

## I. Background

## A. Protein sorting

1. signal sequences

## II. Cotranslational Translocation

- A. Signal sequence
- B. Signal-recognition particle (SRP)
- C. SRP receptor
- D. SRP & SRPr binding
- E. Translocon

## III. Integral Membrane Proteins

- A. Topological classes
  1. Type I
  2. Type II
  3. Type III
  4. Type IV
- B. GPI anchored proteins
  - a. transamidase
- C. Hydrophathy profile

## IV. Post-translational Modifications in ER &amp; Golgi

## A. Glycosylation

1. O-linked
  - 1) glycosyltransferases
    - a. mucin-type O-linked
    - b. proteoglycans
  2. N-linked
    - a. oligosaccharyl transferase
      - 1) N-X-S or N-X-T
    - b. glycosidases
    - c. tunicamycin

## B. Disulfide bonds

1. protein disulfide isomerase

## C. Folding

1. lectins
2. Genetic Control
  - a. Ire1
3. Destruction

## V. Folded Translocation

## A. Nucleus Background

- a. double membrane
  - 1) inner membrane
    - a) nuclear lamina
    - 2) outer membrane
  - b. nuclear pores
    - 1) Nuclear Pore Complex
      - a) nucleoporins

## B. Nuclear localization signal

## C. Transport proteins

1. Importin
2. Ran-GTP

- D. Nuclear export
- 1. exportin

#### VI. Unfolded Translocation

- A. translocon
- B. Driving force
- 1. Sec63
- 2. BiP
- C. Mitochondria Transport
- 1. Overview
- 2. Matrix Targeting sequence
- 3. Process - initial step
- a. import receptor
- b. TOM
- 4. E for transport
- a. Hsc70
- 5. Folding