Positive Charges Determine the Topology and Functionality of the Transmembrane Domain in the Chloroplastic Outer Envelope Protein Toc34

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Abstract. The chloroplastic outer envelope protein Toc34 is inserted into the membrane by a COOH-terminal membrane anchor domain in the orientation Ncyto-Cin. The insertion is independent of ATP and a cleavable transit sequence. The cytosolic domain of Toc34 does not influence the insertion process and can be replaced by a different hydrophilic reporter peptide. Inversion of the COOH-terminal, 45-residue segment, including the membrane anchor domain (Toc34Cinv), resulted in an inverted topology of the protein, i.e., Nin-Ccyto. A mutual exchange of the charged amino acid residues NH2- and COOH-proximal of the hydrophobic α-helix indicates that a double-positive charge at the cytosolic side of the transmembrane α-helix is the sole determinant for its topology. When the inverted COOH-terminal segment was fused to the chloroplastic precursor of the ribulose-1,5-bisphosphate carboxylase small subunit (pS34Cinv), it engaged the transit sequence–dependent import pathway. The inverted peptide domain of Toc34 functions as a stop transfer signal and is released out of the outer envelope protein translocation machinery into the lipid phase. Simultaneously, the NH2-terminal part of the hybrid precursor remained engaged in the inner envelope protein translocon, which could be reversed by the removal of ATP, demonstrating that only an energy-dependent force but no further ionic interactions kept the precursor in the import machinery.

The majority of chloroplast proteins are encoded for in the nucleus, synthesized in the cytosol, and post-translationally imported into the organelle. The standard import route is facilitated by two protein translocation machineries located in the chloroplastic outer and inner envelope membranes, which act jointly during translocation (Schnell and Blobel, 1993; Alessen et al., 1994). Proteins destined to be translocated into chloroplasts carry NH2-terminal transit sequences, which are both necessary and sufficient for targeting and import. The transit sequence recognition is mediated by a protease-sensitive receptor polypeptide located in the outer envelope membrane (Cline et al., 1985). A low ATP concentration, between 5–50 μM, is required for binding of the precursor protein to the outer envelope translocon in a step not well characterized yet, while high concentrations, between 50–1,000 μM, are necessary for complete translocation into the chloroplasts (Flügge and Hinz, 1986; Schindler et al., 1987; Olsen et al., 1989).

In contrast, proteins destined for the chloroplastic outer envelope are generally synthesized without an NH2-terminal cleavable transit sequence but are targeted by internal information (Salomon et al., 1990; Li et al., 1991; Ko et al., 1992; Fischer et al., 1994; Kessler et al., 1994; Seedorf et al., 1995; Li and Chen, 1996; Chen and Schnell, 1997), except for the 86- and 75-kD subunits of the translocon at the chloroplastic outer envelope (Toc86,1 Hirsch et al., 1994; Kessler et al., 1994; Toc75, Tranel et al., 1995; Tranel and Keegstra, 1996). The Toc75 precursor uses the general import machinery during the initial phase of the import. It is processed in the stroma to an intermediate-size form while the majority of the mature protein remains in the chloroplastic outer membrane by an unknown stop transfer mechanism. In a second phase of translocation, the protein is inserted into the membrane and terminally processed in a poorly understood way. The transit sequence–independent targeting and insertion seem to require neither protease-sensitive polypeptide components of the outer envelope membrane nor the hydrolysis of ATP. Current

1. Abbreviations used in this paper: chl, chlorophyll; inv, inverse; mS, small subunit of ribulose-1,5-bisphosphate carboxylase; pS, precursor form; Tic, translocon at the inner envelope membrane of chloroplasts; TIM, translocation intermediates; Toc, translocon at the outer envelope membrane of chloroplasts.
evidence suggests that none of the protein translocon components of the chloroplastic outer envelope is involved in the presequence-independent insertion pathway (Salomon et al., 1990; Li et al., 1991; Ko et al., 1992; Fischer et al., 1994; Seedorf et al., 1995; Li and Chen, 1996; Chen and Schnell, 1997; for review see Soll et al., 1992). This insertion mechanism is different from the presequence-independent integration of proteins into the mitochondrial outer membrane, which uses subunits of the TOM machinery (for review see Lill and Neupert, 1996).

An issue that remains to be solved is how these membrane proteins achieve their correct topology. (a) Does the specific lipid composition influence targeting and topology of the chloroplastic outer envelope proteins, and (b) what is the role of the membrane spanning domains of these envelope polypeptides? Or, (c) do soluble segments of the protein influence its insertion and topology? Toc34 has a well-established topology, N\text{cyto}-C\text{cyt}, with only one transmembrane segment. The \(\alpha\)-helical hydrophobic membrane anchor is located close to the COOH terminus of the protein, which leaves only 3 kD of the polypeptide exposed to the intermembrane space. The major portion of the protein, including its GTP-binding domain, is cytosolic. Insertion of Toc34 translation product into the chloroplastic outer envelope requires the presence of the hydrophobic transmembrane segment (Seedorf et al., 1995; Chen and Schnell, 1997; Li and Chen, 1997). This insertion process renders Toc34 resistant to extraction at pH 11.5. Productive folding or assembly, which can be followed by the appearance of an 8-kD protease-protected fragment, seems to be stimulated by the presence of nucleoside-triphosphates and a protease-sensitive envelope membrane component (Seedorf et al., 1995; Chen and Schnell, 1997). Resistance to alkaline extraction together with the appearance of the 8-kD proteolytic fragment can be used to follow the productive insertion process of Toc34.

In this study, we demonstrate that a double-positive charge at the cytosolic side of the \(\alpha\)-helical membrane anchor dictates the topology of Toc34. Furthermore, this domain acts as a functional stop transfer signal for the outer envelope protein translocon. The signal anchor sequence is released laterally from the translocon into the lipid phase of the membrane.

**Materials and Methods**

**Chloroplastic Protein Import Assays**

Chloroplasts were isolated from pea leaves of 10–12-d-old plants by standard procedures and purified further on silica-sol gradients (Waegemann and Soll, 1995). Standard import assays contained chloroplasts equivalent to 30 \(\mu\)g chl in 100 \(\mu\)l import buffer (330 mM sorbitol, 50 mM Hepes-KOH, pH 7.6, 3 mM MgCl\(_2\), 10 mM methionine, 10 mM cysteine, 20 mM potassium glutonate, 10 mM NaHCO\(_3\), 2% [wt/vol] BSA) and 1–10% of reticulocyte lysate in vitro–synthesized \(^{35}\)S-labeled proteins. Translocation reactions were initiated by the addition of organelles and allowed to continue for 10 min at ambient temperature. Chloroplasts were recovered from the import reaction by centrifugation (5,000 \(\times\) g, 5 min, 4°C) through a Percoll cushion (40% [vol/vol] Percoll in 330 mM sorbitol, 50 mM Hepes-KOH, pH 7.6), washed once in Hepes-sorbitol (Waegemann and Soll, 1991), and used for further treatments.

Chloroplasts were treated with thermolysin before a translocation experiment with 750 \(\mu\)g protease/mg chl for 30 min on ice in 330 mM sorbitol, 50 mM Hepes-KOH, pH 7.6, 3 mM MgCl\(_2\), and 0.5 mM CaCl\(_2\). The reaction was terminated by the addition of 10 mM EDTA. Intact chloroplasts were recovered on silica-sol gradients and washed twice as described above. When chloroplasts were treated with thermolysin after an import experiment, organelles were incubated at a final concentration of 100 \(\mu\)g protease/mg chl for 15 min at 4°C. The reaction was stopped by the addition of 10 mM EDTA, and the chloroplasts were recovered by centrifugation and washed once (Joyard et al., 1983; Clène et al., 1984).

When indicated, chloroplasts were treated with trypsin (Marshall et al., 1990; Lübeck et al., 1996). Chloroplasts equivalent to 200 \(\mu\)g chl were incubated with 200 \(\mu\)g trypsin (10,700 Na-benzoyl-l-arginine ethyl ester U/mg from bovine pancreas) for 60 min at 25°C in 50 mM Hepes-KOH, pH 8.0, 0.1 mM CaCl\(_2\), and various sorbitol concentrations (200 mM = hypertonic; 330 mM = isotonic; 600 mM = hypertonic conditions). The reaction was stopped by the addition of 1 mM PMSF and a fivefold molar excess of soybean trypsin inhibitor. Intact chloroplasts were recovered by centrifugation through a Percoll cushion and washed twice under isotonic conditions as described above. PMSF was present at all washing steps. Chlorophyll concentration was determined, and equal amounts of organelles were loaded onto the SDS-PAGE on a chl basis.

After hypotonic lysis in 10 mM Hepes-KOH, pH 7.6, chloroplasts were separated into a soluble and a total membrane fraction by centrifugation for 10 min at 165,000 \(\times\) g. Soluble proteins were precipitated by 10% TCA. In some cases, lysed organelles were treated with 0.1 M Na\(_2\)CO\(_3\), pH 11.5, for 10 min on ice. Insoluble material was recovered by centrifugation before and, soluble proteins were precipitated by 10% TCA. Import products were analyzed by SDS-PAGE (Laemmli, 1970) followed by fluorography (Bonner and Laskey, 1974).

**Construction of cDNAs Coding for Toc34 Hybrid Proteins and Synthesis of Labeled Proteins**

cDNAs coding for the different hybrid proteins, including single amino acid exchanges, were constructed by recombinant PCR (Higuchi, 1990). Toc34Cinv was constructed using two extra-long (105-bp) primers, which coded for the inverted sequence of the Toc34 COOH terminus (see Fig. 1). Primers were cloned into a vector suitable for in vitro transcription and controlled by DNA sequencing (Sanger et al., 1977). The original cDNAs coding for Toc34 and the precursor form (pS) have been described (Klein and Salvucci, 1992; Seedorf et al., 1995). In vitro transcription was done using either T7 or SP6 RNA-polymerase, as outlined before (Salomon et al., 1990). Proteins were synthesized in a reticulocyte lysate system in the presence of \(^{14}\)C]methionine and \(^{35}\)S]cysteine (1,175 Ci/mmol) for 1.5 h at 30°C. Overexpression of unlabeled or radioactively labeled proteins was done in Escherichia coli BL21 (DE3) cells using the pET vector system (Novagen Corp., Madison, WI) (Waegemann and Soll, 1995). Proteins were isolated from inclusion bodies and solubilized in 8 M urea. The final urea concentration in the import assays never exceeded 80 mM.

**Chloroplastic ATP Levels**

Intact chloroplasts were incubated in import buffer in the presence of 2 mM ATP and 2 \(\mu\)Ci [\(\gamma\)–\(^{32}\)P]ATP for 10 min at 25°C. An aliquot was removed, and chloroplasts were separated from the aqueous medium by centrifugation (15 s, 9,000 rpm) through a silicon oil layer (Wirtz et al., 1980). Residual chloroplasts were recovered by centrifugation and resuspended in import buffer in the absence of ATP. Aliquots were removed from the organelle suspension at different time intervals and recovered by centrifugation through silicon oil. Chloroplasts were extracted with 80% dimethylketon. Nucleotides were separated by thin layer chromatography on glass plates coated with polyethyleneimine cellulose using 0.5 M KH\(_2\)PO\(_4\) as solvent. ATP-containing spots were localized in UV light and scraped off the plates, and radioactivity was determined after addition of scintillation cocktail in a scintillation counter.

**Results**

Toc34 has an established topology, i.e., N\text{cyto}-C\text{cyt} with a single hydrophobic \(\alpha\)-helical region at the COOH terminus acting as a membrane anchor domain. Our aim was to study the role of the membrane anchor region in the topology of Toc34 and its influence on the protein import pathway. Therefore, the hybrid protein Toc34Cinv was constructed containing 45 amino acids of the COOH terminus,
including the membrane anchor domain, but in an inverted sequence (for details see Fig. 1). Accurate targeting and insertion of Toc34 into the chloroplastic outer envelope can be proven by a combination of two experimental approaches: (a) Toc34 is largely resistant to extraction at pH 11.5 in situ (Seedorf et al., 1995); and (b) Toc34 is sensitive to thermolysin in situ, except for an 8-kD COOH-terminal fragment (Seedorf et al., 1995). When Toc34Cinv translation product was incubated with intact chloroplasts, the protein integrated into the outer envelope membrane in a largely protease-resistant form (Fig. 2 A, bottom, lane 2), in contrast to the wild-type Toc34 (Fig. 2 A, top, lane 2). Protease resistance of inserted hybrid protein is compatible with the sequence of Toc34Cinv because the protein contains only three amino acids COOH-terminal of the inverted membrane anchor (see Fig. 1). The appearance of proteolytic fragments, which are formed from a subpopulation of Toc34Cinv, was most likely due to an incomplete transfer of the hydrophilic domain across the outer envelope membrane. Toc34Cinv inserted into the membrane in a way that renders it largely resistant to extraction at pH 11.5, i.e., it behaves as an integral membrane protein like the wild-type Toc34 (Fig. 2 A, lanes 3 and 4). The proteolytic fragments generated from inserted Toc34Cinv were also resistant to alkaline extraction, demonstrating that the membrane anchor region had integrated into the lipid bilayer even in those cases where translocation was not complete. Insertion of Toc34 translation products into chloroplasts yielded the expected results (Seedorf et al., 1995; Chen and Schnell, 1997), i.e., the integrated protein was sensitive to thermolysin but resistant to alkaline extraction (Fig. 2 A, top, lanes 3–6).

Next we wanted to know if the hydrophilic NH₂-terminal part of Toc34 influences the insertion reaction. It was replaced by the mature form of the Rubisco small subunit (mS), which is a soluble stroma-localized protein. Both hybrid proteins, mS34C and mS34Cinv, inserted indistinguishably in the chloroplastic outer envelope to the analogic “wild-type” Toc34’s (Fig. 2, A and B), i.e., inserted mS34C and mS34Cinv were resistant to extraction at pH 11.5 (Fig. 2 A, lanes 9 and 10), and mS34C was susceptible to a thermolysin treatment while mS34Cinv proved to be largely resistant (Fig. 2 A, lanes 7 and 8). The proteolytic fragments of inserted mS34Cinv are most likely due to incomplete translocation of the NH₂-terminal polypeptide chain across the outer envelope membranes. Solubilization of chloroplast membranes by detergent before protease treatment resulted in complete proteolysis (not shown).

To obtain further experimental evidence for the putative topology of the different Toc34 proteins, they were synthesized in a reticulocyte lysate system in the presence of [³H]leucine. The 8-kD proteolysis-resistant fragment (see above) can only be detected in experiments using [³H]leucine-labeled Toc34 because the COOH terminus does not contain any methionine or cysteine residues. The appearance of the 8-kD fragment either from [³H]Toc34

Figure 1. Peptide domain combination of various hybrid proteins used in this study and their nomenclature. The membrane anchor (MA) is underlined. Numbers indicate amino acid positions of the Toc34 sequence from pea (Seedorf et al., 1995).

Figure 2. The COOH-terminal segment of Toc34 is the sole determinant for the protein topology. (A) ³⁵S-labeled Toc34, Toc34Cinv, mS34C, and mS34Cinv translation product (TL) was used in standard translocation experiments (2 mM ATP, 25°C, 10 min) with purified pea chloroplasts (equivalent to 30 μg chl). After completion of the translocation experiment, organelles were either not treated or treated with the protease thermolysin as indicated. Chloroplasts were lysed in hypertonic buffer (lanes 3–6) and separated into a soluble (S) and insoluble (M) fraction by centrifugation after extraction at pH 11.5 with 0.1 M Na₂CO₃. (B) Toc34, Toc34Cinv, mS34C, and mS were synthesized in a reticulocyte lysate system in the presence of [³H]leucine and used in a parallel translocation experiment under identical conditions as described for A. SDS-PAGE gels were loaded with equal amounts of organelles on a chl basis. A 12.5% (wt/vol) acrylamide-containing separation gel was used in A, while a 15% gel was used in B. Translocation product (TL) is shown as internal standard, 10% of which was added to a standard translocation reaction. Fluorograms are shown.
or [3H]mS34C translation product shows that both poly-
peptides obtained a topology N\textsubscript{cyto}-C\textsubscript{cyto}, i.e., indistinguish-
able to Toc34 in situ (Fig. 2 B, lanes 1 and 2, and 5 and 6).
Thermolysin treatment of inserted [3H]Toc34Cinv yielded
no 8-kD fragment, but did yield a fragmentation pattern
similar to the [35S]-labeled Toc34Cinv (Fig. 2 B, lanes 3 and
4). mS does not interact with chloroplasts to a detectable
amount (Fig. 2 B, lane 7). From these data, we conclude
that the Toc34 COOH-terminal 45 amino acids seem to be
the only determinant for chloroplast targeting, and they
simultaneously dictate the topology of the inserted pro-
tein. The inverted COOH terminus of Toc34 can function
as a bona fide membrane anchor in the chloroplastic outer
membrane but simultaneously inverts the topology of the
protein into N\textsubscript{in}-C\textsubscript{cyto}.

As demonstrated above, the COOH-terminal region of
Toc34 contains an outer envelope targeting domain re-
gardless of its orientation. In light of this, we asked if the
stroma-targeting envelope transfer signal present in the
transit sequence of pS is able to override the membrane
anchoring information present in the COOH terminus of
Toc34. We therefore compared the import and binding
of preSSU with its derived hybrid proteins pS34C and
pS34Cinv. Under conditions that allow import to occur at
optimal rates (25°C and 2 mM ATP), pS is efficiently
translocated into the chloroplasts and processed to the
mature form (Fig. 3 A, lanes 1 and 2). The mature form mS
is recovered as a protease-protected soluble protein (Fig. 3
A, lanes 3–6). The ratio of radioactivity recovered in pS
and processed mature mS was 1:19, as determined by
laser densitometry of exposed x-ray films (n = 5). pS34C is
imported into intact chloroplasts with a yield similar to
preSSU. It is processed like the wild-type protein to yield
mS34C (Fig. 3 B, lanes 1 and 2). The ratio of chloroplast-
bound precursor and mature mS34C was 2:8 (n = 5).

Upon subfractionation of the chloroplasts, an equal distri-
bution of mS34C is observed between the total membrane
and the soluble protein fraction (Fig. 3 B, lanes 3 and 4).
However, mS34C only adheres to the membranes since it
is completely extractable at pH 11.5 (Fig. 3 B, lanes 5 and
6). In comparison to preSSU, the binding of pS34C is only
slightly increased, indicating that most but not all of the
hybrid precursor might enter the standard import route.
Translocation still takes place when pS34Cinv translation
product was incubated with intact chloroplasts under stan-
dard import conditions. However, the processing resulted
primarily in a product of intermediate size (iS34Cinv) and
only in a very small amount of mature mS34Cinv (Fig. 3 C,
lanes 1 and 2). The ratio between pS34Cinv, iS34Cinv, and
mS34Cinv was 6:3:1 (n = 5). iS34Cinv was completely re-
covered in the total membrane fraction (Fig. 3 C, lanes 3
and 4). Upon alkaline extraction, 80% of iS34Cinv was re-
covered in the pellet fraction while 20% could be ex-
tacted (Fig. 3 C, lanes 5 and 6). These latter results indi-
cated that most of iS34Cinv had inserted into the lipid
bilayer of the outer or inner envelope while a smaller por-
tion of iS34Cinv remained in a proteinaceous environ-
ment, e.g., the protein translocation pore. When the outer
and inner envelope membranes were separated from chlo-
oplasts after pS34Cinv translocation, iS34Cinv was re-
covered in the outer envelope (Fig. 3 E). The translocation
efficiency of pS34Cinv is less than for pS and pS34C because
more pS34Cinv remained bound to the chloroplast surface
in a protease-accessible manner (Fig. 3 B, lanes 1 and 2).
On the other hand, insertion of mS34Cinv translation
product into the outer envelope attains a largely protease-
protected topology (Fig. 2, lanes 5). It is processed like the
wild-type protein to yield mS34C (Fig. 3 B, lanes 1 and 2).

The general import pathway into chloroplasts by the
Toc and Tic machinery requires protease-sensitive recep-
tor components at the chloroplastic surface, which recog-
nize the NH\textsubscript{2}-terminal transit sequences. In contrast, the
insertion of chloroplastic outer envelope proteins, which
do not contain a presequence, is independent of a pro-
tease-sensitive surface component. To investigate further
if and to what extent pS34C and pS34Cinv used the stan-
dard import pathway, chloroplast surface–exposed precu-
sor receptors were removed by treatment with the pro-

Figure 3. The inverted
COOH-terminal segment of
Toc34 functions as a stop
transfer domain at the chlo-
roplast outer envelope.
(A–D) Indicated [35S]-labeled,
in vitro–translated (A–C) or
overexpressed (D) proteins
were incubated with intact
pea chloroplasts under stan-
dard import conditions. Af-
after completion of the import
reaction, chloroplasts were
either not treated (lane 1) or
treated (lane 2–6) with the
protease thermolysin. Or-
ganelles were fractionated into
a total membrane (M, lane 3)
or soluble protein (S, lane 4)
fraction. Chloroplasts were
incubated with 0.1 M
Na\textsubscript{2}CO\textsubscript{3} before separation
into an alkaline-insoluble (M, lane 5) or -soluble (S, lane 6)
fraction. (E) Chloroplastic outer and inner envelope membranes
were fractionated after a standard import reaction as described
by Keegstra and Youssif (1986). Chloroplasts were treated with
thermolysin after import. Fluorograms are shown.

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tease thermolysin. The binding of pS to protease-shaved chloroplasts was reduced to 10% in comparison to untreated organelles (Fig. 4, C, lanes 1 and 3; and D). In contrast, the amount of binding of pS34C or pS34Cinv was not significantly influenced by a protease pretreatment, indicating that both proteins inserted largely via the COOH-terminal membrane anchor region under conditions that discourage the Toc pathway. Under import conditions, (2 mM ATP and 25°C), thermolysin-shaved chloroplasts were also blocked in complete translocation of pS (Fig. 4 C, lanes 7–10). The residual import yield varied between 10 and 15% (n = 6; Fig. 4 D) for protease-pretreated chloroplasts. In parallel experiments, the efficiencies of pS34C import into thermolysin-shaved chloroplasts as measured by the appearance of processed mS34C dropped to 15–20% of control imports (Fig. 4, A, lanes 7–10; and D). The import yield was even less for the translocation of pS34Cinv into protease-pretreated chloroplasts (Fig. 4 B, lanes 7–10). Only 5% iS34Cinv was still detected (Fig. 4 D). The precursor proteins pS34C and pS34Cinv, which were bound to protease-shaved chloroplasts, were resistant to alkaline extraction, indicating that they inserted by the COOH terminus into the outer envelope membrane.

To distinguish in vitro between presequence-dependent and -independent interactions with the organellar surface, i.e., via the COOH-terminal α-helix, chloroplasts were incubated with pS, pS34C, and pS34Cinv, respectively, under binding conditions (50 μM ATP and 4°C) for 10 min, followed by a chase period under translocation conditions (2 mM ATP and 25°C). pS bound to the chloroplast surface and translocation intermediates (TIM 3/4), which are well characterized for the pS import pathway (Waegeemann and Soll, 1991, 1996), were detected upon thermolysin treatment (Fig. 4 C, lanes 1 and 2, and 13 and 14). The bound pS could be chased into the chloroplasts upon the addition of ATP and raising the temperature to 25°C, demonstrating that the binding state was productive (Fig. 4 C, lane 15). pS34C and pS34Cinv also bound to the chloroplast surface with a yield similar to pS (Fig. 4, A and B, lanes 1 and 13). Upon thermolysin treatment, the translocation intermediates (TIM 3/4) could be detected (Fig. 4, B and C, lanes 2 and 14). However, chloroplast-bound hybrid proteins did not translocate into chloroplasts to any significant extent upon establishing import conditions (Fig. 4, A and B, lane 15), probably because they had also inserted during the course of the incubation period into the chloroplast outer envelope with the COOH-terminal α-helical region. This is supported by the observation that the surface-bound pS34C and pS34Cinv were resistant to alkaline extraction (not shown). These results indicate that conditions that do not allow import, i.e., low ATP and low temperature, favor the insertion of Toc34 via the α-helical membrane anchor region.

From the data presented in Fig. 4, we conclude that pS34C and pS34Cinv use preferentially the Toc and Tic complex-dependent pathway under experimental conditions that favor this pathway (2 mM ATP, 25°C) and functional receptor polypeptides. Whenever we manipulate the optimal requirements by lowering the temperature or the ATP concentration or by removing precursor recognition sites, the membrane insertion route via the COOH-terminal membrane anchor becomes more prominent.

pS34Cinv is imported into chloroplasts at 2 mM ATP and processed to an intermediate-size form (see Fig. 4 B), most likely by the stromal-processing protease. Therefore, it should also engage the Tic complex during this process. Furthermore, iS34Cinv behaves as an integral membrane protein that cofractionates with the chloroplastic outer envelope (Fig. 3 E) when the inner and outer membrane are separated by shearing forces (Keegstra and Youssif, 1986).
To test if iS34Cinv is in contact with the Tic complex, we compared the sensitivity of the imported and processed forms of pS, pS34C, and pS34Cinv, to thermolysin and trypsin, respectively. Under strictly controlled conditions, thermolysin only shaves proteins off the chloroplastic surface, while trypsin penetrates the outer envelope membrane and gains access to the intermembrane space and to proteins exposed on the surface of the inner membrane (Marshall et al., 1990; Lübeck et al., 1996; see also Fig. 5 C). mS and mS34C were resistant to both thermolysin and trypsin (Fig. 5 A, lanes 1–6), indicating that they had reached the stroma. In contrast, iS34Cinv was resistant to thermolysin but sensitive to trypsin (Fig. 5 A, lanes 7–9). A proteolytic fragment was generated by trypsin (iSΔC; Fig. 5 A, lane 9), which was recovered in the total soluble protein fraction of chloroplasts (Fig. 5 B, lane 4). These data indicate that trypsin removed the membrane anchor of iS34Cinv, so that translocation into the stroma could proceed through the inner envelope translocon. The predicted localization of iS34Cinv, i.e., anchored in the outer envelope membrane while simultaneously engaging the Tic complex, was further established by carrying out the trypsin treatment under different osmotic conditions. The rationale behind this approach is that at hypertonic conditions (0.6 M sorbitol), a retraction of the inner envelope membranes from the outer and a widening of the intermembrane space occurs because H₂O is extruded from the stroma into the surrounding medium (Block et al., 1983; see Fig. 5 C). Under these conditions, iS34Cinv should become trypsin sensitive. On the other hand, isotonic or slightly hypotonic conditions do not result in a membrane rearrangement, and trypsin treatment should result in the appearance of iSΔC. This was found to be the case (Fig. 5 B). Chloroplast-inserted iS34Cinv was completely degraded by trypsin treatment under hypertonic conditions (Fig. 5 B, lanes 9 and 10). At isotonic and hypotonic conditions, iSΔC remained protease protected (Fig. 5 B, lanes 3 and 4, and 7 and 8). iSΔC is not protease resistant per se, as demonstrated by treating the soluble iSΔC with trypsin (Fig. 5 B, lane 6), which results in complete proteolysis. We conclude that the inverted COOH-terminal segment of Toc34 can serve as a stop transfer signal for the Tic machinery. The stop transfer signal is recognized by the outer envelope translocon and laterally released into the plane of the membrane. Simultaneously, the iS part of the protein continuously interacts with the inner envelope translocon.

The latter prediction was analyzed using the following experimental procedures. Intact chloroplasts were preincubated either in the absence or presence of E. coli expressed pS (Fig. 6 A), pS34Cinv (Fig. 6 B), or Toc34 (Fig. 6 C) under import conditions for 5 min. Organelles were recovered by centrifugation and washed once in import buffer in the presence or absence of 2 mM ATP. A second round of protein translocation was started by the addition of 35S-labeled pS reticulocyte lysate translation product. Import of 35S-labeled pS was only slightly diminished when chloroplasts were preincubated in the presence of overexpressed pS, in comparison to a mock preincubation in the absence of overexpressed pS (Fig. 6 A, lanes 1 and 2, and 5 and 6). When 35S-labeled pS reticulocyte translocation was added simultaneously with E. coli expressed unlabeled pS, binding and translocation of 35S-labeled pS was almost completely abolished (Fig. 6 A, lanes 3 and 4). This demonstrates that the heterologously expressed precursor could compete with the reticulocyte lysate–synthesized precursor. In contrast, 35S-labeled pS import into chloroplasts that were preincubated with pS34Cinv was very strongly inhibited (Fig. 6 B, lanes 1 and 6), independent of the presence of pS34Cinv during the second round of translocation. Binding of pS to the chloroplast surface was not influenced significantly (Fig. 6 B, lane 5), indicating that inhibition occurred in a later step of the translocation.
reaction, e.g., at the level of the Tic complex. This is corroborated by the observation (Fig. 6 B, lanes 7 and 8) that the inhibitory effect of pS34Cinv preincubation was completely reverted when chloroplasts were reisolated and washed in the absence of ATP before the second round of translocation. The ATP concentration in chloroplasts drops to below 10 μM ATP (Fig. 6 D) during the reisolation and washing procedure. This low level of ATP is not sufficient to support translocation (Olsen et al., 1989). We conclude that an ATP-dependent pulling force in the stroma, e.g., exerted by hsp70 or hsp100, looses its grip on the partly translocated protein, upon which the precursor retracts from the inner envelope translocon. The Tic complex then becomes available for new rounds of translocation.

Heterologously expressed pS34Cinv competes effectively with 35S-labeled pS reticulocyte lysate translation product when added simultaneously to chloroplasts (Fig. 6 B, lanes 3 and 4). This further shows that pS34Cinv enters the standard import pathway. Chemical amounts of Toc34 can neither compete for pS nor for pS34Cinv import (Fig. 6 C), corroborating our earlier notion that under optimal import conditions (2 mM ATP, 25°C), the presequence-dependent standard import route is preferred to the membrane insertion pathway.

Our data so far indicated two findings: (a) Inversion of the COOH-terminal 45 amino acids of Toc34 causes a re-orientation of the membrane topology; and (b) the inverted COOH-terminal segment functions as a stop transfer signal for the Tic machinery. However, it remained to be established if the orientation determinants reside in the transmembrane or in the flanking sequences. As can be seen in Fig. 7, two positively charged amino acids are placed in direct vicinity (position-1 and -3) of the cytosolic side of the Toc34 transmembrane segment, while in the case that two positively charged amino acids are closely spaced in the luminal segment of Toc34, they are directly accompanied by a negatively charged residue. This double-positive charge might represent a determinant for Toc34 orientation. Therefore, a mutual charge exchange was performed (see Fig. 7 A), which resulted in a Toc34∧ and Toc34Cinv∧, corroborating our earlier notion that under optimal import conditions (2 mM ATP, 25°C), the presequence-dependent standard import route is preferred to the membrane insertion pathway.

In contrast to wild-type Toc34, Toc34∧ now inserts into the chloroplastic outer envelope in a way that renders...
As shown above (Fig. 3), pS34C can be imported into chloroplasts, where it is processed and recovered as soluble protein in the stroma. When we tested the binding and import properties of pS34C<sub>C<sub>++</sub></sub>, we found that it was no longer able to translocate fully into chloroplasts (Fig. 7 C, lanes 5 and 6). Instead, under import condition it was processed to iS34C<sub>C<sub>++</sub></sub> and recovered in the membrane fraction, resistant to extraction at pH 11.5 (Fig. 7 C, lanes 7 and 8). Under binding conditions, the translocation intermediates TIM 3/4 indicated that pS34C<sub>C<sub>++</sub></sub> had entered the standard import pathway (Fig. 7 C, lanes 1 and 2). We conclude that charge distribution around the transmembrane segment represents a stronger sorting signal than orientation and hydrophobicity of the membrane domain.

**Discussion**

Toc34 is anchored in the chloroplastic outer envelope by a single transmembrane hydrophobic membrane anchor in an orientation N<sub>Cyto</sub>-C<sub>in</sub>. If we compare the region of 28 amino acids NH<sub>2</sub>-terminal preceding and COOH-terminal following the membrane anchor, a charge difference of 4:7 positive charges can be observed. Hence, it is not obvious that posttranslational insertion of Toc34 obeys the positive inside rule (von Heijne and Gavel, 1988; von Heijne, 1989). Charge differences as sole determinant for the orientation of a membrane anchor region is difficult to reconcile because it would most likely require an intact electrochemical potential across the chloroplastic outer envelope (Andersson and von Heijne, 1994; Rapoport et al., 1996). At present, the evidence suggests that the outer envelope lacks a significant electrochemical potential (Joyard et al., 1991). However, the specific lipid composition of the chloroplastic outer envelope results in a strongly negatively charged membrane surface (Joyard et al., 1991). Positively charged amino acids selectively positioned to a transmembrane segment might be prominent determinants for the membrane orientation of a polypeptide. Our results establish that two positive charges in close proximity to the transmembrane segment dictate the topology of Toc34. The interconversion of the positive charge “motif” (Fig. 7) NH<sub>2</sub>- and COOH-proximal to the transmembrane segment resulted in the interconversion of the Toc34 topology. From these observations we conclude that although the hydrophobic α-helix of Toc34 is necessary for its insertion (Seedorf et al., 1995), positively charged amino acids flanking this region are the major determinants for its orientation. Characterization of the insertion behavior of other single membrane spanning outer envelope proteins will be necessary to validate and generalize this conclusion.

The general protein import machinery of the chloroplastic outer envelope can translocate proteins containing integral membrane components of either the inner envelope or the thylakoid membrane. Consequently, amino acid sequences with the potential to form hydrophobic α-helices pass freely through the outer envelope translocon. The 2-oxoglutarate-malate translocator, a 12-helix-motif transporter of the inner envelope, uses the general import machinery as well as the three α-helices containing thylakoid localized light harvesting chlorophyll a/b binding protein (Hobe et al., 1994; Weber et al., 1995). The presence of a

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**Figure 7.** Positive charge distribution determines the functionality of the COOH-terminal segment of Toc34. (A) Charged amino acids (bold) directly proximal to the α-helical membrane anchor (MA, underlined) were mutually exchanged. (B) Intact chloroplasts were incubated with Toc34 (lanes 1 and 2), Toc34<sub>C<sub>++</sub></sub> (lanes 3 and 4), Toc34Cinv (lanes 5 and 6), and Toc34Cinv<sub>C<sub>++</sub></sub> (lanes 7 and 8) translation product for 10 min under standard conditions. Chloroplasts were either not treated or treated with the protease thermolysin after completion of the insertion reaction (Thl post, as indicated). (C) pS34C<sub>C<sub>++</sub></sub> translation product was incubated with intact chloroplasts under binding (lanes 1–4) or import (lanes 5–8) conditions for 10 min as outlined on top of the figure. Chloroplasts were treated subsequently with thermolysin as outlined before. Reaction products were separated by extraction at pH 11.5 into a membrane (M, lanes 3 and 4) and a soluble (S, lanes 7 and 8) protein fraction. The position of the translocation intermediates (Tim 3/4) is indicated. Fluorograms are shown in B and C.
stop transfer signal functional at the outer envelope protein translocon would obviously be deleterious to organelle biogenesis. Chloroplast preproteins localized in the thylakoid lumen contain a bipartite transit sequence. The NH₂-proximal part functions as an envelope translocation domain, while the COOH-proximal part functions as thylakoid transfer domain and dictates which translocation route it follows, i.e., a Sec-dependent or ΔpH-dependent route (for review see Robinson and Mant, 1997). Positive charges placed around a hydrophobic domain determines the translocation pathway to be used. The hydrophobic domain in thylakoid lumen presequences is much shorter, 10–13 amino acids, than in Toc34. Positively charged amino acids in combination with a critical length of a hydrophobic transmembrane segment might represent a necessary combination of signals that is recognized by the Tic machinery. The unique import pathway of preToc75 into the outer envelope membrane seems to involve a stop transfer signal that is present in the COOH-proximal part of the cleavable transit sequence (Tranel and Keegstra, 1996). However, the nature of this signal is not clear, nor is it clear whether it is deciphered by the Tic or the Tic machinery. Hydrophobic stop transfer regions from mouse immunoglobulin M or vesicular stomatitis virus glycoprotein, which are both necessary and sufficient to halt translocation of proteins across the endoplasmatic reticulum, do not halt translocation of proteins into chloroplasts (Lubben et al., 1987).

Our results (Figs. 3–6) clearly demonstrate that pS34-Cinv is jointly imported by the Tic and the Tic complexes and not via a Tic-independent pathway. The recognition of the stop transfer signal must therefore occur within the Tic complex. A fraction of processed iS34Cinv is still in a proteinaceous environment, maybe in the pore component Toc75. Most of the protein is resistant to extraction at pH 11.5, which is indicative for the insertion and interaction with the lipid moiety of the membrane. Our results and those obtained for preToc75 (Tranel and Keegstra, 1996) indicate that the Tic complex harboring the precursor polypeptide translocation intermediate does not influence whether or not the Tic machinery can release a transmembrane segment into the lipid bilayer. Translocation arrest leaves the hydrophobic transmembrane segment in the pore of the Tic complex, where it might have sufficient time to “sense” the lipid environment, which induces the release into the bilayer. The mechanism of release of the protein into the plane of the membrane is not known but could be analogous to that proposed by Martoglio et al. (1995) and Liao et al. (1997) (for review see Siegel, 1997).

The partially processed polypeptide iS34Cinv remains locked in the Tic complex in the presence of ATP while the membrane anchor has already inserted into the lipid bilayer of the outer membrane. Simultaneously, the Tic complex becomes available for a new round of precursor recognition and translocation initiation (Fig. 6). This indicates that in intact chloroplasts, the Tic complex can function independently of the Tic machinery, corroborating earlier observations (Waegeemann and Soll, 1991) that isolated purified outer envelope vesicles are also functional in specific precursor recognition and partial translocation.

The ATP-dependent lock that keeps iS34Cinv in the Tic complex could be represented by the action of hsp100, which interacts on the stromal site with the Tic complex (Akita et al., 1997; Nielsen et al., 1997). The engagement of the Tic complex by iS34Cinv leads to a block of these translocation sites and to the inhibition of further import. By the removal of ATP, the inhibition is released because iS34Cinv retracts from the Tic complex and becomes exposed to the intermembrane space, indicating that no further ionic interactions keep the protein in the Tic complex. The chloroplastic inner envelope contains an active ATP/ADP carrier that might be responsible for the rapid drop in chloroplastic ATP levels (Neuhaus et al., 1997). iS34Cinv can probably not reengage the Tic machinery because it possesses only an incomplete import signal.

Intermediate processing of the presequence of SSU can occur if the proper site is not available to the stromal-processing peptidase (Archer and Keegstra, 1993). About 120 amino acids are spaced between the authentic processing site and the beginning of the membrane anchor domain. This should suffice to guarantee faithful processing if both the outer and inner envelope translocons were in close proximity to each other during the entire import event. Our data indicate that upon the release of the transmembrane segment into the plane of the membrane, the joint translocation sites dissociate and clear away from each other. This scenario would require a much longer spacing between the outer envelope and the Tic complex, which could explain our results (see also Fig. 5 C). It seems also unlikely that iS34Cinv is inserted into the outer membrane in a loop structure for two reasons: (a) iSSU34Cinv is resistant to the protease thermolysin, and (b) a loop structure would require a topology Ncyto-Cinv, but we have demonstrated (Fig. 2) that mS34Cinv inserts in a topology Ncyto-Ccyto.

In the constructs pS34C and pS34Cinv, competition exists between two targeting signals, which could lead the polypeptides on alternative translocation pathways. Under experimental conditions that are optimized for import (2 mM ATP and 25°C), pS34C and pS34Cinv use preferentially the standard import pathway via the Tic and Tic complex. Removal of the presequence receptors from the chloroplast surface shifts pS34C and pS34Cinv to the outer envelope insertion pathway via the COOH-terminal hydrophobic region. Similarly, preincubation of chloroplasts under binding conditions (50 μM ATP at 4°C) for 10 min also results in a shift to the Tic-independent insertion pathway. While both pathways may operate simultaneously, our data establish that transport conditions can be selected that favor one over the other.

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