

Succinyl-Coenzyme A Synthetase and its Role in δ -Aminolevulinic Acid Biosynthesis in *Euglena gracilis*¹

Sandra M. Mayer and Samuel I. Beale*

Division of Biology and Medicine, Brown University, Providence, Rhode Island 02912

ABSTRACT

Euglena gracilis cells synthesize the key tetrapyrrole precursor, δ -aminolevulinic acid (ALA), by two routes: plastid ALA is formed from glutamate via the transfer RNA-dependent five-carbon route, and ALA that serves as the precursor to mitochondrial hemes is formed by ALA synthase-catalyzed condensation of succinyl-coenzyme A and glycine. The biosynthetic source of succinyl-coenzyme A in *Euglena* is of interest because this species has been reported not to contain α -ketoglutarate dehydrogenase and not to use succinyl-coenzyme A as a tricarboxylic acid cycle intermediate. Instead, α -ketoglutarate is decarboxylated to form succinic semialdehyde, which is subsequently oxidized to form succinate. Desalted extract of *Euglena* cells catalyzed ALA formation in a reaction that required coenzyme A and GTP but did not require exogenous succinyl-coenzyme A synthetase. GTP could be replaced with ATP. Cell extract also catalyzed glycine- and α -ketoglutarate-dependent ALA formation in a reaction that required coenzyme A and GTP, was stimulated by NADP⁺, and was inhibited by NAD⁺. Succinyl-coenzyme A synthetase activity was detected in extracts of dark- and light-grown wild-type and nongreening mutant cells. *In vitro* succinyl-coenzyme A synthetase activity was at least 10-fold greater than ALA synthase activity. These results indicate that succinyl-coenzyme A synthetase is present in *Euglena* cells. Even though the enzyme may play no role in the transformation of α -ketoglutarate to succinate in the atypical tricarboxylic acid cycle, it catalyzes succinyl-coenzyme A formation from succinate for use in the biosynthesis of ALA and possibly other products.

The phytoflagellate *Euglena gracilis* synthesizes the key heme and Chl precursor, ALA², from glutamate via the transfer RNA-dependent five-carbon pathway in the plastids of green cells. Unlike other oxygenic organisms, *Euglena* also has the ability to form ALA via the route used by animals, yeasts, and some bacteria, in which the condensation of glycine and succinyl-coenzyme A is catalyzed by ALA synthase (succinyl-CoA:glycine C-succinyltransferase [decarboxylating], EC 2.3.1.37) (1, 10). ALA that serves as precursor to Chl is synthesized solely by the five-carbon route in *Euglena* (15, 22), whereas ALA that is used for mitochondrial tetrapyrrole synthesis is formed exclusively by the ALA synthase route (22). Certain nongreening *Euglena* mutant strains that do not undergo light-induced plastid differentiation appear to

lack the ability to form ALA from glutamate *in vivo*, and these strains rely on the ALA synthase route for all tetrapyrrole precursor biosynthesis (13, 22).

In the mitochondria of animals and yeasts, the ALA precursor, succinyl-CoA, is formed as the product of the α -ketoglutarate dehydrogenase reaction, a step of the tricarboxylic acid cycle. *Euglena* has been reported to have an atypical tricarboxylic acid cycle. The α -ketoglutarate dehydrogenase complex (EC 1.2.4.2, EC 2.3.1.61, EC 1.6.4.3) is apparently absent, and succinyl-CoA is not an intermediate of succinate formation from α -ketoglutarate in the tricarboxylic acid cycle of *Euglena* (8, 18). Instead, *Euglena* mitochondria contain α -ketoglutarate decarboxylase (EC 4.1.1.), which decarboxylates α -ketoglutarate to form succinic semialdehyde (17). Another enzyme, succinic semialdehyde dehydrogenase (succinate-semialdehyde:NADP oxidoreductase, EC 1.2.1.16), then oxidizes the succinic semialdehyde to form succinate (20).

The absence of α -ketoglutarate dehydrogenase in *Euglena* raises the question of the source of succinyl-CoA substrate for ALA synthesis via the ALA synthase reaction. Therefore, we examined desalted *Euglena* extracts for the presence of succinyl-CoA synthetase (succinic thiokinase and succinate:CoA ligase, EC 6.2.1.4 and EC 6.2.1.5, respectively). Activity was found in light- and dark-grown wild-type and nongreening mutant cell extracts. *Euglena* extract formed ALA from α -ketoglutarate plus glycine via a coupled reaction in the presence of the cofactors required for α -ketoglutarate decarboxylase, succinic semialdehyde dehydrogenase, succinyl-CoA synthetase, and ALA synthase.

MATERIALS AND METHODS

Growth of Cells

Axenic cultures of *Euglena gracilis* Klebs var Pringsheim wild-type strain Z and nongreening mutant strain W₁₄ZNaIL were grown in glucose-based heterotrophic medium in the light or complete darkness as previously described (1). Cells referred to as dark grown had been subcultured in continuous darkness for at least 1 year before these experiments.

Cell Extraction for Enzyme Preparation

Cell cultures in the exponential growth phase were thoroughly chilled on ice under the light conditions in which they were grown. All subsequent operations were performed at 0 to 4°C. Cells were harvested by centrifugation, washed, resuspended, and disrupted by sonication as previously described (12). Cell debris and unbroken cells were removed by centrif-

¹ Supported by National Science Foundation grant DCB91-03253.

² Abbreviations: ALA, δ -aminolevulinic acid; PALP, pyridoxal-5-phosphate.

ugation for 10 min at 10,000g. The clarified supernatant fluid was fractionated by differential $(\text{NH}_4)_2\text{SO}_4$ precipitation between 35 and 60% of saturating concentration of $(\text{NH}_4)_2\text{SO}_4$, in the presence of 5 mM EDTA and 0.004% PMSF, and desalted by passage through Sephadex G-25 that was preequilibrated and eluted with column buffer (1.0 M glycerol, 50 mM Tricine, 15 mM MgCl_2 , 1.0 mM DTT, 0.02 mM PALP [pH 7.9]). The protein-containing effluent was supplemented with 0.004% PMSF and stored at -75°C .

ALA Synthase Assay

ALA synthase was assayed by modifications of a previously described method (5). Incubation was for 30 min at 40°C in 1 mL of medium (75 mM Hepes, 50 mM glycine, 5 mM EDTA, 0.1 mM PALP [pH 7.8]) supplemented with either 0.1 mM succinyl-CoA, prepared by the method of Simon and Shemin (19), or a succinyl-CoA-generating system consisting of 25 mM succinate, 1 mM GTP, 0.2 mM CoA, and sufficient succinyl-CoA synthetase to catalyze the formation of 0.2 μmol succinyl-CoA per min. In some incubations where indicated, 5 mM levulinate was included to inhibit the conversion of ALA to porphobilinogen. Reactions were terminated by addition of one-tenth volume of 1.0 M citric acid and one volume of 10% (w/v) SDS followed by heating for 3 min at 95°C . ALA was isolated on Dowex 50W-X8 (Na) as previously described (23), ethylacetoacetate was added, and the solutions were heated to 95°C for 15 min to form 1-methyl-2-carboxyethyl-3-propionic acid pyrrole (11). The product was quantitated spectrophotometrically after reaction with an equal volume of Ehrlich-Hg reagent (21), using a Cary model 219 spectrophotometer (Varian Instruments, Palo Alto, CA). The A_{553} of control samples containing heat-denatured desalted extract was subtracted from those of incubated samples to determine net A_{553} values, and ALA was calculated from a standard curve.

In some cases where indicated, the Dowex 50W-X8 ALA isolation step was omitted. These reactions were terminated by addition of 50 μL of 100% (w/v) aqueous TCA, precipitated proteins were removed by centrifugation, and the supernatant fluid was reacted with ethylacetoacetate and Ehrlich-Hg reagent as described above.

Assay for ALA Formation from Glycine and Succinate

These assays were done essentially as described above for ALA synthase, except that neither succinyl-CoA nor a succinyl-CoA synthetase system was provided. Instead, the incubation mixture was supplemented with 25 mM succinate, 0.2 mM CoA, and 1 mM GTP or ATP.

Assay for ALA Formation from Glycine and α -Ketoglutarate

Incubation was for 30 min at 43°C in 1 mL of medium (1 M glycerol, 75 mM glycine, 50 mM Tricine, 10 mM α -ketoglutarate, 5 mM MgCl_2 , 5 mM KCl, 5 mM EDTA, 5 mM levulinate, 1 mM GTP, 0.5 mM NAD^+ , 0.5 mM NADP^+ , 1 mM DTT, 0.2 mM thiamine pyrophosphate, 0.1 mM PALP, 0.1 mM CoA [pH 8.2]). Reactions were terminated, and ALA was isolated and quantitated as described above.

Succinyl-CoA Synthetase Assay

Incubation was for 30 min at 43°C in 1 mL of medium (1 M glycerol, 50 mM Tricine, 25 mM succinate, 10 mM MgCl_2 , 5 mM GTP, 1 mM DTT, 0.2 mM CoA, 0.02 mM PALP [pH 8.0]). Reactions were terminated by cooling to 0°C .

Succinyl-CoA was quantitated by modifications of the method of Lipmann and Tuttle (9), in which succinamic acid is chemically generated from succinyl-CoA and determined by the formation of a colored complex with Fe^{3+} . To the cooled incubation mixture was added 0.25 mL of a 1:1 (v/v) mixture of 56% (w/v) aqueous NH_2OH and 28% (w/v) aqueous NaOH. Next, 0.25 mL of 0.2 M Na-acetate (pH 5.4) buffer was added, and the solution was mixed and allowed to stand at room temperature for 10 min. Then, 0.25 mL of 24% (w/v) aqueous TCA was added, followed by 0.25 mL of 6 N HCl. Finally, 0.25 mL of 10% (w/v) $\text{FeCl}_3 \cdot 6\text{H}_2\text{O}$ in 0.1 N HCl was added. After standing at room temperature for 5 min, the solution was clarified by centrifugation and the A_{540} was determined. Net absorbance was determined by subtraction of the A_{540} of samples derived from incubations containing heat-denatured *Euglena* protein. Concentration was calculated from a standard curve based on samples containing known amounts of succinamic acid, prepared from succinic anhydride as described previously (9).

Other Procedures

Cell population densities were determined with a Coulter Counter (model ZBI, Coulter Electronics). Protein concentrations were determined by the dye-binding method of Bradford (2) using BSA as the standard.

Chemicals

Succinyl-CoA synthetase (porcine heart succinic thiokinase) was purchased from Sigma. All chemicals were from Sigma, Fisher, and Research Organics.

RESULTS

ALA Formation in the Presence and Absence of Added Succinyl-CoA Synthetase

In previous studies of ALA synthase activity in *Euglena* extracts, the incubation mixtures were either supplied with chemically synthesized succinyl-CoA or supplemented with commercial succinyl-CoA synthetase and GTP to ensure the availability of the ALA synthase substrate, succinyl-CoA (1, 4-6, 16). *Euglena* extract was tested for the ability to catalyze ALA synthesis in the absence of added succinyl-CoA synthetase. Unsupplemented extract formed nearly as much ALA as did extract supplemented with succinyl-CoA synthetase (Table I). The reaction required nondenatured protein and nucleoside triphosphate.

Nucleoside Triphosphate Requirement for ALA Formation

Succinyl-CoA synthetases from different sources have different nucleoside triphosphate requirements, some requiring GTP and others ATP (14). *Euglena* extract supplied with

Table I. Effect of Added Succinyl-CoA Synthetase on ALA Synthesis

Desalted protein extract from dark-grown wild-type *Euglena* cells was assayed for ALA synthase-catalyzed ALA-forming ability at 40°C in the presence and absence of added succinyl-CoA synthetase. Except where indicated, incubations (1 mL) contained 75 mM Hepes (pH 7.8), 50 mM glycine, 25 mM succinate, 5 mM EDTA, 1 mM GTP, 0.2 mM CoA, 0.1 mM PALP, sufficient succinyl-CoA synthetase to catalyze the formation of 0.2 μ mol succinyl-CoA per min, and 1 mg of *Euglena* protein. In this experiment, ALA was not purified by Dowex 50W-X8 chromatography before determination.

Incubation Content	ALA Formation	
	nmol mg ⁻¹ protein	%
Complete	9.4	100
- <i>Euglena</i> protein	0.0	0
-Succinyl-CoA synthetase	8.2	87
-GTP	0.0	0

either of these nucleoside triphosphates formed approximately equal amounts of ALA (Table II).

Glycine- and α -Ketoglutarate-Dependent ALA Formation

Euglena extract was tested for the ability to form ALA from α -ketoglutarate via the coupling of four enzymic steps: α -ketoglutarate decarboxylase, succinic semialdehyde dehydrogenase, succinyl-CoA synthetase, and ALA synthase. Significant ALA formation occurred only in incubations containing α -ketoglutarate, glycine, GTP, and CoA (Table III). Although the reaction was stimulated by NADP⁺, an absolute requirement could not be established. The residual activity in the absence of added NADP⁺ was probably caused by carryover of endogenous pyridine nucleotide through the purification steps of differential (NH₄)₂SO₄ precipitation and gel filtration, as has been previously observed for other pyridine nucleotide-requiring *Euglena* enzymes (13). In contrast to the stimulation by NADP⁺, a slight inhibition by NAD⁺ was observed.

Under incubation conditions similar to those used for the above assay, ALA synthesis from glycine and chemically synthesized succinyl-CoA did not require nucleoside triphos-

Table II. Nucleoside Triphosphate Requirement for ALA Synthesis

Desalted protein extract from dark-grown wild-type *Euglena* cells was assayed for ALA synthase-catalyzed ALA-forming ability at 40°C in the presence and absence of added GTP and ATP. Incubations (1 mL) contained 75 mM Hepes (pH 7.8), 50 mM glycine, 25 mM succinate, 5 mM EDTA, 1 mM GTP or ATP, 0.2 mM CoA, 0.1 mM PALP, and 1 mg of *Euglena* protein. In this experiment, ALA was not purified by Dowex 50W-X8 chromatography before determination.

Nucleoside Triphosphate	ALA Formation	
	nmol mg ⁻¹ protein	%
GTP	6.44	100
ATP	6.56	102
None	0.13	2

Table III. Glycine- and α -Ketoglutarate-Dependent ALA Synthesis

Desalted protein extract from dark-grown wild-type *Euglena* cells was assayed for ALA formation from α -ketoglutarate and glycine at 43°C. Except where indicated, the incubation mixture (1 mL) contained 1 M glycerol, 50 mM Tricine (pH 8.2), 75 mM glycine, 10 mM α -ketoglutarate, 5 mM MgCl₂, 5 mM KCl, 5 mM EDTA, 5 mM levulinate, 1 mM GTP, 0.5 mM NAD⁺, 0.5 mM NADP⁺, 1 mM DTT, 0.2 mM thiamine pyrophosphate, 0.1 mM PALP, 0.1 mM CoA, and 5.4 mg of *Euglena* protein.

Incubation Content	ALA Formation	
	nmol mg ⁻¹ protein	%
Complete	1.54	100
- α -Ketoglutarate	0.03	2
-Glycine	0.02	1
-GTP	0.02	1
-CoA	0.05	3
-NADP ⁺	0.41	27
-NAD ⁺	1.92	125
-NAD ⁺ , -NADP ⁺	0.59	38

phate, and neither GTP nor ATP markedly increased the amount of ALA formed in this reaction (Table IV).

Succinyl-CoA Synthetase Activity

Succinyl-CoA synthetase activity was present in *Euglena* extracts (Table V). Full activity required nondenatured protein, succinate, CoA, and GTP. Apparent activity was also present when ATP was used instead of GTP as the nucleoside triphosphate, but with ATP, strong dependence on added succinate and incubation at temperatures above 0°C could not be demonstrated (Table VI). Some differences in activity were measured in extracts from light- and dark-grown wild-type and nongreening mutant cells (Table VII). The activity levels were lower, but the differences between cell types were greater, in the GTP-linked enzyme assays than in the ATP-linked assays. In all cases, the measured *in vitro* succinyl-CoA synthetase activity was at least 10 times greater than ALA synthase activity.

Table IV. Influence of ATP and GTP on ALA Synthesis from Succinyl-CoA

Desalted protein extract from dark-grown wild-type *Euglena* cells was assayed for ALA formation from chemically synthesized succinyl-CoA at 43°C. Except where indicated, the incubation mixture (1 mL) contained 1 M glycerol, 75 mM glycine, 50 mM Tricine (pH 8.2), 5 mM MgCl₂, 5 mM EDTA, 5 mM levulinate, 1 mM DTT, 0.1 mM PALP, 0.1 mM succinyl-CoA, and 2.8 mg of *Euglena* protein.

Incubation Content	ALA Formation	
	nmol mg ⁻¹ protein	%
Complete	1.57	100
+1 mM GTP	1.50	96
+5 mM GTP	1.92	122
+1 mM ATP	1.10	70
+5 mM ATP	1.81	115

Table V. GTP-Linked Succinyl-CoA Synthetase Activity

Desalted protein extract from dark-grown wild-type *Euglena* cells was assayed for succinyl-CoA synthetase activity at 43°C. Except where indicated, incubation mixture (1 mL) contained 1 M glycerol, 50 mM Tricine (pH 8.0), 25 mM succinate, 10 mM MgCl₂, 5 mM GTP, 1 mM DTT, 0.2 mM CoA, 0.02 mM PALP, and 1 mg of *Euglena* protein.

Incubation Content	Succinyl-CoA Formation	
	nmol mg ⁻¹ protein	%
Complete	225	100
Complete (incubated at 0°C)	49	21
-Succinate	44	20
-CoA	5	2
-GTP	2	1

DISCUSSION

It was previously demonstrated that *Euglena* has the ability to form ALA by both the transfer RNA-dependent five-carbon pathway and by ALA synthase-catalyzed condensation of glycine with succinyl-CoA (1, 13, 22). *In vivo*, the five-carbon pathway is the sole source of Chl precursors in *Euglena* (15, 22), and this pathway is inactive in dark-grown cells (12, 22). The ALA synthesized from glycine, on the other hand, is the sole source of mitochondrial tetrapyrrole precursors in *Euglena* and may also supply other cellular tetrapyrrole end products in the dark (22).

Even though ALA synthesis from glycine has been shown to occur *in vivo* and ALA synthase activity has been detected *in vitro*, there still remained some uncertainty whether ALA synthase can function *in vivo*. The uncertainty arose from the existence of an atypical tricarboxylic acid cycle in *Euglena*. In the atypical cycle, the α -ketoglutarate dehydrogenase and succinyl-CoA synthetase steps are bypassed, and α -ketoglutarate is converted to succinate via succinate semialdehyde (Fig. 1) (8, 17, 18, 20). Because succinyl-CoA is not an intermediate in the atypical tricarboxylic acid cycle, it was possible that this compound is absent from *Euglena* cells and that ALA is formed from glycine *in vivo* via some other route.

The ability of *Euglena* to synthesize succinyl-CoA and to use this compound for ALA biosynthesis was demonstrated in three ways. First, desalted cell extracts catalyzed ALA formation in the absence of added succinyl-CoA synthetase.

Table VI. Apparent ATP-Linked Succinyl-CoA Synthetase Activity

Desalted protein extract from dark-grown wild-type *Euglena* cells was assayed for succinyl-CoA synthetase activity at 43°C using 5 mM ATP as the nucleoside triphosphate. Except where indicated, incubation mixture (1 mL) contained 1 M glycerol, 50 mM Tricine (pH 8.0), 25 mM succinate, 10 mM MgCl₂, 5 mM ATP, 1 mM DTT, 0.2 mM CoA, 0.02 mM PALP, and 1 mg of *Euglena* protein.

Incubation Content	Succinyl-CoA Formation	
	nmol mg ⁻¹ protein	%
Complete	750	100
Complete (incubated at 0°C)	693	92
-Succinate	625	83
-CoA	6	1
-ATP	6	1

Table VII. Succinyl-CoA Synthetase Activity in Extracts of Light- and Dark-Grown Wild-Type and Nongreening Mutant *Euglena* Cells

Desalted protein extract from light- and dark-grown wild-type and nongreening mutant *Euglena* cells was assayed for GTP- and ATP-linked succinyl-CoA synthetase activity at 43°C. Incubations (1 mL) contained 1 M glycerol, 50 mM Tricine (pH 8.0), 25 mM succinate, 10 mM MgCl₂, 5 mM GTP or ATP, 1 mM DTT, 0.2 mM CoA, and 0.02 mM PALP. The GTP-linked assays contained 1.2 mg of *Euglena* protein and the ATP-linked assays contained 0.27 to 0.41 mg of *Euglena* protein for which the purification step of differential (NH₄)₂SO₄ precipitation had been omitted.

Cell Extract Type	Succinyl-CoA Formation	
	GTP linked	ATP linked
	nmol mg ⁻¹ protein	
Dark-grown wild type	238	649
Light-grown wild type	120	418
Dark-grown nongreening mutant	108	522
Light-grown nongreening mutant	104	390

The reaction required CoA and a nucleoside triphosphate. Second, succinyl-CoA synthetase activity was detected in cell extracts. The activity level was more than adequate to supply substrate for ALA synthase. Third, cell extracts catalyzed glycine- and α -ketoglutarate-dependent ALA synthesis. This activity required CoA and nucleoside triphosphate. Nucleoside triphosphate was not required for ALA synthesis from glycine and chemically synthesized succinyl-CoA. The nucleoside triphosphate requirement for ALA synthesis from α -ketoglutarate indicates that the succinyl-CoA is not formed from α -ketoglutarate via the α -ketoglutarate dehydrogenase reaction because neither that reaction nor ALA synthase has a requirement for nucleoside triphosphate. The CoA requirement indicates that succinyl-CoA is a required precursor to ALA and that succinic semialdehyde cannot be converted to ALA directly. The stimulation of the reaction by NADP⁺ also supports the proposal that succinic semialdehyde must first be oxidized to succinate before succinate is converted to ALA. These results thus support the previous reports indicating that *Euglena* contains an atypical tricarboxylic acid cycle that lacks α -ketoglutarate dehydrogenase and that α -ketoglutarate is converted to succinate via an alternative route that does not involve succinyl-CoA (17, 18).

Succinyl-CoA synthetases have been classified into two groups based on the nucleoside triphosphate specificity (7, 14, 24, 25). The enzymes obtained from most animal cells, Gram-positive bacteria, and cyanobacteria (EC 6.2.1.4) are more active with GTP, whereas the enzymes from yeasts, plants, and most Gram-negative bacteria (EC 6.2.1.5) are more active with ATP. Some bacterial enzymes have been reported to have no marked activity difference with ATP and GTP (3). Enzymes of both catalytic classes can exist in "large" or "small" sizes, depending on the number of protein subunits they contain (14). ALA synthesis in *Euglena* extracts was supported approximately equally by GTP and ATP. This result indicates that both nucleoside triphosphates support sufficient succinyl-CoA synthesis to provide this ALA synthase substrate at nonlimiting concentration. Both nucleoside triphosphates also supported succinyl-CoA synthetase activ-

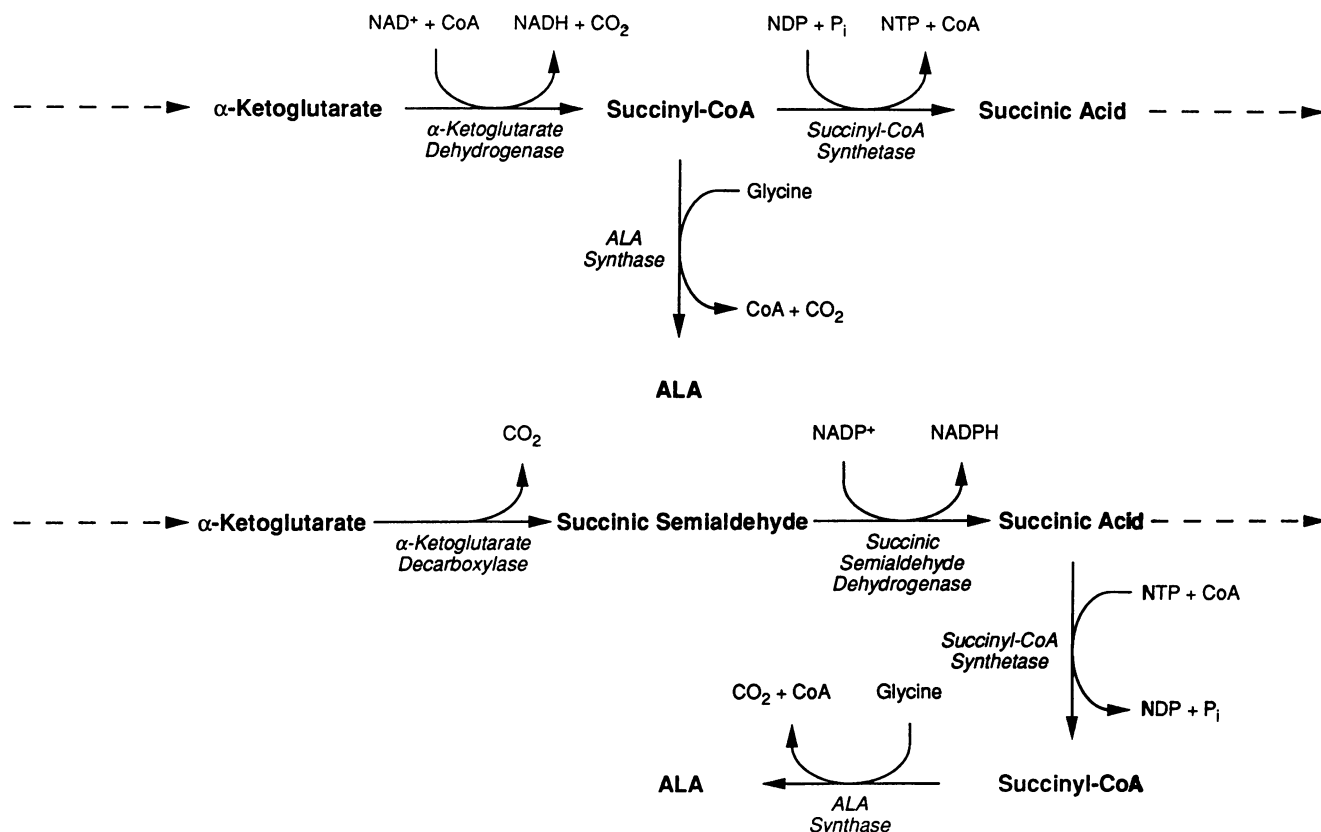


Figure 1. Reactions of the tricarboxylic acid cycle between α -ketoglutarate and succinate and ALA formation via the ALA synthase-catalyzed reaction. Depending on the species, the required nucleoside di- and triphosphates for succinyl-CoA synthetase (indicated as NDP and NTP) may be either the adenine or guanine compounds. Top, Reactions that occur in most organisms that form ALA via the ALA synthase reaction; bottom, reactions that occur in *Euglena*.

ity. The measured activity was somewhat greater in the ATP-linked enzyme assay than in the GTP-linked assay. However, because a strong dependence on added succinate or elevated temperature could not be demonstrated for the ATP-linked enzyme assay, it could not be ascertained whether some portion of the measured ATP-linked activity is nonspecific or artifactual. Therefore, we hesitate to classify the *Euglena* succinyl-CoA synthetase on the basis of its nucleoside triphosphate preference at this time.

ALA synthase activity is much higher in extracts of dark-grown wild-type cells than in light-grown cell extracts, and the extractable activity declines precipitously within the first few hours after transfer of dark-grown cells to the light (4, 6). Certain nongreening mutant *Euglena* strains appear to lack the capacity to form ALA from glutamate *in vivo* (22). In these strains, ALA synthase activity remains high in the light or dark (1). It has been proposed that ALA formation via the ALA synthase pathway in *Euglena* is regulated by the availability of ALA that is synthesized in the plastid via the five-carbon pathway (4). When the plastids are biosynthetically active, they may export heme or a heme precursor for incorporation into other cellular hemoproteins, thus lowering the requirement for mitochondrial ALA synthesis. Conversely, when the plastids are quiescent or inactive, ALA synthase may satisfy most or all cellular requirements for heme pre-

cursors. Relatively small differences were found in succinyl-CoA synthetase activity in extracts of dark- and light-grown wild-type and nongreening mutant *Euglena* cells. This result indicates that the reciprocal relationship that exists between the state of plastid development and ALA synthase activity is specific to that enzyme and is not a general pattern exhibited by mitochondrial biosynthetic enzymes.

In conclusion, the results reported here indicate that succinyl-CoA synthetase is present in *Euglena* cells. Even though the enzyme may play no role in the transformation of α -ketoglutarate to succinate in the atypical tricarboxylic acid cycle of *Euglena*, it can catalyze succinyl-CoA formation from succinate for use in the biosynthesis of ALA (Fig. 1) and possibly other products.

ACKNOWLEDGMENTS

We thank S. Rieble for helpful discussions and A.G. Smith for bringing to our attention the reports of the atypical tricarboxylic acid cycle in *Euglena*.

LITERATURE CITED

1. Beale SI, Foley T, Dzelzkalns V (1981) δ -Aminolevulinic acid synthase from *Euglena gracilis*. Proc Natl Acad Sci USA 78: 1666-1669
2. Bradford MM (1976) A rapid and sensitive method for the

- quantitation of microgram quantities of protein utilizing the principle of protein-dye binding. *Anal Biochem* **72**: 248–254
3. **Burnham BF** (1963) Purification and characterization of succinyl CoA synthetase from *Rhodospseudomonas spheroides*. *Acta Chem Scand* **17**: S123-S128
 4. **Corriveau JL, Beale SI** (1986) Influence of gabaculine on growth, chlorophyll synthesis, and δ -aminolevulinic acid synthase activity in *Euglena gracilis*. *Plant Sci* **45**: 9–17
 5. **Dzelzkalns V, Foley T, Beale SI** (1982) δ -Aminolevulinic acid synthase of *Euglena gracilis*: physical and kinetic properties. *Arch Biochem Biophys* **216**: 196–203
 6. **Foley T, Dzelzkalns V, Beale SI** (1982) δ -Aminolevulinic acid synthase of *Euglena gracilis*: regulation of activity. *Plant Physiol* **70**: 219–226
 7. **Kelly CJ, Cha S** (1977) Nucleotide specificity of succinate thiokinases from bacteria. *Arch Biochem Biophys* **178**: 208–217
 8. **Kitaoka S, Nakano Y, Miyatake K, Yokota A** (1989) Enzymes and their functional locations. In DE Buetow, ed, *The Biology of Euglena*, Vol IV. Academic Press, New York, pp 1–135
 9. **Lipmann F, Tuttle LC** (1945) A specific micromethod for the determination of acyl phosphates. *J Biol Chem* **159**: 21–28
 10. **Lombardo, ME, Araujo LS, Juknat AA, Batlle AM del C** (1988) Effect of illumination on growth, chlorophyll content and δ -aminolevulinic acid synthesis in *Euglena gracilis*. *Comp Biochem Physiol* **91B**: 279–284
 11. **Mauzerall D, Granick S** (1956) The occurrence and determination of δ -aminolevulinic acid and porphobilinogen in urine. *J Biol Chem* **219**: 435–445
 12. **Mayer SM, Beale SI** (1990) Light regulation of δ -aminolevulinic acid biosynthetic enzymes and tRNA in *Euglena gracilis*. *Plant Physiol* **94**: 1365–1375
 13. **Mayer SM, Beale SI, Weinstein JD** (1987) Enzymatic conversion of glutamate to δ -aminolevulinic acid in soluble extracts of *Euglena gracilis*. *J Biol Chem* **262**: 12541–12549
 14. **Nishimura JS** (1986) Succinyl-CoA synthetase structure-function relationships and other considerations. *Adv Enzymol Relat Areas Mol Biol* **58**: 141–172
 15. **Okazaki T, Kurumaya K, Sagae Y, Kajiwarra M** (1990) Studies on the biosynthesis of corrinoids and porphyrinoids IV. Biosynthesis of chlorophyll in *Euglena gracilis*. *Chem Pharm Bull* **38**: 3303–3307
 16. **Rieble S, Beale SI** (1988) Enzymatic transformation of glutamate to δ -aminolevulinic acid by soluble extracts of *Synechocystis* sp. 6803 and other oxygenic prokaryotes. *J Biol Chem* **263**: 8864–8871
 17. **Shigeoka S, Nakano Y** (1991) Characterization and molecular properties of 2-oxoglutarate decarboxylase from *Euglena gracilis*. *Arch Biochem Biophys* **288**: 22–28
 18. **Shigeoka S, Onishi T, Maeda K, Nakano Y, Kitaoka S** (1986) Occurrence of thiamin pyrophosphate-dependent 2-oxoglutarate decarboxylase in mitochondria of *Euglena gracilis*. *FEBS Lett* **195**: 43–47
 19. **Simon EJ, Shemin D** (1953) The preparation of S-succinyl coenzyme A. *J Am Chem Soc* **75**: 2520
 20. **Tokunaga M, Nakano Y, Kitaoka S** (1976) Separation and properties of the NAD-linked and NADP-linked isozymes of succinic semialdehyde dehydrogenase in *Euglena gracilis* Z. *Biochim Biophys Acta* **429**: 55–62
 21. **Urata G, Granick S** (1963) Biosynthesis of α -aminoketones and the metabolism of aminoacetone. *J Biol Chem* **238**: 811–820
 22. **Weinstein JD, Beale SI** (1983) Separate physiological roles and subcellular compartments for two tetrapyrrole biosynthetic pathways in *Euglena gracilis*. *J Biol Chem* **258**: 6799–6807
 23. **Weinstein JD, Mayer SM, Beale SI** (1986) Stimulation of δ -aminolevulinic acid formation in algal extracts by heterologous RNA. *Plant Physiol* **82**: 1096–1101
 24. **Weitzman PDJ, Jaskowska-Hodges H** (1982) Patterns of nucleotide utilisation in bacterial succinate thiokinases. *FEBS Lett* **143**: 237–240
 25. **Weitzman PDJ, Kinghorn HA** (1980) Succinate thiokinase from cyanobacteria. *FEBS Lett* **114**: 225–227