

# Photosynthetic Light Reactions in C<sub>4</sub> Photosynthesis

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## Abstract

The most productive wild plants and crops use C<sub>4</sub> photosynthesis (Brown, 1999). C<sub>4</sub> photosynthesis requires the coordinated functions of two cell types in leaves, namely mesophyll (M) - and bundle-sheath (BS) - cells (Hatch, 1987). Atmospheric CO<sub>2</sub> is initially fixed by phosphoenolpyruvate carboxylase (PEPC) in M cells. The resulting products, C<sub>4</sub> acids are transported into BS cells where CO<sub>2</sub> is released by decarboxylation of C<sub>4</sub> acids and refixed by ribulose-1, 5-bisphosphate carboxylase/oxygenase (RuBisCO) functioning in the Calvin cycle (C<sub>3</sub> cycle). Since this process increase CO<sub>2</sub> levels at the site of RuBisCO in the BS cells, the oxygenase reaction of RuBisCO is largely reduced and RuBisCO can achieve maximal catalytic activity of CO<sub>2</sub> fixation. Therefore, C<sub>4</sub> plants have higher potential efficiencies in the use of light, water and nitrogen than C<sub>3</sub> plants (Long, 1999). C<sub>4</sub> plants have evolved from ancestral C<sub>3</sub> plants (Sage *et al.*, 2011). In C<sub>4</sub> plants, not only the CO<sub>2</sub> metabolism but a manner of light reactions which is a process to produce ATP and NADPH used for CO<sub>2</sub> metabolism has been changed. However, mechanism of light reactions and its physiological roles on C<sub>4</sub> photosynthesis are not yet fully understood. Here, we introduce latest findings and perspectives of physiological roles of light reactions in C<sub>4</sub> photosynthesis.

*Keywords: C<sub>4</sub> photosynthesis, cyclic electron flow, Flaveria, Zea mays*

## The structure of mesophyll and bundle-sheath chloroplasts: the regulation of grana formation

Light reactions take place on thylakoid membranes in chloroplasts. The thylakoid membranes are arranged in stacked (grana) and unstacked (stroma lamellae) in C<sub>3</sub> plants. Supercomplex of photosystem (PS) II and light-harvesting complex of photosystem II (LHC-II) are localized in grana thylakoids while most of PSI and ATP synthase are localized in stroma lamellae (Albertsson, 2001). The grana stacks are more developed in low light intensity than in high light intensity, the changes are assumed to optimize the photosynthetic efficiency in C<sub>3</sub> plants (Ballantine and Forde, 1970). In NADP-malic enzyme (ME) type C<sub>4</sub> plants such as *Zea mays*, *Sorghum bicolor* and *Flaveria trinervia*, grana thylakoids are observed in the M chloroplasts like chloroplasts of C<sub>3</sub> plants, but BS cells possess chloroplasts where grana thylakoids are largely decreased, as shown in Fig. 1 (Laetsch *et al.*, 1965, Laetsch and Price, 1969, Hofer *et al.*, 1992). Interestingly, grana formation was observed in BS chloroplasts of maize leaves at very low light intensity but was completely abolished at normal light intensity (Brangeon, 1973). Decrease in grana stacks of BS chloroplasts is likely induced by high light signal and those systems would have been recruited from C<sub>3</sub> plants during C<sub>4</sub> evolution.

LHC-II is suggested to be involved in grana formation by binding thylakoid membranes in C<sub>3</sub> plants (Allen and Forsberg, 2001, Standfuss *et al.*, 2005). LHC-II is mainly present as trimer on grana thylakoids. It is reported that in *Flaveria trinervia* the amounts of LHC-II polypeptides in BS chloroplasts is only slightly reduced compared with that in M chloroplasts (Hofer *et al.*, 1992) and a significant amount of LHC-II polypeptides are also detected in fractionated stroma lamellae (Munekage *et al.*, 2010). Moreover, in maize, LHC-II trimer is identified in BS thylakoid membranes

using Blue-Native PAGE, although BS chloroplasts of Maize are almost grana-free (Romanowska *et al.*, 2008). These results suggest that amounts of LHC-II polypeptide and the presence of LHC-II trimer are not sufficient for grana formation. Perhaps the additional mechanism is involved in the formation of thylakoid membranes with reduced grana in BS chloroplasts.

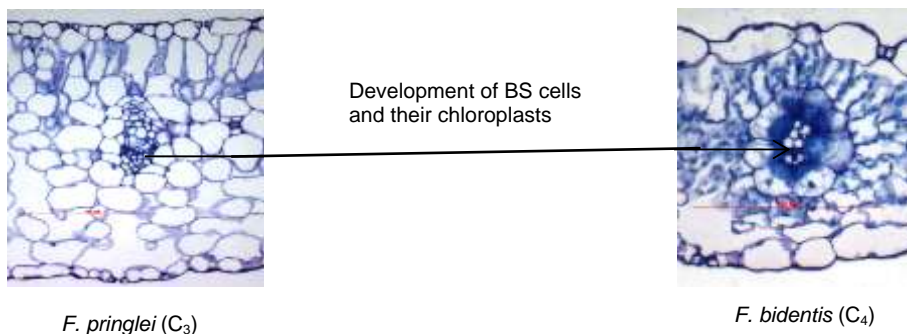


Figure 1. Light micrograph's of leaf transverse section of *Flaveria species*.

### Regulation of linear electron flow and cyclic electron flow around PSI in C<sub>4</sub> photosynthesis

Linear electron flow operates to produce the both ATP and NADPH. In linear electron flow, the electrons excised from water molecules at PSII are transferred to PSI through the cytochrome *b<sub>6</sub>f* complex, and finally NADPH is produced. Protons are transported across the thylakoid membranes from stroma side to lumen side during the process of electron transfer. This proton gradient is used for ATP synthesis (Fig. 2). On the other hand, cyclic electron flow around PSI (CEF1) contributes to only ATP production. In CEF1, electrons are returned from acceptor side of PSI to plastoquinone which mediate electron transport from PSII to cytochrome *b<sub>6</sub>f* complex (Shikanai, 2007). This electron transport contributes to form proton gradient without NADPH production. Two CEF1 pathways have been identified in C<sub>3</sub> plants.

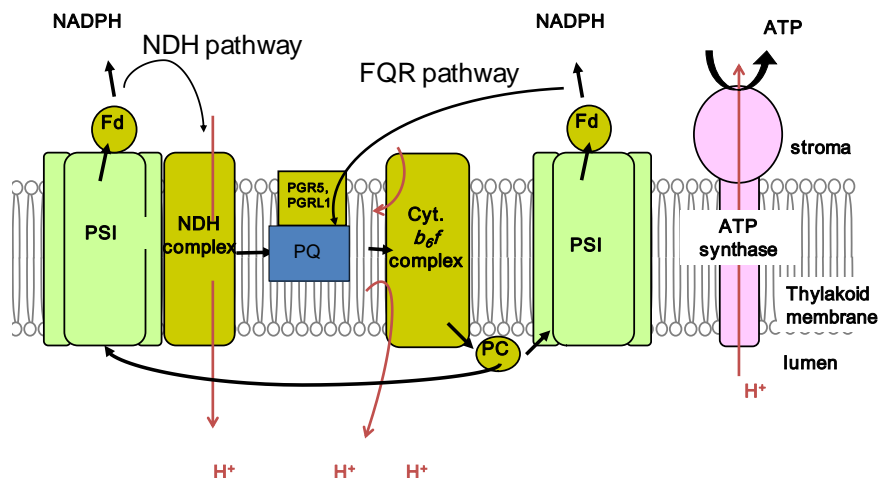


Figure 2. Schematic model of PS1 cyclic electron transport.

The first pathway is called as ferredoxin-plastoquinone reductase (FQR) pathway involving PROTON GRADIENT REGULATION 5 (PGR5) and PGRL1 (Munekage *et al.*, 2002, Dalcorso *et al.*, 2008). The other pathway is NAD(P)H dehydrogenase (NDH) pathway involving a plastidial NDH

complex composed of the eleven plastid-coded subunits (NDH-A to -K) and the over 15 nuclei-encoded subunits (Peng *et al.*, 2011). In C<sub>3</sub> plants, the FQR pathway contributes for maintaining the production ratio of ATP/NADPH and for photoprotection, whereas the NDH pathway functions in stress resistance in C<sub>3</sub> plants (Munekage *et al.*, 2004, Wang *et al.*, 2006, Munekage *et al.*, 2008). Localization of the NDH complex and PGR5 on stroma lamellae and distribution of PSII and PSI on thylakoid membrane suggest that CEF1 takes place in stroma lamellae (Rumeau *et al.*, 2005, Munekage *et al.*, 2010).

In C<sub>4</sub> plants, increased CEF1 activity and decreased linear electron transport is induced. In NADP-ME type C<sub>4</sub> plants, BS chloroplasts of mature leaves show the inhibition of PSII activity (Woo *et al.*, 1970). The composition of PSII complex in BS chloroplasts is investigated by many researchers using *Zea mays*, *Sorghum bicolor* and *Flaveria trinervia*. Several studies are suggested that the subunits of PSII complex, especially those of oxygen evolving complex are largely reduced in BS chloroplasts compared with M chloroplasts (Sheen *et al.*, 1987, Oswald *et al.*, 1990, Hofer *et al.*, 1992). BS chloroplasts produce only limited amounts of ATP and NADPH from linear electron flow activity, because in this chloroplasts, PSII activity is significantly inhibited (Schuster *et al.*, 1985). Since NADPH is supplied by decarboxylation of C<sub>4</sub> acids by NADP-ME, only ATP is required to be produced in BS chloroplasts of NADP-ME type. The ATPs are considered to be produced by CEF1 activity. Significant activity of CEF1 compared with C<sub>3</sub> plants has been reported in *Zea mays*, *Sorghum bicolor* (Herbert *et al.*, 1990, Asada *et al.*, 1993). The higher amounts of NDH complex and PGR5 in BS cell than M cell also supports activation of CEF1 in BS cell of NADP-ME type C<sub>4</sub> plants (Kubicki *et al.*, 1996, Munekage *et al.*, 2010). On the other hand, ATP demands are increased in M chloroplasts in NAD-ME type C<sub>4</sub> metabolism. In *Salsola laricina* and *Halocharis gossypina* carrying NAD-ME type C<sub>4</sub> photosynthesis, lower grana index and higher ratio of PSI/PSII in M chloroplasts than BS chloroplasts are reported (Voznesenskaya *et al.*, 1999). Moreover, subunits of NDH complex are enriched in M chloroplasts of NAD-ME type C<sub>4</sub> plants (Takabayashi *et al.*, 2005). Activated CEF1 likely contributes to the ATP production in C<sub>4</sub> photosynthesis. However, for final conclusion, it is need to investigate phenotype of transgenic plants in which genes involved in CEF1 are knocked down.

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