PLANT EVOLUTION

The inevitability of C₄ photosynthesis

Elements of C₄ photosynthesis—a complex adaptation that increases photosynthetic efficiency—may have evolved first to correct an intercellular nitrogen imbalance, and only later evolved a central role in carbon fixation.

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Related research article Mallmann J, Heckmann D, Bräutigam A, Lercher MJ, Weber APM, Westhoff P, Gowik U. 2014. The role of photorespiration during the evolution of C_4 photosynthesis in the genus *Flaveria. eLife* **3**:e02478. doi: 10.7554/ eLife.02478

Image C_4 plants (Fb and Ft) have higher levels of certain enzymes than plants that use other forms of photosynthesis

Notes that the evolution of complex innovations remains one of the most challenging problems in biology (Lynch, 2007; Wagner, 2014). Insights often stem from experimental lab studies that manipulate systems under 'directed evolution' (Weinreich et al., 2006; Blount et al., 2012; Finnigan et al., 2012). However, complex traits that have evolved many times over in independent lineages present a different—yet equally powerful—opportunity to infer the evolutionary trajectories of novel traits.

In flowering plants, C_4 photosynthesis is a wellstudied, complex adaptation that has independently evolved over 60 times (*Sage et al., 2011*). Many key, shared stages along the C_4 evolutionary trajectory have been identified by studying multiple C_4 -evolving plant groups (e.g., *Kennedy*) et al., 1980; Ku et al., 1983; Vogan et al., 2007; Williams et al., 2013). Now, in *eLife*, Udo Gowik and colleagues at Heinrich-Heine-Universität including Julia Mallmann and David Heckmann as joint first authors—present a compelling new hypothesis for how the final evolutionary steps were realized (Mallmann et al., 2014).

Although atmospheric carbon dioxide (CO₂) levels are currently rising, the last 30 million years witnessed great declines in CO₂, which has limited the efficiency of photosynthesis. Rubisco, the critical photosynthetic enzyme that catalyses the fixation of CO₂ into carbohydrate, also reacts with oxygen when CO₂ levels are low and temperatures are high. When this occurs, plants activate a process known as photorespiration, an energetically expensive set of reactions that—importantly for this story—release one molecule of CO₂.

 C_4 photosynthesis is a clever solution to the problem of low atmospheric CO₂. It is an internal plant carbon-concentrating mechanism that largely eliminates photorespiration: a 'fuelinjection' system for the photosynthetic engine. C_4 plants differ from plants with the more typical 'C₃' photosynthesis because they restrict Rubisco activity to an inner compartment, typically the bundle sheath, with atmospheric CO₂ being fixed into a 4-carbon acid in the outer mesophyll. This molecule then travels to the bundle sheath, where it is broken down again, bathing Rubisco in CO₂ and limiting the costly process of photorespiration.



Figure 1. An 'Evolvability Landscape' for C_4 photosynthesis. Many intermediate stages along the evolutionary trajectory from C_3 to C_4 are well known (**Sage et al., 2012**). These can be displayed as part of an adaptive fitness landscape, which links biological properties (horizontal axis) with the fitness they produce (right vertical axis; a greater height indicates a greater fitness). The adaptive fitness landscape of the C_4 trajectory was recently modelled as 'Mt. Fuji-like': a steep linear incline with each step along the trajectory bringing small, incremental increases in fitness (*Heckmann et al., 2013*), represented here by the grey dashed line. The gains in relative likelihood of evolving C_4 , or the 'evolutionary accessibility' of the pathway, may not be so linear (left vertical axis; black line). In spite of some limited flexibility in the order of trait acquisition (*Williams et al., 2013*), two intermediate stages are relatively fixed in position along the trajectory and also provide steep increases in C_4 evolvability. One early step, an elevated ratio of bundle sheath: mesophyll cross-sectional area (BS:M ratio) was recently identified as a key trait that preceded multiple parallel realizations of C_4 (*Christin et al., 2013*). Mallman et al. propose a mechanistic interaction between C_2 and C_4 photosynthesis, suggesting that evolution of the C_2 stage of the trajectory greatly increases the probability that full C_4 photosynthesis will quickly follow.

The evolution of the C_4 pathway requires many changes. These include the recruitment of multiple enzymes into new biochemical functions, massive shifts in the spatial distribution of proteins and organelles, and a set of anatomical modifications to cell size and structure. It is complex, and it is also highly effective: C_4 plants include many of our most important and productive crops (maize, sorghum, sugarcane, millet) and are responsible for around 25% of global terrestrial photosynthesis (**Still et al., 2003**).

A key intermediate step in the evolution of C_4 is the establishment of a rudimentary carbonconcentrating mechanism. Termed ' C_2 photosynthesis', this mechanism limits certain reactions of the photorespiratory cycle to the bundle sheath cells. A byproduct of these reactions is CO_2 , creating a slightly elevated CO_2 concentration and increasing Rubisco efficiency in these cells. Though much rarer than C_4 plants, C_2 plants have been discovered in a variety of C_4 -evolving lineages, and are thought to represent a common, if not requisite, intermediate step along the C_4 trajectory (*Sage et al., 2012*).

One implication of a restricted photorespiratory cycle is the development of a severe nitrogen imbalance between the mesophyll and the bundle sheath cells. This occurs because every molecule of CO_2 produced in the bundle sheath is accompanied by a molecule of ammonia. While this nitrogen imbalance has previously been recognised (**Monson and Rawsthorne, 2000**), it has never been closely studied, and certainly never considered as potentially important to the evolutionary assembly of the C₄ pathway.

To investigate this, Mallmann, Heckmann et al. combined a mechanistic model of C_2 physiological function with a metabolic model, which allowed them to predict the buildup of certain metabolites based on the rates of Rubisco and photorespiratory activity. They then modelled the various biochemical pathways that could potentially be induced to balance metabolic fluxes between the mesophyll and bundle sheath cells. This creative combination of models allowed them to evaluate the various metabolic pathways for re-balancing nitrogen in terms of which pathways resulted in the highest biomass yield (a proxy for fitness).

Remarkably, when low levels of C₄ enzyme activity are permitted in the model, key elements of the C4 cycle are favoured as the nitrogenbalancing pathway. What's more, this model predicts that with a C₄ cycle established, increasing the activity of the enzymes results in a linear increase in biomass yield. Allowing for low levels of C₄ enzyme activity is biologically reasonable, as these enzymes are routinely present in C_3 leaves. Mallmann, Heckmann et al. support their model predictions with experimental gene expression data from a set of C_3 , C_2 , C_4 , and other C_3 - C_4 intermediate types in the plant lineage Flaveria, which show elevated C_4 cycle activity even in intermediates that are not using the enzymes to capture carbon.

In other words, once a C_2 cycle is established, the evolution of a fully realized C_4 process is fairly trivial. Once C_4 enzymes are recruited to shuttle nitrogen back to the mesophyll, it is all but inevitable. This can explain in part why C_4 has evolved such a startling number of times, and why many of these origins are highly clustered across the tree of life. Many C_4 evolutionary clusters likely share an ancestor that had already acquired an elevated likelihood of evolving the pathway (**Figure 1**).

This may also explain why C_2 species are so rare relative to C_4 species— C_2 is likely to be a step along the trajectory with a relatively short evolutionary lifespan. At the same time, it raises the question of why a handful of C_2 species are persistent—the C_2 *Mollugo verticillata* group may be up to 15 million years old (*Christin et al., 2011*). A testable hypothesis would be that these C_2 plants have solved their nitrogen problem a different way, thereby limiting their own evolutionary accessibility to C_4 photosynthesis. If so, this highlights the key role of contingency in adaptation, and our growing power to understand and predict macroevolutionary processes.

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