The discovery of plastid-to-nucleus retrograde signaling—a personal perspective

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Abstract DNA and machinery for gene expression have been discovered in chloroplasts during the 1960s. It was soon evident that the chloroplast genome is relatively small, that most genes for chloroplast-localized proteins reside in the nucleus and that chloroplast membranes, ribosomes, and protein complexes are composed of proteins encoded in both the chloroplast and the nuclear genome. This situation has made the existence of mechanisms highly probable that coordinate the gene expression in plastids and nucleus. In the 1970s, the first evidence for plastid signals controlling nuclear gene expression was provided by studies on plastid ribosome deficient mutants with reduced amounts and/or activities of nuclear-encoded chloroplast proteins including the small subunit of Rubisco, ferredoxin NADP+ reductase, and enzymes of the Calvin cycle. This review describes first models of plastid-to-nucleus signaling and their discovery. Today, many plastid signals are known. They do not only balance gene expression in chloroplasts and nucleus during developmental processes but are also generated in response to environmental changes sensed by the organelles.

Keywords Chloroplast signal · Plastid signal · Retrograde signaling · Chloroplast development · Regulation of gene expression

Introduction

It was firmly established already in the early 1970s that plastids of algae like Euglena, Chlamydomonas, and Acetabularia and of higher plants contain their own genomes, RNA polymerase activity to transcribe their genes, as well as ribosomes, tRNAs, and aminoacyl-tRNA synthetases for protein synthesis (Gibor and Granick 1964; Kirk 1971; Tewari 1971; Gillham 1974). Chloroplasts were found to contain multiple copies of their genomes in several nucleoids, the number of which increases with plastid size (Gunning 1965; Herrmann and Kowallik 1970; Kung 1977). Good evidence existed for chloroplast genes to code for chloroplast ribosomal RNA, tRNAs, and the large subunit of Rubisco (ribulose-1,5-bisphosphate carboxylase/oxygenase) (Kirk 1971; Chan and Wildman 1972; Blair and Ellis 1973; Börner 1973; Bogorad 1975). A chlorophyll-binding protein of photosystem I, chloroplast ribosomal proteins, and components of cytochrome complexes were also proposed to be encoded in chloroplast DNA (Armstrong et al. 1971; Herrmann 1971; Mets and Bogorad 1971). It has soon been realized, however, that the size of chloroplast genomes deduced from electron microscopic pictures and renaturation kinetics is much too small to code for all components of the organelar genetic machinery, the photosynthetic apparatus, and the other plastid-localized metabolic pathways (Kolodner and Tewari 1975; Kung 1977). First studies on chromoplast DNA revealed identical size and biochemical properties with chloroplast DNA (Herrmann 1972; Falk et al. 1974). The existence, in higher plants, of several types of plastids like chloroplasts, amyloplasts, chromoplasts, and gerontoplasts (a term coined by Peter Sitte in Sitte 1977), or proplastids raised the question as to the role of plastid and nuclear genomes in cell differentiation, in particular, in the formation of the different plastid types (Sitte 1977). Is the...
plastid type signaled to the nucleus to control plastid-type specific gene activities? Several lines of evidence had already earlier indicated that a large number of genes required for chloroplast biogenesis and function reside in the nuclear genome (Kirk and Tilney-Bassett 1967). Data collected during the early years of the molecular biology of plastids indicated that the proteins of the chloroplast envelope, of photosystems I and II, of the chloroplast ribosomes, and even the subunits of Rubisco are encoded by both chloroplast and nuclear genes (Kirk 1971; Börner 1973; Bogorad 1975). This observation strongly suggested the existence of one or more mechanisms, which coordinate the activity of genes in the genomes of chloroplasts and nucleus. Is the developmental and/or metabolic status of the plastid/chloroplast signaled to the nucleus?—a question that is also connected to the much discussed problem of chloroplast autonomy (Tewari 1971).

Today, we know that the developmental and metabolic states of plastids, in particular of chloroplasts, but also of mitochondria, are signaled to the nucleus via different signaling molecules and signal chains. The activity of nuclear genes is modified in response to these organellar signals to adapt gene expression and other metabolic functions in the organelle, but also processes outside the plastids, to the requirements of development and of changes in the environment (Chan et al. 2016; Kleine and Leister 2016; de Souza et al. 2017). The processes of organello-to-nuclear signaling are often referred to as ‘retrograde signaling’ and the regulation of organellar gene expression by nucleus-encoded proteins as ‘anterograde signaling’ although the latter is (with the exception of a few cases) in a strict sense no ‘signaling’ and should be better termed ‘anterograde control’ or ‘anterograde regulation.’ Organellar signals control transcription and post-transcriptional processes in the nucleo-cytoplasmic compartment and vice versa nuclear-encoded proteins execute and control gene expression at all levels in the organelles.

Coordination of the expression of chloroplast and nuclear genes—first hypotheses

First ideas about the control of the expression of nuclear and plastid genes coding for plastid proteins were developed specifically for the subunits of Rubisco (often called “fraction I protein” at that time; Kawashima and Wildman 1970). John Ellis from the University of Warwick, Coventry, contributed together with his coworkers essential insights into the functional role of chloroplast ribosomes by identifying products of protein synthesis in isolated chloroplasts and etioplasts and by applying specific inhibitors of cytoplasmic or chloroplast protein synthesis. He observed that 2-(4-methyl-2,6-dinitroanilino)-N-methylpropionamide, at concentrations that inhibit rather specifically protein synthesis on cytoplasmic ribosomes, rapidly inhibits the synthesis of both the large and small subunits of fraction I protein on chloroplast and cytoplasmic ribosomes, respectively, in greening detached pea shoots. Based on this finding, Ellis proposed a model “for the co-operation of nuclear and chloroplast genomes in the synthesis of fraction I protein by noting that the small subunit is a positive factor required for the initiation of either transcription of the messenger RNA for the large subunit or for its translation. This model predicts that inhibition of synthesis of small subunit on cytoplasmic ribosomes will lead to a rapid inhibition of synthesis of large subunit on chloroplast ribosomes” (Ellis 1975). Givan and Criddle (1972) proposed also a not further elaborated regulatory mechanism as possible explanation for their observation that inhibition of protein synthesis in the cytoplasm affected not only the formation of small but also of large subunits of Rubisco in Chlamydomonas.

Somewhat later, Ellis (1977) substantially extended his model by formulating two principles for the control of chloroplast protein synthesis: “The multisubunit completion principle”... states that the function of organellar ribosomes is to synthesize some subunits of multisubunit proteins, the other subunits being products of cytoplasmic ribosomes... There is no reported case of a complete protein being made by organellar ribosomes, and the multisubunit completion principle states that this does not occur. “The cytoplasmic control principle”... states that cytoplasmic products control organellar protein synthesis, but that the converse does not occur... Combining these two principles leads to the suggestion that the cytoplasmic products, which control the synthesis of subunits in the organelle, are the same subunits which combine with them to form the complete protein. This is not a necessary conclusion, but it describes the most economical arrangement” (Ellis 1977).

A study on the incorporation of (14C) leucin into total plastid protein of Euglena gracilis cells growing at 27 or 35 °C without or with chloramphenicol or cycloheximide in the medium led to the postulate that proteins synthesized on plastid ribosomes have an influence on the translation in the cytoplasm and vice versa (Brandt 1976). This investigation had several weaknesses and has not been discussed by other authors in the following years.

Expression of nuclear genes for plastid proteins is reduced in plastid ribosome deficient mutants—evidence for plastid-to-nucleus signaling

Studies on the Pelargonium cultivar “Mrs. Parker” provided a first hint of a role of plastids in the coordination of gene activities in the nucleus and chloroplasts, i.e., that in a certain sense the converse of the “cytoplasmic control principle” might happen. This cultivar is a chimera characterized by green leaves with white margins. It eventually develops entirely green and white leaves containing normal wild-type chloroplasts and undifferentiated mutant plastids, respectively.
The cultivar “Mrs. Parker” was the subject of my doctoral studies in Rudolf Hagemann’s lab at Martin Luther University in Halle (Saale), East Germany. I could not detect plastid rRNAs in the white leaves of “Mrs. Parker” and somewhat later not in white leaves of several other Pelargonium cultivars, an observation that was supported by the lack of ribosomes in electron microscopic pictures of the mutant plastids. The plastid ribosome deficiency should lead to the inability of the mutant plastids to express their protein-encoding genes and made this mutant an ideal object of investigations into the functional role of plastid ribosomes (Börner et al. 1972, 1973). The protein with the best evidence for being synthesized on chloroplast ribosomes was the large subunit of Rubisco at that time (Criddle et al. 1970; Chan and Wildman 1972; Blair and Ellis 1973). To gain further support for the supposed lack of ribosomes in the Pelargonium mutant plastids, I isolated therefore soluble proteins from green and white leaves and separated them on native polyacrylamide gels. This method allows for the easy detection of ‘fraction I protein’ (Rubisco). As anticipated, a protein band at the position of fraction I protein consisting of the large and small Rubisco subunits was found in preparations of soluble proteins from green but not from white leaves. The missing Rubisco band was obviously due to the incapacity of ribosome-deficient plastids to synthesize the large subunits. However, since the small subunits of Rubisco are synthesized on cytoplasmic ribosomes, which are functioning in the white leaves, the obvious lack of a corresponding band in the gels was not expected (Börner et al. 1974).

The lack of the small Rubisco subunits in the mutant plastids of “Mrs. Parker” and of other cultivars could have different reasons. One would be that proteases degrade the small subunits because they are not protected by assembling with the large subunits to form an active enzyme complex. Another reason could be a specific mechanism that coordinates the synthesis of the two Rubisco subunits as suggested by Givan and Criddle (1972) for Chlamydomonas and later by Ellis (1975) for higher plants. I hypothesized, however, that there might exist a more general mechanism that coordinates the expression of chloroplast and nuclear genes, i.e., a mechanism that would, in white “Mrs. Parker” leaves, repress or not activate the expression of most or all nuclear genes coding for those proteins that are specifically needed in photosynthetically active chloroplasts.

To rule out the possibility that just degradation of the small subunits is the reason for their absence in plastid ribosome deficient mutant leaves, I decided to determine the activity of such chloroplast-localized enzymes which were known to be synthesized on cytoplasmic ribosomes and would, in contrast to the small Rubisco subunits, not form complexes with other proteins encoded in chloroplast genes and synthesized on chloroplast ribosomes. Unfortunately, Pelargonium leaves (green leaves more, white ones somewhat less) turned out to contain substances, most likely polyphenols, which precipitate and inactivate proteins during the isolation procedure. By modifying the conditions, I was able to keep proteins soluble but not to preserve their activities (Börner et al. 1974). Therefore, I studied albino mutants of other species in 1972 and 1973 and luckily found two barley (Hordeum vulgare L.) mutants, Saskatoon and albostrians, to lack plastid ribosomes and the large and small subunits of Rubisco (Hagemann et al. 1973; Börner et al. 1976). Using sensitive immunoelectrophoresis, we detected traces of Rubisco in young white leaves of “Mrs. Parker” but not in light-grown or etiolated albostrians leaves suggesting residual translational activity in plastids of the Pelargonium mutant and the complete absence of ribosomes in plastids of white leaves of the barley mutant (Börner et al. 1976; Hagemann and Börner 1978; Reichenbächer et al. 1978). Activity tests of Rubisco supported also its absence from white leaves of albostrians and Saskatoon. For further studies, I selected the ferredoxin: NADP$^+$ reductase and the Calvin cycle enzyme phosphoribulokinase. Like the small subunit of Rubisco, these enzymes were proposed to be synthesized on cytoplasmic ribosomes (Armstrong et al. 1971; Ellis and Hartley 1971; Vaisberg et al. 1976). In contrast to the small Rubisco subunit, however, ferredoxin: NADP reductase (at least its soluble form) and phosphoribulokinase were not known to form part of complexes with other proteins. A poorly reacting antibody against ferredoxin: NADP$^+$ reductase from Antirrhinum majus did not detect ferredoxin: NADP$^+$ reductase in white leaves. Activity tests revealed later clearly the existence of this enzyme in white leaves, though with reduced activity; white leaves contain about 60% of the soluble form and less than 10% of the membrane-bound form of the ferredoxin: NADP$^+$ reductase activity compared with green leaves (part of these results were later published in Börner 1981). The activity of phosphoribulokinase was measured together with Klaus-Peter Rindt, who at the time was working on enzymes involved in bacterial photosynthesis in Erich Ohmann’s group at the Institute of General Botany of Martin Luther University. The results of this preliminary study suggested a marked reduction of this enzyme in white primary leaves of albostrians barley, a reduction being much more drastic than in the case of the ferredoxin reductase. Since all these observations supported the hypothesis of a controlling influence of the plastid/chloroplast on the expression of nuclear genes coding for chloroplast proteins, I was already in the mid 1970s convinced that such a regulatory mechanism really operates in leaves.

Quite surprisingly, I got the permission to travel to the UK in March and April 1977 to visit several labs involved in research on chloroplasts and plant mitochondria. This was rather exceptional for a young scientist in East Germany and only possible thanks to the strong support by my PhD supervisor and head of the Institute of Genetics at the University of Halle, Rudolf Hagemann (and certainly thanks to the fact that
my wife and our two daughters would stay at home as a guarantee for my return to East Germany. Most of the time I spent in the lab of William Bradbeer at the King’s College, University of London. Bill was an expert of the biochemistry and physiology of chloroplast development including the Calvin cycle. I brought seeds of Saskatoon and albostrians with me to London, and Bill selected two Calvin cycle enzymes, phosphoribulokinase and glyceraldehyde 3-phosphate dehydrogenase (NADP⁺), to be studied in fractions of soluble proteins from green and white leaves of the barley mutants. In addition, the activities of the cytoplasmically localized isoenzyme, glyceraldehyde 3-phosphate dehydrogenase (NAD⁺), and of phosphoglycerate kinase were determined. The measured activity of phosphoglycerate kinase originated from two isoenzymes, one located in the plastids, the other in the cytoplasm. Already the first series of experiments clearly demonstrated extremely low activities of the plastid-localized enzymes phosphoribulokinase and glyceraldehyde 3-phosphate dehydrogenase (NADP⁺) in white leaves of albostrians while the cytoplasmic enzyme glyceraldehyde 3-phosphate dehydrogenase (NAD⁺) and the jointly measured activities of the plastid and cytoplasmic forms of phosphoglycerate kinase were much less affected. Similar results were obtained with Saskatoon and irrespective of whether etiolated or illuminated seedlings were investigated. These experiments were continued by Yvonne Atkinson in Bill’s lab.

**Plastid RNA as derepressor of nuclear genes?**

Shortly, after my visit to the UK, I presented a poster during the International Conference on the Regulation of Developmental Processes in Plants, which was held in July 1977 in Halle. The poster reported results of studies performed in Halle on the large and small subunits of Rubisco, phosphoribulokinase, and ferredoxin: NADP⁺ reductase together with the hypothesis of regulatory effects of the plastid/chloroplast on the expression of nuclear-encoded plastid proteins (Börner 1977). Based on genetic analyses, the barley mutant alleles Saskatoon and albostrians were originally described as examples of so-called plastome mutator genes, i.e., nuclear alleles that cause mutations in the plastid DNA (Arnason and Walker 1949; Hagemann and Scholz 1962). In contrast to several other plastome mutator genes, which generate various plastome mutations with different phenotypes (reviewed in Börner and Sears 1986), the two barley alleles were thought to induce in each generation the same mutant phenotype characterized by white leaves with undifferentiated, ribosome-deficient plastids (Börner et al. 1976). It is difficult to imagine how a nuclear gene product might induce plastome mutations leading again and again to an identical phenotype. Yet the mutations could theoretically be very drastic alterations of the genome like large deletions similar to what is known from albino seedlings generated by microspore culture (Day and Ellis 1984; Harada et al. 1991), massive alterations in the DNA sequence, or the complete loss of chloroplast DNA. Yet earlier investigations had shown that DNA is present in ribosome-free albostrians plastids (Hagemann and Börner 1978) and later we could exclude large deletions, insertions or rearrangements (Hess et al. 1993). Alternatively, the albostrians and Saskatoon gene products might not induce plastome mutations but be proteins that are needed for the biogenesis and function of chloroplast ribosomes. Once lacking, ribosomes cannot reappear since they depend on their own protein synthesizing capacity to produce the proteins they are composed of, i.e., the lack of chloroplast ribosomes should be stably inherited like a genuine DNA mutation (Börner and Sears 1986; Zubko and Day 1998). We know today that the albostrians allele does indeed not act as plastome mutator, but prevents the formation of plastid ribosomes early in plant development (M. Li, N. Stein, T. Börner, unpubl. data) and the same is probably true for Saskatoon (see also Han et al. 1992 for a similar mutant in maize). In 1977, however, I considered both a mutation in plastid DNA and the lack of ribosomes as equally likely primary cause of the albostrians phenotype. Therefore, my poster presented a hypothetical model in which an altered plastid gene product or its absence affects the expression of nuclear genes coding for chloroplast proteins.

Curious though, Jürgen Feierabend, then at the Ruhr University Bochum, showed during the same conference contrasting results on a poster directly adjacent to my presentation (Feierabend 1977a). Feierabend and coworkers published shortly before the conference that growth at elevated temperature (32 °C) prevents the formation of plastid ribosomes in leaves of rye (Secale cereale L.) (Feierabend and Schrader-Reichhardt 1976; Feierabend and Mikus 1977). The plastid ribosomal deficient leaves are chlorotic; hence, these plants represent phenocopies of the barley mutant albostrians and Saskatoon. Yet, in spite of the chloroplast ribosome deficiency, the chlorotic rye leaves do not exhibit the strikingly low amounts and activities of nuclear-encoded plastid proteins that we have observed in the barley mutants. Chlorotic rye leaves grown at elevated temperature and under illumination contain the plastid ribosomes in the absence of the large subunit. Related to the activities in green leaves grown at permissive temperature (22 °C), the plastid ribosomal deficient leaves exhibit nearly normal activities of 97% (glyceraldehyde 3-phosphate dehydrogenase (NADP⁺)) and 83% (phosphoribulokinase). In addition, other plastid enzymes are present with virtual normal activities (Feierabend 1977a; Feierabend and Schrader-Reichhardt 1976).

In the light of the results reported by Feierabend und coworkers on plastid ribosome deficient rye, it seemed unlikely that the ribosome deficiency and impaired protein synthesis in the mutant barley plastids (or any component/process that depends upon plastid protein synthesis) would be...
responsible for the extremely low amounts and activities of nuclear-encoded plastid proteins in albostrians and Saskatoon. We speculated, therefore, that mutation of the plastid DNA in albostrians and Sakatoon barley might affect nuclear gene expression independently of plastid protein synthesis via altered plastid transcript(s). According to our hypothesis, one or more “derepressors” would be transcribed from plastid DNA (implying that the “derepressors” are plastid RNAs) starting with the beginning of chloroplast development. The “derepressor” would derepress/activate those nuclear genes which code for proteins that are specifically needed in photosynthetically active plastids and would be present in wild-type leaves, also when grown at elevated temperature, but not in white leaves of albostrians barley. Bill Bradbeer presented this hypothesis together with the results of our investigations into enzyme activities in green and white leaves of albostrians and Saskatoon during the International Symposium on Chloroplast Development organized by George Akoyunoglou and Joan Argyroudi-Akoyunoglou and held in July 1978 on the island of Spetsai (Bradbeer and Börner 1978). The idea of RNA transcribed from plastid genes but functioning outside the plastids got support in the same year by a publication in Nature reporting that several tRNA species have their genes within the chloroplast DNA of E. gracilis but are involved in protein synthesis on cytoplasmic ribosomes (McCrea and Hershberger 1978). We included therefore the possibility that plastid-derived RNA controls protein synthesis on cytoplasmic ribosomes in the publication of our ideas about regulatory plastid RNA involved in the transcription of nuclear genes (Bradbeer et al. 1979). Not necessarily the existence of a regulatory plastid RNA, but the idea of an involvement of plastid transcription in signaling to the nucleus got later support by experiments with tagetitoxin and rifampicin, inhibitors of the plastid-encoded RNA polymerase (Lukens et al. 1987; Rapp and Mullet 1991; Woodson et al. 2013), and with nalidixic acid, a prokaryotic DNA gyrase inhibitor that affects plastid DNA replication and transcription (Gray et al. 1995), as well as by the analysis of Arabidopsis mutants defective in plastid sigma factors SIG2 and SIG6 (Woodson et al. 2013).

The contrasting results obtained with ribosome deficient plastids of heat-treated rye seedlings and of barley mutants, respectively, triggered, together with the “regulatory-RNA-hypothesis”, investigations into transcription of plastid DNA in this material. Although to different extent, either type of ribosome-deficient plastids synthesizes RNA (Bünger and Feierabend 1980; Siemenroth et al. 1981), i.e., a general lack of transcription of plastid genes in one type was not the reason for the observed differences. However, these studies revealed the existence of a nuclear-encoded plastid RNA polymerase (later dubbed NEP by Hajdukiewicz et al. 1997) and plastid rRNA was the first identified product of NEP (Siemenroth et al. 1981).

I moved to the Humboldt University at Berlin in 1982 and established there a lab dealing with research on the molecular genetics mainly of plants and cyanobacteria. This was a difficult and tedious task under the East German conditions. Therefore, we could resume research on albostrians barley only at the end of the 1980s and then more extensively after the reunification of the two Germanys. The problem of plastid-to-nucleus signaling became the topic of Wolfgang Hess’ PhD and Habilitation theses.

Intermediates of chlorophyll biosynthesis were among the factors first suggested to be involved in plastid-to-nucleus signaling (Johanningmeier and Howell 1984; Kropat et al. 1997). The biosynthesis of tetrapyrroles including chlorophyll, heme, siroheme, and phytochromobilin starts in plants with the ligation of glutamate to tRNA Glu (Brzezowski et al. 2015). Since tRNA Glu was known to be encoded by a chloroplast gene (Hollingsworth and Hallick 1982; Kuntz et al. 1984), we regarded it as a candidate for a plastid RNA involved in plastid-to-nucleus signaling. This RNA would not need to leave the chloroplast as originally thought (Bradbeer and Börner 1978; Bradbeer et al. 1979) but act inside the plastids via its role in tetrapyrrole synthesis. In collaboration with Gaby Walter and Paul Hoffmann (and later Bernhard Grimm) from our University and with Wolfhart Rädiger from the Ludwig Maximilian University at Munich, we studied the accumulation of tRNA Glu and the activity of several enzymes of the tetrapyrrole pathway in leaves of albostrians barley and in leaves of rye grown at elevated temperature. A remarkable difference between the two types of plastid ribosome deficient material was the extremely low accumulation (not detected by RNA blot hybridization, only by PCR) of tRNA Glu in white albostrians leaves while the chlorotic rye leaves contained substantial amounts of this RNA (Hess et al. 1992a, b; Walter et al. 1995). This difference is certainly caused by the lack of ribosomes in albostrians plastids, as opposed to the presence of a few ribosomes in the plastids of heat-treated rye leaves (Feierabend and Schrader-Reichhardt 1976; Feierabend 1977b). Due to the lack of ribosomes, the plastid-encoded plastid RNA polymerase (PEP) cannot be synthesized and only NEP transcribes the plastid genes in white albostrians leaves (Hess et al. 1993; Zhelyazkova et al. 2012). In contrast, NEP should be active together with the plastid-encoded plastid RNA polymerase (Pep) in the chlorotic rye leaves as suggested by the similar amounts of uridine incorporation into RNA of the plastids at 32 °C and chloroplasts at 22 °C (Bünger and Feierabend 1980). The plastid trnE gene coding for tRNA Glu is transcribed mainly by PEP and only very weakly by NEP (Hanaoka et al. 2003; Zhelyazkova et al. 2012). Thus, the lacking PEP activity will result in the observed low levels of tRNA Glu in white albostrians leaves, which in turn might be one reason for the much lower content of chlorophyll and carotenoids in albostrians plastids vs. plastids of heat-treated leaves (Börner...
and Meister 1980; Yaronskaya et al. 2003; Feierabend 1977b). Even though protein synthesis does not compete with tetrapyrole biosynthesis for tRNA^{Glu} in the ribosome-free albostrians plastids, the very low quantity of tRNA^{Glu} will limit the biosynthesis of tetrapyroles (Yaronskaya et al. 2003). The common precursors of all tetrapyroles are channeled in the direction of heme synthesis in albostrians plastids while the formation of chlorophylls is repressed. In addition to other effects, we observed reduced amounts of Mg-chelatase contrasted by a drastically enhanced accumulation of Fe-chelatase (Yaronskaya et al. 2003).

There is now experimental evidence for a positive effect of increased flux through the heme branch of the tetrapyrole pathway on the transcription of photosynthesis-associated nuclear genes (PhANGs) suggesting that heme might leave the chloroplast and act as a signal molecule (Woodson et al. 2011). Heme might not act as a positive signal in ribosome-free albostrians plastids, since the content of non-covalently bound heme is diminished to only 55 and 38% in etiolated and light/dark-grown white compared with green leaves in spite of the enhanced levels of Fe-chelatase. We concluded that altered levels of intermediates and products of the tetrapyrole pathway—which results at least in part from the impaired transcription of the trnE gene—might be involved in plastid-to-nucleus signaling (Yaronskaya et al. 2003). Recent studies on sigma factor mutants in Arabidopsis confirm a role of trnE transcription in plastid-to-nucleus signaling and PhANG expression (Woodson et al. 2013). Sigma factor SIG2 supports transcription of trnE by PEP (Hanaoka et al. 2003). Woodson et al. (2013) observed that in an Arabidopsis sig2 mutant the levels of tRNA^{Glu}, non-covalently bound heme, and PhANG transcripts were low—a situation similar to what we found in white leaves of albostrians. PhANG expression was restored to control levels by heme overexpression and in a gun1sig2 double mutant (Woodson et al. 2013). GUN1 is a chloroplast-localized PRR protein. Lack of functional GUN1 restores PhANG transcription in abinotic leaves treated with norflurazon or lincomycin (Nott et al. 2006).

### Plastid proteins, metabolites, and hormones as “plastid factors”

The idea of a “derepressor” RNA transcribed from plastid DNA, leaving the plastids and moving to the nucleus to activate silenced genes (Bradbeer and Börner 1978; Bradbeer et al. 1979) originated from the contradictory results obtained with plastid ribosome deficient rye seedlings grown at elevated temperature and plastid ribosome deficient seedlings of mutant barley lines as outlined above. It was obvious that there could be reasons other than mutant chloroplast DNA in albostrians vs. wild-type chloroplast DNA in rye for the low activities of nuclear-encoded plastid enzymes in barley vs. nearly normal activities in heat-treated rye seedlings: species-specific effects, the presence of ribosomes in heat-treated chlorotic rye leaves, or the developmental stage at which the impaired plastid translation becomes effective. A species-specific difference in the response to plastid ribosome deficiency between barley and rye has been ruled out by studies on barley seedlings grown at elevated temperature (Allsop et al. 1981). Another potentially important difference between the mutants and plants grown at elevated temperature is that the white leaves of albostrians (Saskatoon has not been investigated to comparable extent) do not contain Rubisco (Reichenbächler et al. 1978) while the chlorotic rye leaves show residual Rubisco activity (8% of the activity measured in extracts of green leaves of seedlings grown at permissive temperature and under light; Feierabend and Schrader-Reichhardt 1976), i.e., the chlorotic leaves must possess a few plastid ribosomes as noted by Feierabend and Schrader-Reichhardt (Feierabend and Schrader-Reichhardt 1976; Feierabend 1977b). It has been shown much later that an impaired but not fully blocked translation in plastids due to mutation of the nuclear gene for a plastid ribosomal protein (PRPL11) has no effect on the expression of PhANGs (Pesaresi et al. 2006). Moreover, the lack of ribosomes in albostrians barley is induced already at early stages of embryo development whereas the ribosome deficiency in rye is initiated only during germination at elevated temperature and fully effective later in leaf development (Hagemann and Schulz 1962; Feierabend and Schrader-Reichhardt 1976). An effect of plastid translation on nuclear gene expression has been observed also by use of inhibitors, such as lincomycin, streptomycin, and chloramphenicol. Inhibition of translation in plastids affected nuclear gene expression only if applied within the first 2–3 days of seedling development in mustard or tobacco (Oelmüller et al. 1986; Bajracharya et al. 1987; Gray et al. 1995), i.e., at a time point when the plastids in heat-treated rye leaves should still have a substantial quantity of active ribosomes. Thus, the existence of plastid ribosomes and/or the function of plastid protein synthesis early in leaf development might be responsible for the observed missing or small effects on nuclear gene expression in heat-treated rye seedlings. Consequently, one has to consider also factors other than RNA acting as “plastid factors” or “plastid signals” in plastid-to-nucleus signaling.

The evolutionary origin of the chloroplast genome was another reason to entertain doubts about the existence of a chloroplast gene encoding an RNA (or protein) involved in the regulation of gene expression in the nucleus. The hypothesis about the evolution of chloroplasts from cyanobacteria (Mereschkowski 1905; Marzulis 1970) triggered an intense debate and the generation of alternative hypotheses in the 1970s (e.g., Raff and Mahler 1972; Bogorad 1975). Rudolf Hagemann organized in 1977 an international meeting to discuss the various conceptions about the evolution of eukaryotic
cells. It was Peter Sitte (Sitte 1981) who convinced me of the endosymbiotic origin of chloroplasts and mitochondria by his talk during this meeting where he discussed phylogenetic relationships based on sequences of proteins and nucleic acids from chloroplasts, nuclei, and bacteria in a similar way as was published somewhat later by Schwartz and Dayhoff (1978) and Doolittle and Bonen (1981). If the chloroplast genes have evolved from cyanobacterial genes, then it was not easily understandable how these organelles could possess a gene for a regulator of nuclear gene activities. There is certainly the possibility that a chloroplast gene got another or an additional function during evolution. Nevertheless, the endosymbiosis theory weakened our idea from a chloroplast “derepressor” RNA.

I did not get the permission from the responsible East German authorities to take part together with Bill Bradbeer in the 1978 Spetsai meeting on Chloroplast Development, but Rudolf Hagemann did. Hagemann’s talk during this meeting was devoted to our studies on plastid ribosome deficient mutants of Pelargonium and barley. Because of the reasons outlined above, we decided, in agreement with Bill Bradbeer, that Hagemann should also shortly comment in his talk on the low activities and levels of nuclear-encoded chloroplast proteins in the barley mutants and point out that “it has still to be analyzed whether the postulated ‘chloroplast control principle’, i.e., a control by plastids of nuclear gene expression complementing the ‘cytoplasmic control principle’ formulated by Ellis (1977), is exerted by RNA or ‘derepressor’ proteins or mechanisms not necessarily based on RNA or proteins, e.g., by plant hormones” (Hagemann and Börner 1978).

In 1978, we regarded plastid RNA, proteins, hormones, and other metabolites as alternative candidates for a plastid factor controlling PhANG expression in the nucleo-cytoplasmic compartment (Bradbeer and Börner 1978; Hagemann and Börner 1978; Fig. 1). RNA remained our favorite (Bradbeer et al. 1979), but only for a short time. Research over the last 30 years on plastid-to-nucleus signaling has made it likely that all these types of molecules originating from plastids/chloroplasts might play roles in plastid-to-nucleus signaling (Chan et al. 2016; Kleine and Leister 2016; de Souza et al. 2017).

Support for the hypothesis of plastid-to-nucleus signaling from further studies during the 1980s

A couple of years later, the idea of signals from plastids to nucleus got support from studies on albino mutants and plants treated with herbicides or inhibitors of chloroplast transcription and translation, which demonstrated that impaired chloroplast development reduces the steady state levels of mRNAs transcribed from PhANGs (most analyzed were LHCP transcripts) and the transcription of PhANGs as demonstrated by run-on assays (Harper et al. 1984; Mayfield and Taylor 1984, 1987; Batschauer et al. 1986; Oelmüller and Mohr 1986; Oelmüller et al. 1986; Burgess and Taylor 1988; Giuliano and Scolnik 1988).

Similar to growth under elevated temperatures, treatment of seedlings with inhibitors of carotenoid biosynthesis, like the herbicide norflurazon, results in chlorosis and chloroplast ribosome deficiency (Bartels et al. 1967). In contrast to heat-treated seedlings, but like albostrians and Saskatoon, the herbicide-treated seedlings of ryegrass and mustard (Sinapis alba L.) are characterized by extremely low glycerolaldehyde-phosphate dehydrogenase (NADP+) activities (Feierabend and Schubert 1978; Reiss et al. 1983). Moreover, the activities of the peroxisomal enzymes glycolate oxidase and hydroxypyruvate reductase are very low. Since the effects on plastid and peroxisomal enzymes could only be observed in seedlings grown under white light but not in dark-grown seedlings, they were regarded as consequences of photodamage of the proteins due to the absence of protective carotenoids and not discussed under the aspect of plastid signals to the nucleus (Feierabend and Kemmerich 1983; Reiss et al. 1983). Feierabend and Kemmerich (1983) hypothesized that lipid peroxidation specifically transmits photooxidative damage via membrane contacts from plastids to peroxisomes. Later, it was shown that the expression of nuclear genes for the peroxisomal glycolate oxidase, catalase and hydroxypyruvate reductase is indeed under control of plastid signal(s) (Schwartz et al. 1992; Boldt et al. 1997).

Reviewing research on regulatory interactions between nuclear and plastid genomes from the first reports on
albostrians barley until the end of the 1980s, William Taylor (1989) speculated “What sort of molecule might the chloroplast signal be?...It appears to be a positive regulator, given that Cab is transcriptionally inactive in cells with nongreen plastids or with photooxidatively destroyed chloroplasts...It could easily be a small molecule produced by chloroplast metabolic activity.”

Concluding remarks

This was a wise prediction since in the following 25 years several metabolic pathways in chloroplasts could be shown to be the origin of signals controlling the activities of nuclear genes. Impaired chloroplast transcription and/or chloroplast translation have striking impacts on the expression of nuclear genes (see above) and chloroplast proteins or RNAs are still today discussed as source of plastid signals (or more precisely as start points of signal transduction chains) to the nucleus. However, a block of chloroplast gene expression due to mutation, treatment with inhibitors or photodestruction exerts massive influences on chloroplast development, photosynthesis, pigment synthesis, and other metabolic pathways in the plastids. Such plants are very suitable to detect effects of plastid-to-nucleus signaling but less ideal for identifying specific signaling molecules and signaling pathways. Joan Chory and co-workers found a way to overcome this situation by treating transgenic Arabidopsis plants with norflurazon and screening for rare mutations that lead to the expression of reporter genes under the control of a CAB/LHCP promoter (Susak et al. 1993). In this and similar ways up to now six GUN (genomes uncoupled) mutants have been identified, five of which point specifically to the tetrapyrrole biosynthesis as source of molecules involved in plastid-to-nucleus signaling (Woodson et al. 2011). Plastid gene expression and tetrapyrrole biosynthesis can be regarded as source of “biogenic signals,” which report the developmental state of chloroplasts to the nucleus (Pogson et al. 2008). Further insights into plastid-to-nucleus signaling were provided by studies on specific metabolic processes and components. Singlet oxygen (Wagner et al. 2004), H₂O₂ (Maruta et al. 2012), the redox state of the photosynthetic electron transport chain (Pfalz et al. 2012), 3′-phosphoadenosine 5′-phosphate (Estavillo et al. 2011), the isoprenoid precursor methylerythritolcycloisiphosphate (Xiao et al. 2012), β-cycloclinal (Ramel et al. 2012), and other potential candidates (Chi et al. 2015; Tian 2015; Chan et al. 2016; Kleine and Leister 2016; de Souza et al. 2017) were added to the list of “operational signals” (Pogson et al. 2008), which report changes in the status of the chloroplast in response to the environment to the nucleus. Changes in Ca²⁺ levels might act as both biogenic and operational signal (Guo et al. 2016; de Souza et al. 2017). Also, hormones with plastids as major sites of their biosynthesis like cytokinins, ABA, jasmonic acid, and others are essential players in the interactions between plastids/chloroplasts and nucleus (Tian 2015; Chan et al. 2016) and coordinate the expression of nuclear and plastid genes during chloroplast development, senescence and in response to stress (Zubo et al. 2008, 2011; Yamburenko et al. 2015; Cortleven et al. 2016). Further components of the signal transduction pathways from plastids to the nucleus have been identified; in most cases, these processes remain poorly understood (Chan et al. 2016; Kleine and Leister 2016; de Souza et al. 2017).

Soon after its discovery, plastid-to-nucleus signaling was shown to affect not only the expression of nuclear genes encoding chloroplast-localized proteins as originally assumed (Bradbeer and Börner 1978; Hagemann and Börner 1978) but also the gene for nitrate reductase, a cytoplasmic enzyme (Börner et al. 1986; Oelmüller et al. 1988; Mohr et al. 1992; Hess et al. 1994). Somewhat later was reported that the expression of genes for peroxisomal enzymes (Schwartz et al. 1992; Boldt et al. 1997) and even mitochondrial DNA and RNA levels (Hedtke et al. 1999) are under control of plastid signal(s). The drastic increase in transcription and transcript levels of the gene(s) for chalcone synthase, a cytoplasmic enzyme regarded as fully unrelated to plastids/chloroplasts, as well as the enhanced expression of other stress- and defense-related genes in white albostrians leaves (Hess et al. 1994, 1998) were first indications for roles of plastid-to-nucleus signaling in the response of plants to biotic and abiotic stresses.

Research on plastid-to-nucleus signaling started with the observation of missing small subunits of Rubisco in plastid ribosome deficient mutants. Now, it is obvious that plastid signals are not only involved in harmonizing gene expression in plastids and nucleus for undisturbed chloroplast development. They are also generated to communicate with other regulatory networks and to alter the expression of thousands of genes at the level of transcription and posttranscriptional levels in order to adapt the metabolism inside the plastids and in other parts of the cell to environmental changes sensed by the plastids/chloroplasts (Chan et al. 2016; Kleine and Leister 2016; Woodson 2016; de Souza et al. 2017). Plastid-to-nucleus signaling that we once termed “chloroplast control principle” and speculated it might depend on a chloroplast gene product thereby giving the chloroplast some kind of autonomy (Hagemann and Börner 1978; Bradbeer and Börner 1978; Bradbeer et al. 1979) is, like replication and expression of chloroplast genes, completely under control of the nucleus.

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Compliance with ethical standards

Conflict of interest The author declares that he has no conflict of interest.
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References

Allsop D, Atkinson YE, Bradbeer JW (1981) The effect of nonpermissive temperature on chloroplast development in barley. In: Akoyunoglou G (ed) Proceedings of the International Congress on Photosynthesis. Chloroplast development, vol 5. Balaban International Science Services, Philadelphia, pp 787–795

Armstrong JJ, Surzycki SJ, Moll B, Levine RP (1971) Genetic transcription and translation specifying chloroplast components in Chlamydomonas reinhardtii. Biochemist 10:692–701

Arnason TJ, Walker GWR (1949) An irreversible gene-induced plastid mutation. Canad J Res, Sect C 27:172–178

Bajracharya D, Bergfeld R, Hatzfeld W-D, Klein S, Schopfer P (1987) Regulatory involvement of plastids in the development of peroxisomal enzymes in the cotyledons of mustard (Sinapis alba L.) seedlings. J Plant Physiol 126:421–436

Bartels PG, Matsuda K, Siegel A, Weier TE (1967) Chloroplastic ribosome formation: inhibition by 3-amino-1,2,4-triazole. Plant Physiol 42:736–741

Batschauer A, Mösinger E, Kreuz K, Dörm I, Apel K (1986) The implications of a plastid-derived factor in the transcriptional control of nuclear genes encoding the light-harvesting chlorophyll a/b protein. Eur J Biochem 154:625–634

Blair GE, Ellis RJ (1973) Protein synthesis in chloroplasts. I. Light-driven synthesis of the large subunit of fraction I protein by isolated pea chloroplasts. Biochim Biophys Acta 319:223–234

Bogorad L (1975) Evolution of organelles and eukaryotic genomes. Science 188:891–898

Bödtker T, Knoth R, Herrmann F, Hagemann R (1974) Struktur und Funktion der genetischen Information in den Plastiden. X. Das Fehlen von cytoplasmatischer und plastidaler Proteinsynthese beim Aufbau der Plastiden. VI. Zur Funktion von Plastiden-DNA, Kern-DNA, plastid ribosome-deficient mutant. In: Abstracts International Conference on Regulation of Developmental Processes in Plants, Halle (Saale), p 62

Börner T (1973) Struktur und Funktion der genetischen Information in den Plastiden. VI. Zur Funktion von Plastiden-DNA, Kern-DNA, cytoplasmatischer und plastidaler Proteinsynthese beim Aufbau der Chloroplasten - eine tabellarische Übersicht. Biol Zbl 92:545–561

Börner T (1977) The absence of cytoplasmically synthesized proteins in a plastid ribosome-deficient mutant. In: Abstracts International Conference on Regulation of Developmental Processes in Plants, Halle (Saale), p 62

Börner T (1981) Enzymes of plastid ribosome-deficient mutants. Ferredoxin: NADP+ reductase. Biochem Physiol Pflanzen 176: 737–743

Börner T, Meister A (1980) Chlorophyll and carotenoid content of ribosome-deficient plastids. Photosynthesica 14:588–593

Börner T, Sears BB (1986) Plastome mutants. Plant Mol Biol Rep 4:69–92

Börner T, Knoth R, Herrmann F, Hagemann R (1972) Struktur und Funktion der genetischen Information in den Plastiden. V. Das Fehlen von ribosomal RNA in den Plastiden der Plastommutante "Mrs. Parker" von Pelargonium zonale Ait. Theor Appl Genet 42:3–11

Börner T, Herrmann FH, Hagemann R (1973) Plastid ribosome deficient mutants in Pelargonium zonale. FEBS Lett 37:17–19

Börner T, Knoth R, Herrmann F, Hagemann R (1974) Struktur und Funktion der genetischen Information in den Plastiden. X. Das Fehlen von Fraktion-I-Protein in den weißen Plastiden einiger Sorten von Pelargonium zonale. Biochem Physiol Pflanzen 165:429–432

Börner T, Schumann B, Hagemann R (1976) Biochemical studies on a plastid ribosome-deficient mutant of Hordeum vulgare. In: Bücher T, Neupert W, Sebald W, Werner S (eds) Genetics and biogenesis of chloroplasts and mitochondria. Elsevier / North-Holland Medical Press, Amsterdam, pp 41–48

Börner T, Mendel RR, Sichmann J (1986) Nitrate reductase is not accumulated in chloroplast-ribosome-deficient mutants of higher plants. Planta 169:202–207

Bradbeer JW, Börner T (1978) Activities of glyceraldehyde-phosphate dehydrogenase (NADP+) and phosphoribulokinase in two barley mutants deficient in chloroplast ribosomes. In: Akoyunoglou G, Argyroudi-Akoyunoglou I (eds) Chloroplast development. North Holland, Amsterdam, pp 727–732

Bradbeer JW, Atkinson YE, Börner T, Hagemann R (1979) Cytoplasmic synthesis of plastid poly peptides may be controlled by plastid-synthesised RNA. Nature 279:816–817

Brandt P (1976) Regulation of the Synthesis of Plastidenproteinen in Euglena gracilis, Stamm Z. Planta 130:81–83

Brzezowska P, Richter AS, Grimm B (2015) Regulation and function of tetrapyrrole biosynthesis in plants and algae. Biochim Biophys Acta 1847:968–985

Bünger W, Feierabend J (1980) Capacity for RNA synthesis in 70S ribosome-deficient plasts of heat-bleached rye leaves. Planta 149:163–169

Burgess DG, Taylor WC (1988) Chloroplast photooxidation affects the transcription of a nuclear gene family. Mol Gen Genet 214:89–96

Chao PH, Wildman SG (1972) Chloroplast DNA codes for the primary structure of the large subunit of fraction I protein. Biochim Biophys Acta 277:677–680

Chan XX, Phua SY, Crisp P, McQuinn R, Pogson BJ (2016) Learning the languages of the chloroplast: retrograde signaling and beyond. Annu Rev Plant Biol 67:25–53

Chi W, Feng P, Ma J, Zhang L (2015) Metabolites and chloroplast retrograde signalling.Curr Opin Plant Biol 25:32–38

Cortleven A, Marg I, Yambureno MV, Schlicke H, Hill K, Grimm B, Schaller GE, Schmülling T (2016) Cyanobacteria regulates the etioplast-chloroplast transition through the two-component signaling system and activation of chloroplast-related genes. Plant Physiol 172:464–478

Criddle RS, Dau B, Kleinkopf DE, Huffaker RC (1970) Differential synthesis of ribulose diphosphate subunits. Biochem Biophys Res Commun 41:621–627

Day A, Ellis THN (1984) Chloroplast DNA deletions associated with wheat plants regenerated from pollen: possible basis for maternal inheritance of chloroplasts. Cell 39:359–368

De Souza A, Wang J-Z, Dehesh K (2017) Retrograde signals: integrators of interorganellar communication and orchestrators of plant development. Annu Rev Plant Biol 68. doi:10.1146/annurev-arplant-042916-041007

Doolittle WF, Bonen L (1981) Molecular sequence data indicating an endosymbiotic origin for plastids. Ann N Y Acad Sci 361:248–259

Ellis RJ (1975) Inhibition of plastid protein synthesis by lincomycin and 2-(4-methyl-2,6-dinitroanilino)-N-methylpropionamide. Phytochemistry 14:89–93

Ellis RJ (1977) Protein synthesis by isolated chloroplasts. Biochim Biophys Acta 463:285–315

Ellis RJ, Hartley MR (1971) Sites of synthesis of chloroplast proteins. Nature New Biol 233:193–196

Estavillo GM, Crisp PA, Porsasrirwong W, Wirtz M, Collinge D, Carrie C, Giraud E, Whelan J, David P, Javot H, Brelley C, HELL R, Marin E, Pogson BJ (2011) Evidence for a SAL1-PAP chloroplast retrograde pathway that functions in drought and high light signaling in Arabidopsis. Plant Cell 23:3992–4012

Falk H, Liedvogel B, Sitte P (1974) Circular DNA in isolated chloroplasts. Z Naturforschg 29c:541–544
Feierabend J (1977a) Cooperation of cytoplasmic and plastidic protein synthesis in the biogenesis of chloroplasts. In: Abstracts of the International Conference on Regulation of Developmental Processes in Plants, Halle (Saale), p 61
Feierabend J (1977b) Capacity for chlorophyll synthesis in heat-bleached 70S ribosome-deficient yeast cells. Planta 155:83–88
Feierabend J, Kemmerich P (1983) Mode of inference of chlorosis-inducing herbicides with peroxisomal enzyme activities. Physiol Plant 57:346–351
Feierabend J, Mikus M (1977) Occurrence of a high temperature sensitive mutant. Planta 137:20–27
Feierabend J, Schrader-Reichhardt U (1976) Biochemical differentiation of plastids and other organelles in yeast cells with a high-temperature-deficient of plastid ribosomes. Planta 129:133–145
Feierabend J, Schubert B (1978) Comparative investigation of the action of several chlorosis-inducing herbicides on the biogenesis of chloroplasts and leaf microbodies. Plant Physiol 61:1077–1022
Gibor A, Granick S (1964) Plastids and mitochondria: inheritable systems. Science 145:890–897
Gillham NW (1974) Genetic analysis of the chloroplast and mitochondrial gene expression. In: Abstracts of the International Conference on Regulation of Developmental Processes in Plants, Halle (Saale), p 61
Givan AL, Criddle RS (1972) Ribulose diphosphate carboxylase from Chlamydomonas reinhardtii. Purification, properties and mode of synthesis in the cell. Arch Biochem Biophys 149:153–154
Glowatzki E, Scolnik PA (1988) Transcription of two photosynthesis-associated nuclear gene families correlates with the presence of chloroplasts in leaves of the variegated tomato ghost mutant. Plant Physiol 86:7–9
Givan AL, Criddle RS (1972) Ribulose diphosphate carboxylase from Chlamydomonas reinhardtii. Purification, properties and mode of synthesis in the cell. Arch Biochem Biophys 149:153–154
Gray JC, Somarajah R, Zabron AA, Duckett CM, Khan MS (1995) Chloroplast control of nuclear gene expression. In: Mathis P (ed) Photosynthesis, from light to biosphere, vol 3. Kluwer, Dordrecht, pp 543–550
Gunning BES (1965) The fine structure of chloroplast stroma following adenosine-octetotidex fixation. J Cell Biol 2:79–93
Guo H, Feng P, Chi W, Sun X, Xu X, Li Y, Ren D, Lu C, Rocha JD, Leister D, Zhang L (2016) Plastid-nucleus communication involves calcium-modulated MAPK signalling. Nature Comms 7:12173
Hagemann R, Börner T (1978) Plastid ribosome-deficient mutants of higher plants as a tool in studying chloroplast biogenesis. In: Akoyunoglou G, Argyroudi-Akoyunoglou JG (eds) Chloroplast development. North Holland, Amsterdam, pp 709–720
Hagemann R, Scholz F (1962) Ein Fall Gen-induzierter Mutationen des Plasmoytys bei Gerste. Züchter 32:50–59
Hagemann R, Börner T, Knot B (1973) Plastid ribosome deficiency in plastid mutants of Pelargonium and Hordeum. Genetics 74(suppl): 103–104
Hajdukiewicz PT, Allison LA, Maliga P (1997) The two RNA polymerases encoded by the nuclear and the plastid compartments transcribe distinct groups of genes in tobacco plastids. EMBO J 16:4041–4048
Han CD, Coe EH Jr, Martienssen RA (1992) Molecular cloning and characterization of ioiap (ij), a pattern stripping gene of maize. EMBO J 11:4037–4046
Hanaoka M, Kanamaru K, Takahashi H, Tanaka K (2003) Molecular genetic analysis of chloroplast gene promoters dependent on SIG2, a nuclear-encoded sigma factor for the plastid-encoded RNA polymerase, in Arabidopsis thaliana. Nucl Acids Res 31:7090–7098
Harada T, Sato T, Asaka D, Matsukawa I (1991) Large-scale deletions of rice plastid DNA in anther culture. Theor Appl Genet 81:157–161
Harpster MH, Mayfield SP, Taylor WC (1984) Effects of pigment-deficient mutants on the accumulation of photosynthetic proteins in maize. Plant Mol Biol 3:258–263
Hedtke B, Wagner L, Börner T, Hess WR (1999) Inter-organelar crosstalk in higher plants: impaired chloroplast development affects mitochondrial gene an transcript levels. Plant J 19:635–643
Herrmann F (1971) Genetic control of pigment-protein complexes I and II of the plastid mutant enalba-1 of Antirrhinum majus. FEBS Lett 19:267–269
Herrmann RG (1972) Do chloroplasts contain DNA? II. Isolation and characterization of DNA from chloroplasts, chloroplasts, mitochondria and nuclei of Narcissus. Protoplasma 74:7–17
Herrmann RG, Kowallik KV (1970) Multiple amounts of DNA related to the size of chloroplasts II. Comparison of electron-microscopic and autoradiographic data. Protoplasma 69:365–372
Hess WR, Blank-Huber M, Fiedler B, Börner T, Rüdiger W (1992a) Chlorophyll synthetase and chloroplast rRNA0glu are present in heat-induced, ribosome-deficient plastids. J Plant Physiol 139:427–430
Hess WR, Schendel R, Rüdiger W, Börner T (1992b) Protochlorophyllide oxidoreductase and chlorophyll synthetase are present in a barley mutant unable to synthesize d-aminolevulinic acid from rRNA0glu. Planta 188:19–27
Hess WR, Prombona A, Fiedler B, Subramanian AR, Börner T (1993) Chloroplast rps13 and the rpoB/C1/C2 gene cluster are strongly transcribed in ribosome-deficient plastids: evidence for a functioning non-chloroplast encoded RNA polymerase. EMBO J 12:563–671
Hess WR, Müller A, Nagy F, Börner T (1994) Ribosome-deficient plastids affect transcription of light-induced nuclear genes: evidence for a plastid-derived signal. Mol Gen Genet 242:305–312
Hess WR, Goz R, Börner T (1998) Analysis of randomly selected cDNAs reveals the expression of stress- and defense-related genes in the barley mutant albostrans. Plant Sci 133:191–201
Hollingsworth MJ, Hallick RB (1982) Euglena gracilis chloroplast transfer RNA transcription units. Nucleotide sequence analysis of the RNA-tRNAHis-rRNArnp-tRNA0glu-rRNA0gly gene cluster. J Biol Chem 257:12795–12799
Johanne Meier U, Howell SH (1984) Regulation of light-harvesting chlorophyll-binding protein mRNA accumulation in Chlamydomonas reinhardtii. Possible involvement of chlorophyll synthesis precursors. J Biol Chem 259:13541–13549
Kawashima N, Wildman SG (1970) Fraction I protein. Annu Rev Plant Physiol 21:325–358
Kirk JTO (1971) Chloroplast structure and biogenesis. Annu Rev Biochem 40:161–196
Kirk JTO, Tilney-Bassett RA (1967) The plastids. Freeman, London
Kleine T, Leister D (2016) Retrograde signalling: organelles go network-oriented. Biochim Biophys Acta 1857:1313–1325
Kolodner R, Tewari KK (1975) The molecular size and conformation of the chloroplast DNA from higher plants. Biochim Biophys Acta 402:372–390
Kropat J, Oster U, Rüdiger W, Beck CF (1997) Chlorophyll precursors are signals of chloroplast origin involved in light induction of nuclear heat-shock genes. Proc Natl Acad Sci U S A 94:14168–14172
Kung SD (1977) Expression of chloroplast genomes in higher plants. Annu Rev Plant Physiol 28:401–437
Kunz M, Weil JH, Steinmetz A (1984) Nucleotide sequence of a 2 kb BamHI fragment of Vicia faba chloroplast DNA containing the genes for threonine, glutamic acid and tyrosine transfer RNAs. Nucleic Acids Res 12:5037–5047
Lukens JH, Mathews DE, Durbin RD (1988) Effect of tagetitoxin on the levels of ribulose 1,5-bisphosphate carboxylase, ribosomes, and RNA in plastids of wheat leaves. Plant Physiol 84:808–813
Margulis L (1970) Origin of eukaryotic cells. Yale University Press, New Haven
Maruta T, Noshi M, Tanouchi A, Tamoi M, Yabuta Y, Yoshimura K, Ishikawa T, Shigeoka S (2012) H2O2-triggered retrograde signaling from chloroplasts to nucleus plays specific role in response to stress. J Biol Chem 287:11717–11729
Mayfield SP, Taylor WC (1984) Carotenoid-deficient maize seedlings fail to accumulate light-harvesting chlorophyll a/b binding protein (LHCP) mRNA. Eur J Biochem 144:79–84

Mayfield SP, Taylor WV (1987) Chloroplast photooxidation inhibits the expression of a set of nuclear genes. Mol Gen Genet 208:309–314

McCrea JM, Hershberger CL (1978) Chloroplast DNA codes for tRNA from cytoplasmic polyribosomes. Nature 274:717–719

Mereschkowski KS (1905) Über Natur und Ursprung der Chloroplasten. Zool Stud 14:419–465

Oelmüller R, Schuster C, Mohr H (1988) Physiological characterization of a plastidic signal required for nitrate-induced appearance of nitrite reductase gene expression by light, nitrate and a plastidic factor. Bot Acta 105:81–89

Oelmüller R, Mohr H (1986) Photooxidative destruction of chloroplasts and their consequences for expression of nuclear genes. Planta 167:106–113

Oelmüller R, Levitan I, Bergfeld R, Rajasekhar VK, Mohr H (1986) Expression of nuclear genes as affected by treatments acting on plastids. Planta 168:482–492

Oelmüller R, Schuster C, Mohr H (1988) Physiological characterization of a plastidic signal required for nitrate-induced appearance of nitrate and nitrite reductases. Planta 174:75–83

Pesaresi P, Masiero S, Eubel H, Bhushan S, Glaser E, Salamini F, Leister D (2006) Nuclear photosynthetic gene expression is synergistically modulated by rates of protein synthesis in chloroplasts and mitochondria. Plant Cell 18:970–991

Pfalz J, Liebers M, Hirth M, Grühler B, Holzegel U, Schröter Y, Dietzel L, Pfannschmidt T (2012) Environmental control of plant nuclear gene expression by chloroplast redox signals. Front Plant Sci 3:257

Pogson BJ, Woo NS, Förster B, Braun HP, Bhushan S, Glaser E, Salamini F, Leister D (2006) Nuclear photosynthetic gene expression is synergistically modulated by rates of protein synthesis in chloroplasts and mitochondria. Plant Cell 18:970–991

Pogson BJ, Woo NS, Förster B, Small ID (2008) Plastid signalling to the nucleus and beyond. Trends Plant Sci 13:602

Raf R, Müller HR (1972) The non symbiotic origin of mitochondria. Science 177:575–582

Ramel F, Birtic S, Ginies C, Soubigou-Taconnat L, Triantaphylidès C, Wohlenberg T, Link G, Thien W, Mohr H (1983) Photooxidative destruction of chloroplasts and its consequences for cytosolic enzyme levels and plant development. Plant Mol Biol 17:813–823

Reichenbächer D, Börner T, Richter J (1978) Untersuchungen am Fraktion-I–Protein der Gerste mit Hilfe quantitativer Immunelektrophorese. Biochem Physiol Pflanzen 172:53–60

Reiss T, Bergfeld R, Link G, Thien W, Mohr H (1983) Photooxidative destruction of chloroplasts and its consequences for cytosolic enzyme levels and plant development. Plant Mol Biol 17:59–80

Schwartz RM, Dayhoff MO (1978) Origins of prokaryotes, eukaryotes, mitochondria, and chloroplasts, Science 199:395–403

Schwarz BW, Daniel SG, Becker WM (1992) Photooxidative destruction of chloroplasts leads to reduced expression of peroxisomal NADH-dependent hydroxypropionate reductase in developing cucumber cotyledons. Plant Physiol 99:681–685

Siemenroth A, Wollgiehn R, Neininger A, Seith B (1992) Control of nitrate reductase and nitrite reductases. Planta 174:75

Siemenroth A, Wollgiehn R, Neumann D, Börner T (1981) Synthesis of LHCP mRNA. Eur J Biochem 144:79–84

Sitte P (1977) Chromoplasten—bunte Objekte der modernen Zellbiologie. Biologie in Unserer Zeit 7:65–74

Sitte P (1981) Die Endosymbionten-Hypothese—eine kritische Betrachtung zur Zellevolution. Nova Acta Leopoldina NF 56, No 251:41–58

Susek RE, Ausubel FM, Chory J (1993) Signal transduction mutants of Arabidopsis uncouple nuclear CAB and RBCS gene expression from chloroplast development. Cell 74:787–799

Taylor WC (1989) Regulatory interactions between nuclear and plastid genomes. Annu Rev Plant Physiol Plant Mol Biol 40:211–233

Tewari KK (1971) Genetic autonomy of extranuclear organelles. Annu Rev Plant Physiol 22:141–168

Tian L (2015) Recent advances in understanding carotenoid-derived signalling molecules in regulating plant growth and development. Front Plant Sci 6:790

Vaisberg AJ, Schiff JA, Li L, Freedman Z (1976) Events surrounding the early development of Euglena chloroplasts. VIII. Photocontrol of the source of reducing power for chloramphenicol reduction by the ferredoxin-NADP reductase system. Plant Physiol 57:594–601

Wagner D, Przybyla D, Op den Camp R, Kim C, Landgraf F, Lee KP, Würsch M, Laloï C, Nater M, Hideg E, Apel K (2004) The genetic basis of singlet oxygen-induced stress responses of Arabidopsis thaliana. Science 306:1183–1185

Walter G, Müller A, Hoffmann P, Börner T (1995) Tryptophylpyrrole synthesis in leaves and roots of a barley albina mutant with extremely low amount of plastid glutamyl-RNA. In: Mathis P (ed) Photosynthesis from light to biosphere, vol Vol. III. Kluwer Academic Publ, Dordrecht, pp 937–940

Woodson JD (2016) Chloroplast quality control—balancing energy production and stress. New Physiol 212:36–41

Woodson JD, Perez-Ruiz JM, Chory J (2011) Heme synthesis by plastid ferrochelatase I regulates nuclear gene expression in plants. Curr Biol 21:897–903

Woodson JD, Perez-Ruiz JM, Schmitz RJ, Ecker JR, Chory J (2013) Sigma factor-mediated plastid retrograde signals control nuclear gene expression. Plant J 73:1–13

Xiao Y, Savchenko T, Baidoo EE, Chehab WE, Hayden DM, Tolstikov V, Corwin JA, Kliebenstein DJ, Keasling JD, Dehesh K (2012) Retrograde signaling by the plastidial metabolite MeEPP regulates expression of nuclear stress-response genes. Cell 149:1525–1535

Yamamura MF, Zubo YO, Börner T (2015) Abscisic acid affects transcription of chloroplast genes via protein phosphatase 2C-dependent activation of nuclear genes: repression by guanosine-3'5'-bisphosphate and activation by sigma factor 5. Plant J 82:1030–1041

Yaronskaya E, Ziemann V, Walter G, Averina N, Börner T, Grimm B (2012) The primary transcriptome of barley chloroplasts: numerous noncoding RNAs and the dominating role of the plastid-encoded RNA polymerase. Plant Cell 24:123–136

Zubko MK, Day A (1998) Stable albinism induced without mutagenesis: a model for ribosome-free plastid inheritance. Plant J 5:265–271

Zubko YO, Iamburencu MV, Selivankina SY, Shakerova FM, Avalbaev AM, Kudryakova NV, Zubkova NK, Liere K, Kulaeva ON, Kusnetsov VV, Börner T (2008) Cytokinins stimulates chloroplast transcription in detached barley leaves. Plant Physiol 148:1082–1093

Zubko YO, Iamburencu MV, Kusnetsov VV, Börner T (2011) Methyl jasmonate, gibberelic acid, and auxin affect transcription and transcript accumulation of chloroplast genes in barley. J Plant Physiol 168:1335–1344