Procoat, the Precursor of M13 Coat Protein, Inserts Post-Translationally into the Membrane of Cells Infected by Wild-Type Virus

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In growing cells infected by wild-type coliphage M13, the synthesis of procoat protein is completed before it inserts into the plasma membrane and is converted to coat protein.

The synthesis and membrane insertion of coliphage M13 coat protein (gene 8 product) has membrane assembly $(8, 14)$. Coat protein is membrane assembly $(0, 14)$. Coat protein is made as a precursor, termed procoat, with 23 additional $NH₂$ -terminal residues (10). Procoat is only synthesized by polysomes which are not is only synthesized by polysomes which are not attached to the plasma membrane (5). In earlier studies, cells infected by M13 or by M13 amber
7 mutant virus were pulse-labeled with $\int_0^3 H$ lproline, chased with nonradioactive proline, fractionated, and analyzed by sodium dodecyl sultionated, and analyzed by sodium dodecyl sulate-polyacrylamide gel electrophoresis and fluorography (2-5). The amber 7 mutant was used
because the conversion of procoat to coat protein was slower in cells infected by this virus than in cells infected by wild-type M13 (5) . In amber 7 M13-infected cells, pulse-labeled procoat is found in the cytoplasm. Procoat then binds to found in the cytoplasm. Procoat then binds to the inner surface of the plasma membrane and, in the presence of the transmembrane electro-
chemical potential, inserts into the bilayer; it is then cleaved to coat protein plus leader peptide. However, amber 7-infected cells accumulate in- H^2 is a subset H^2 infected cells accumulate in t_{t} are not include the architecture (9), although they continue to metabolize. This has synthesis occurs by the same pathway in these cells as in cells infected by wild-type virus. che as in cells intered by who-type virus.

Soluble procoat and procoat bound to the inner surface of the plasma membrane have also been detected in pulse-labeled cells infected by
all time M19 (0.5) at 9790. However, meat of wild-type M15 (2-5) at 37 C. However, most of
he labeled present were closedy convented to the labeled procoat was already converted to Thus, the internal procoat seen in these experi-Thus, the internal procoat seen in these experiments was not proven to be the precursor of coat

We now report the kinetics of procoat synthe-
is mombrane insertion, and proteclutie proc. sis, membrane msertion, and proteolytic proc-
saine in calls infected by wild type M19 et 40% essing in cells infected by wild-type M13 at 42° C, where procoat is metabolized more slowly that
4.3790. A clear presumer product relationship at 370C. A clear precursor-product relationship

is found between procoat and coat proteins. Procoat is only found inside the cell, hidden from

FIG. 1. E. coli HJM114 (13) was grown at 42° C in medium containingM9 salts (7), glucose, and vitamin B_1 to an absorbance at 600 nm of 0.4 and infected with wild-type M13 at a multiplicity of 100. After 1 h, 3.2 ml of culture was pulse-labeled with $\int^3 H$]proline $(100 \ \mu$ Ci, 110 Ci/mmol, New England Nuclear Corp.) for 15 s and chased with 1.6 mg of proline. Portions (0.8 ml) were chilled with crushed ice, collected by centrifugation (10 min, 4° C, 30,000 \times g), and suspended in 100 μ l of a solution of 10% (wt/vol) sucrose, ³⁰ mM Tris-chloride (pH 8.1), ¹⁰ mM EDTA, and ¹ mg of lysozyme per ml. After 30 min at 0°C, one-half of each fraction was mixed with trypsin (0.23 mg in $100 \mu l$ of 20% sucrose-30 mM Tris-chloride [pH 8.1]-10 mM EDTA) and incubated for 30 min at 0° C. After addition of soybean trypsin inhibitor $(0.4 \text{ mg}, \text{World})$ ington Diagnostics Corp.), proteolyzed spheroplasts i ugen Diagnostics Corp.), proteolyzed spheroplasts
were centrifuged (0°C, 15,000 \times g, 10 min) mixed with 500 μ l of 5% trichloroacetic acid, and prepared for sodium dodecyl sulfate-polyacrylamide gel electrophoresis as previously described (3). Spheroplasts mich were not treated with trypsin were precipitated with 500 μ l of 5% trichloroacetic acid. Lanes 1, 3, 5, and 7 were wandat no trypsin treatment, lanes 2, 4,
in and 8 were with transin treatment. Observings 6, and 8 were with trypsin treatment. Chase times were as follows: 5 s, lanes ¹ and 2; 25 s, lanes 3 and 4; 45 s, lanes 5 and 6; 75 s, lanes 7 and 8. P, Protein.

external protease by the plasma membrane. It is post-translationally inserted into the membran and converted to coat protein.
Escherichia coli HJM114 was grown at 42°C

in minimal medium, infected by \overline{M} 13, and pulselabeled for 15 s with $[^3H]$ proline. Unlabeled proline was added at the end of the pulse period. After 5, 25, 45, and 75 s of chase, portions of the culture were rapidly cooled by addition to ice. culture were rapidly cooled by addition to ice. Each portion was converted to spheroplasts at 00C, and one-half of each spheroplast preparation was digested with trypsin. Each sample was analyzed by sodium dodecyl sulfate-polyacryl-1). After only 5 s of chase (lane 1), very little of 1). After only σ s of chase (lane 1), very little σ the procoat was converted to the coat protein. The NH2 terminus of the coat protein is exposed on the outer surface of the membrane (12, 13).
Trypsin cleaves off the first 8 residues (13), including the sole proline residue of coat protein at position $6(11)$. This renders the remaining 42 at position σ (11). This renders the remaining σ residues of coat protein invisible on the fluor graph (lane 2). Procoat, gene 5 protein (1), and ϵ other cytoplasmic proteins of higher molecular weights are protected from the trypsin by the plasma membrane (2).
During the succeeding 60-s chase, coat protein

was formed as the procoat was lost (lanes $1, 3, 5$, and 7). This strongly suggests a precursor-product relationship. This relationship has been seen in each of several such experiments and has been confirmed by quantitative densitometry of the fluorographs. At each time of chase, the procoat was inaccessible to trypsin, whereas the coat was inaccessible to trypsin, whereas the coat protein was trypsin sensitive (lanes 2, 4, 6, and 8). Although coat protein inside the cell or transassembly intermediate if cleavage occurred before or after insertion, respectively, neither spefore or after insertion, respectively, neither sp cies was abundant enough to be detected in these experiments.
Although these experiments were qualita-

tively similar to those reported for amber 7 M13infected cells, $E.$ coli was not killed by wild-type virus infection. Cells infected at 42°C continued to grow (Fig. 2) and shed virus. Although the to grow (Fig. 2) and shed virus. Although the rate of virus production was much lower at $42⁰$ than at 37° C, a cen culture infected by M13 and
movies overwight at 499C aboved a 10⁴ feld is grown overing the 42 C showed a 10-fold is

crease in virus titer. This post-translational assembly pathway has been reproduced in an in vitro protein synthesis incubation (J. M. Goodman, C. Watts, and W. Wickner, manuscript in preparation) with polytoplasmic membrane vesicles from E , coli. The toplasmic membrane vesicles from E. coll. The collection of availability of pure procoat (P. Silver and W. Wickner, manuscript in preparation, and leader
montidees (C. Zwisinghi and W. Wickner, J. Die peptidase (C. Zwizinski and W. Wickner, J. Biol.

FIG. 2. Growth of E. coli JHM114 in medium con-
taining M9 salts, 1% glucose, and vitamin B₁ at 3
and 49°C was methoded by the abundance at 30 and 42° C was monitored by the absorbance at 600 nm (A_{800}) . Where indicated, cells were infected by wild-type M13 at a multiplicity of 10. Symbols: \bigcirc , uninfected cells at 37°C; \bullet , infected cells at 37°C; uninfected cells at 37°C; 0, infected cells at 49°C; \overline{a} , which cells at \overline{a} \overline{b} , \overline{a} , infected cells at \overline{a} .

Chem., in press) may allow these reactions to be studied with defined components.

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