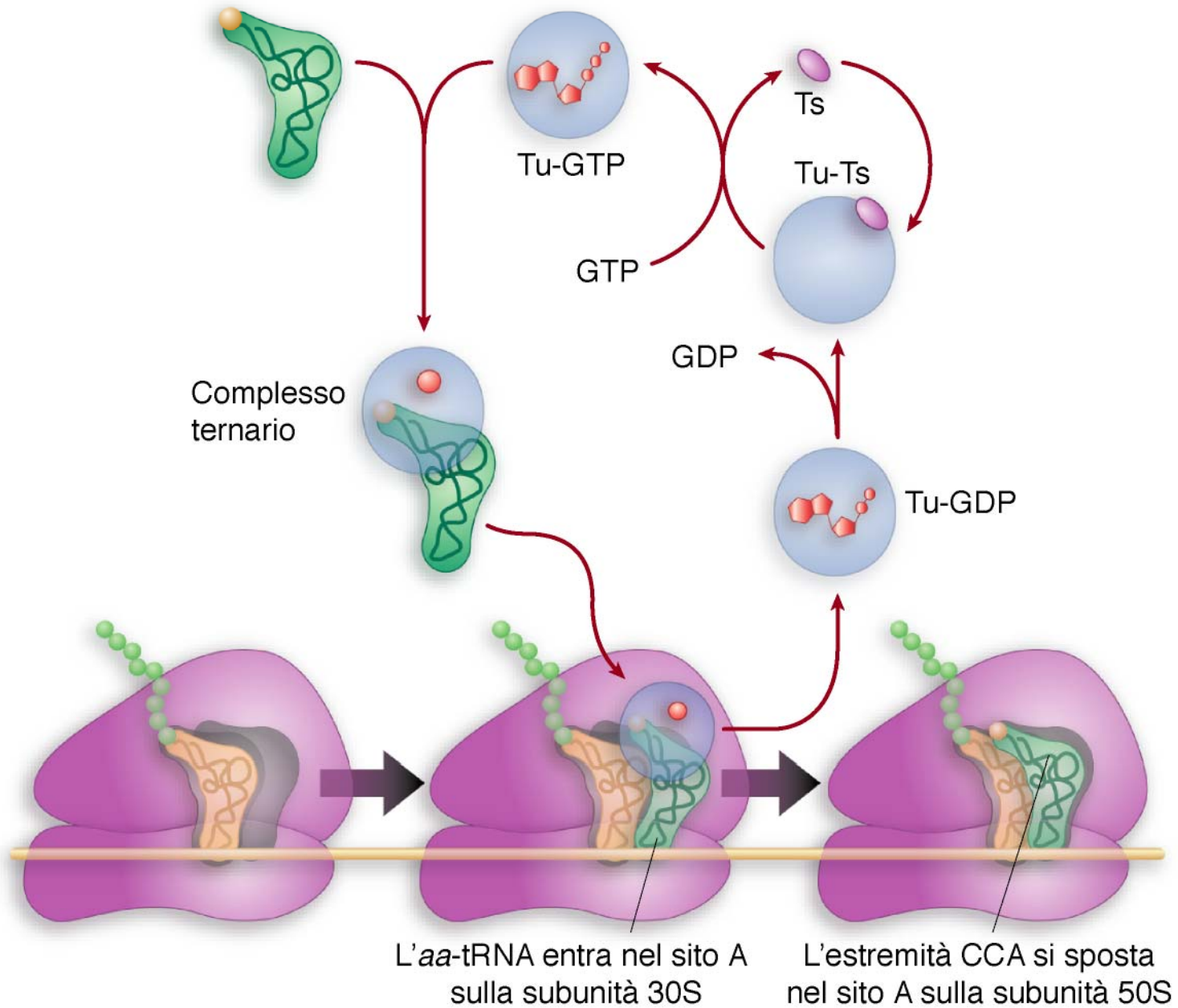


Traduzione: l'allungamento

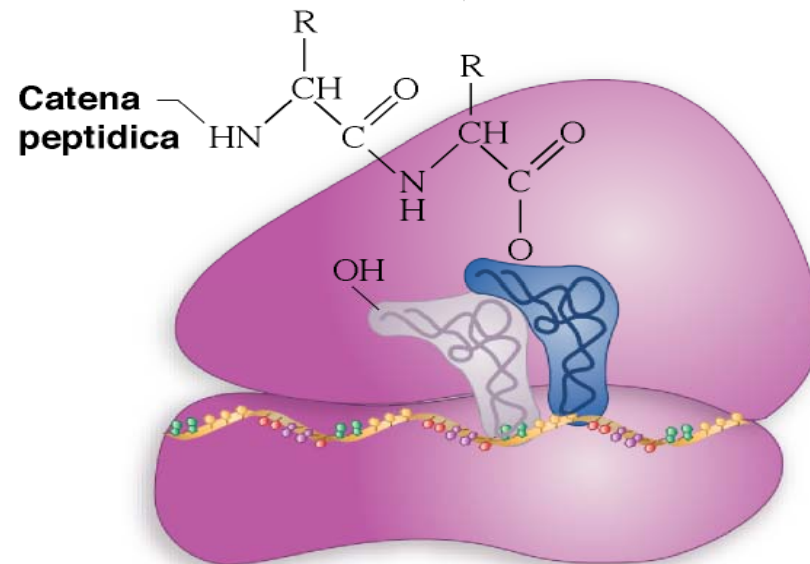
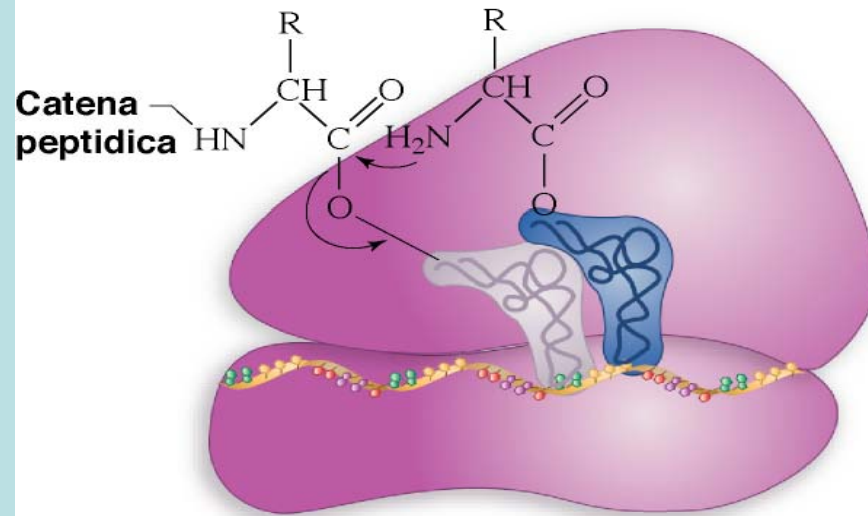
Fattori di allungamento

<i>Procarioti</i>	<i>Eucarioti</i>	
EF-Tu	eEF1α	trasporto aa-tRNA
EF-Ts	eEF1$\beta\gamma$	riciclo
EF-G	eEF2	traslocazione

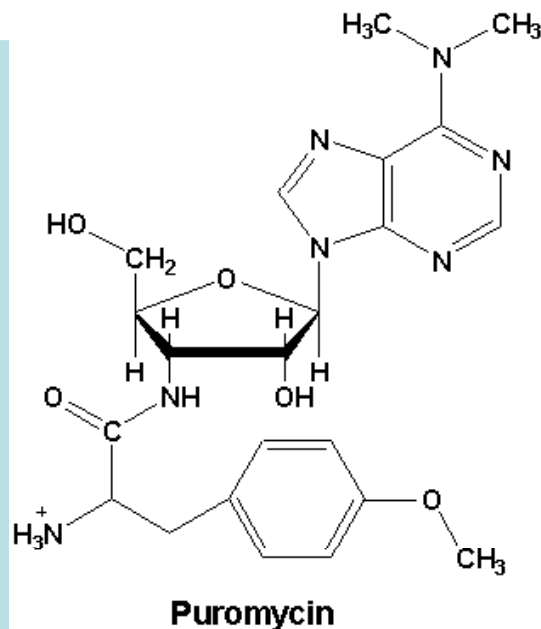
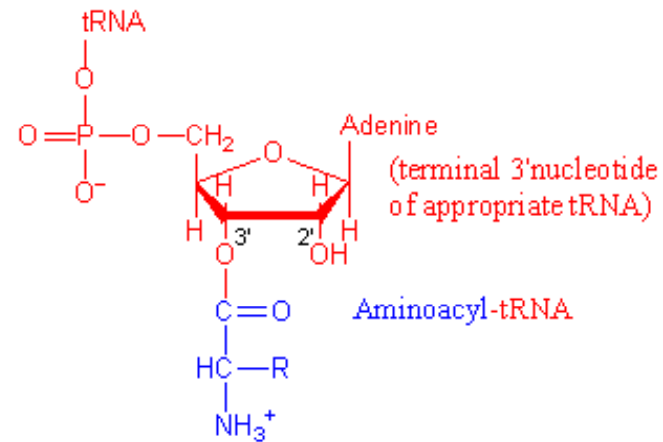
EF-Tu ricicla fra una forma legata al GTP e una legata al GDP



Il polipeptide nascente viene trasferito all'aa-tRNA

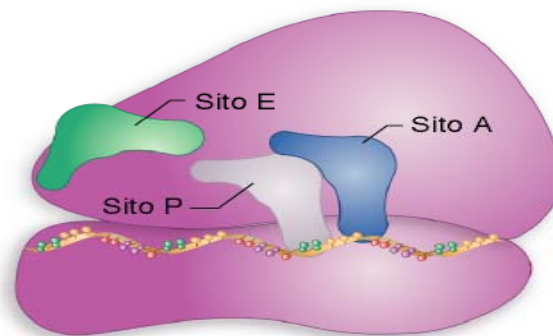


La puromicina è simile
a un amminoacil-tRNA

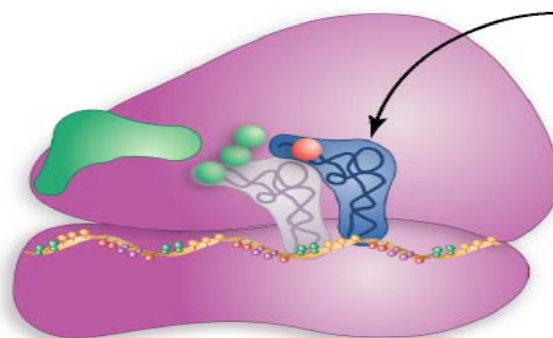


Puromycin is an aminonucleoside antibiotic, derived from the *Streptomyces alboniger* bacterium, that causes premature chain termination during translation taking place in the ribosome. It is not selective for either prokaryotes or eukaryotes.

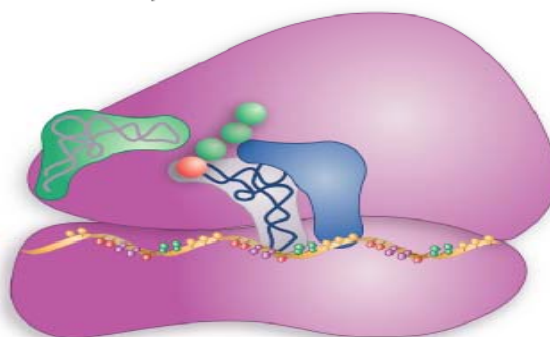
Il tRNA si sposta in 3 siti del ribosoma



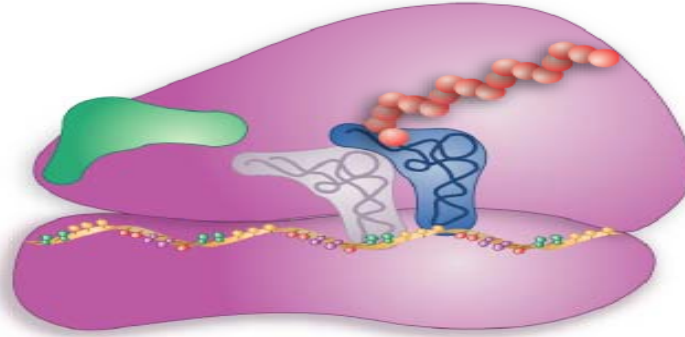
Pre-traslocazione:
il peptidil-tRNA è nel sito P;
l'amminoacil-tRNA entra nel sito A



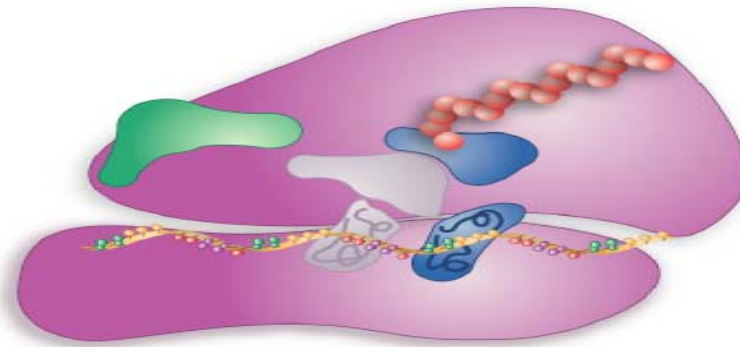
Post-traslocazione:
il tRNA deaccilato si sposta nel sito E;
il peptidil-tRNA si sposta nel sito P



La traslocazione avviene in due fasi

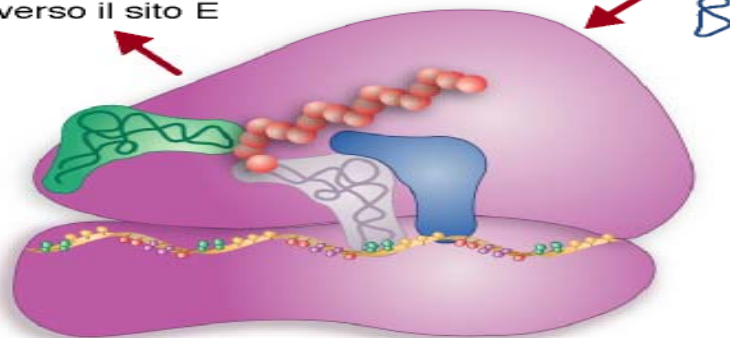


La subunità 50S
si sposta rispetto
alla 30S



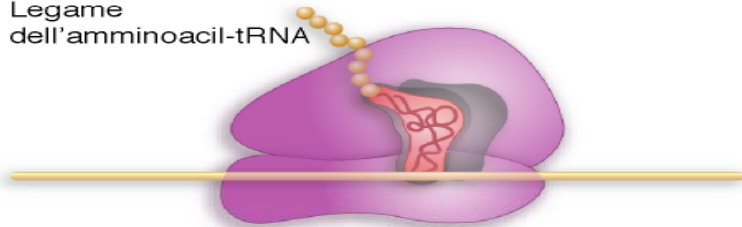
Il tRNA scarico esce
attraverso il sito E

aa-tRNA
in entrata



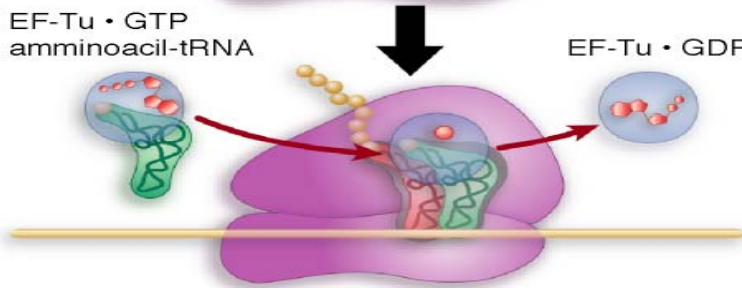
I fattori EF-Tu ed EF-G si legano alternativamente al ribosoma

Legame dell'amminoacil-tRNA

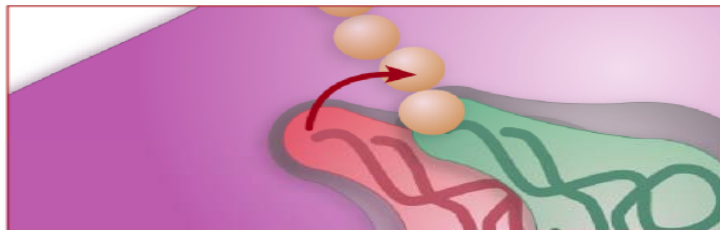


EF-Tu • GTP
amminoacil-tRNA

EF-Tu • GDP



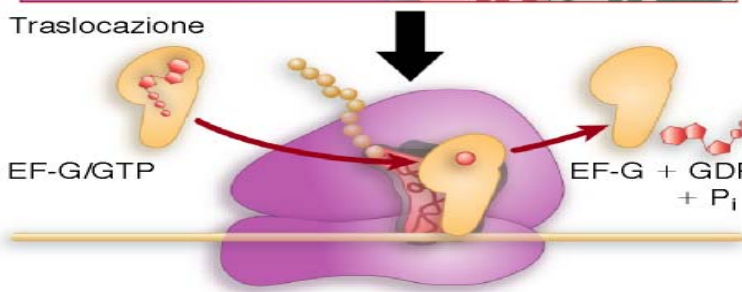
Sintesi del legame peptidico



Traslocazione

EF-G/GTP

EF-G + GDP + P_i

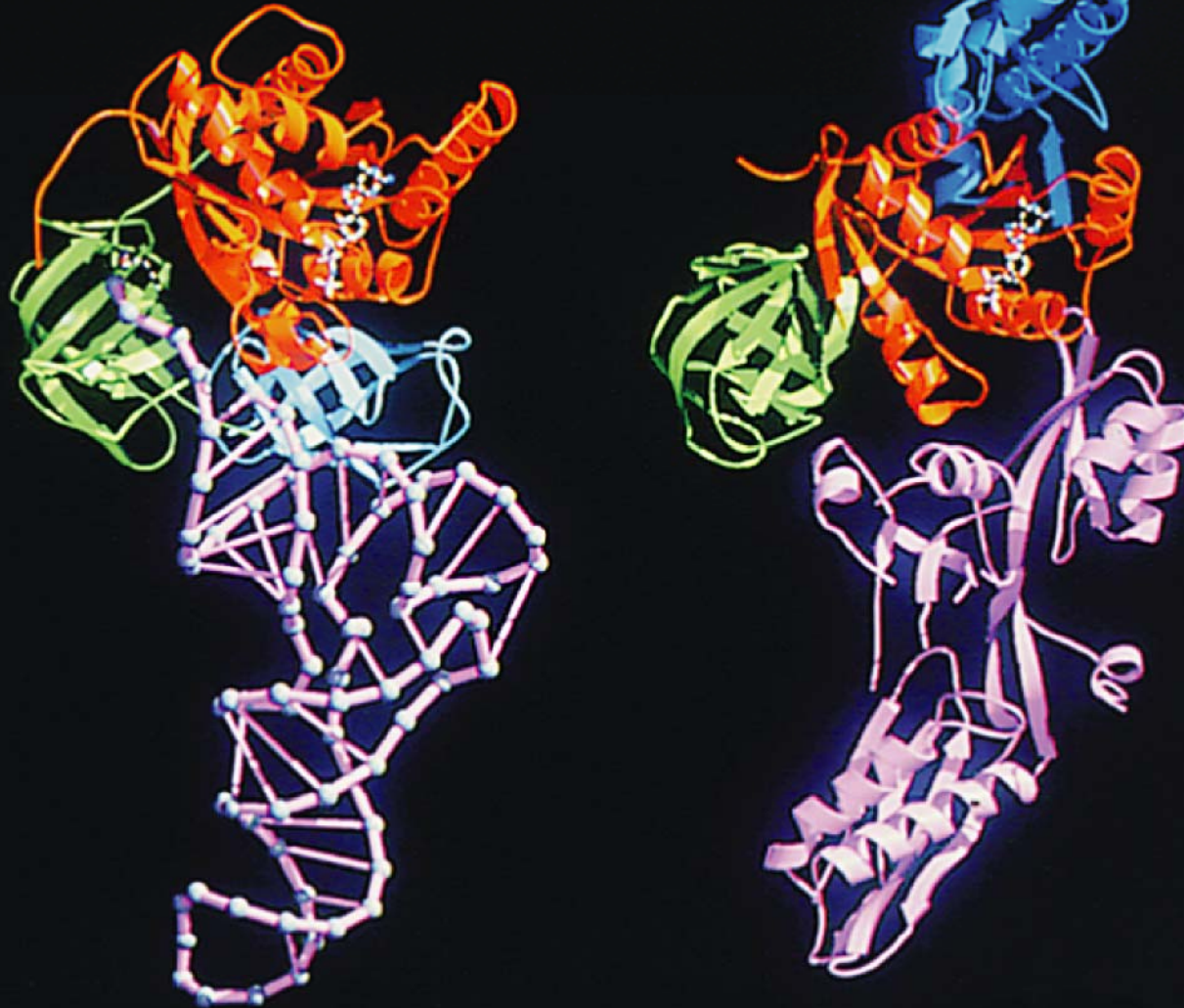


The factor **EF-G** catalyzes the translocation of the tRNA and mRNA down the ribosome at the end of each round of polypeptide elongation. Homologous to EF-Tu + tRNA, EF-G also binds to the ribosome in its GTP-bound state. When it associates with the A site, EF-G causes the tRNA previously occupying that site to occupy an **intermediate A/P position** (bound to the A site of the small ribosomal subunit and to the P site of the large subunit), and the tRNA in the P site is shifted to a **P/E hybrid** state. EF-G hydrolysis of GTP causes a conformation change that forces the A/P tRNA to fully occupy the P site, the P/E tRNA to fully occupy the E site (and exit the ribosome complex), and the mRNA to shift three nucleotides down relative to the ribosome due to its association with these tRNA molecules. The GDP-bound EF-G molecule then dissociates from the complex, leaving another free A-site where the elongation cycle can start again

La struttura di EF-G imita l'amminoacil-tRNA

Amminoacil-tRNA • EF-Tu • GTP

EF-G



Decoding by the 70S ribosome

Rebecca M Voorhees and T. Martin Schmeing

Ramakrishnan Laboratory
MRC-Laboratory of Molecular Biology

Traduzione: terminazione

Fattori di terminazione

<i>Procarioti</i>	<i>Eucarioti</i>	
RF1	eRF	riconoscimento UAA, UAG (ocra, ambra)
RF2	“	riconoscimento UGA, UAA (opale)
RF3	eRF3	GTPase
RRF		rilascio

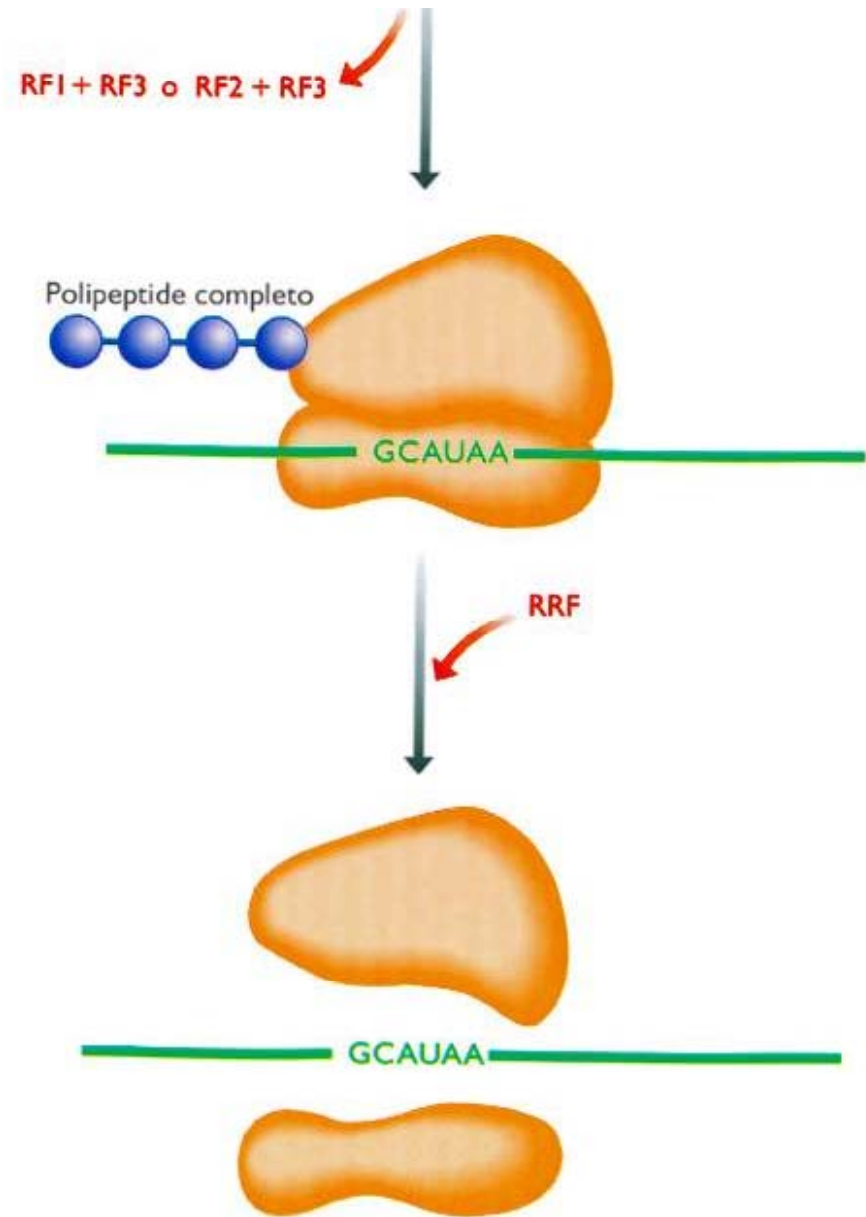
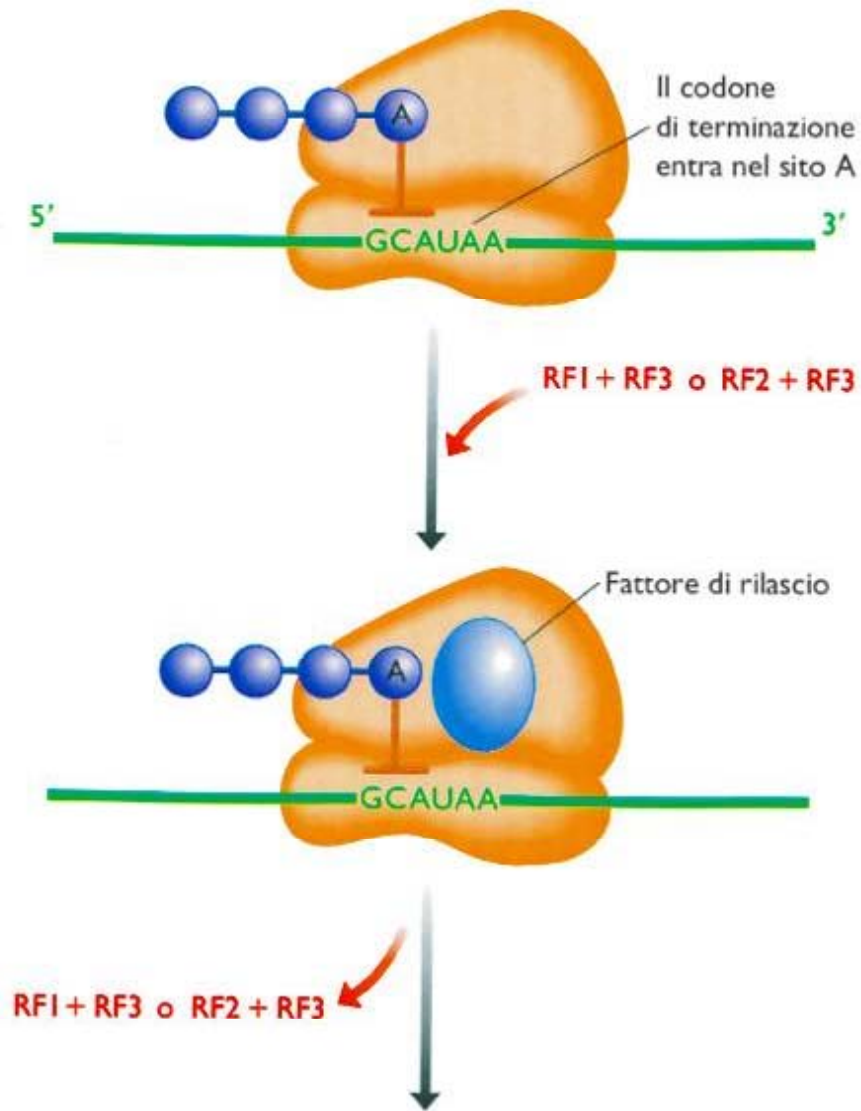
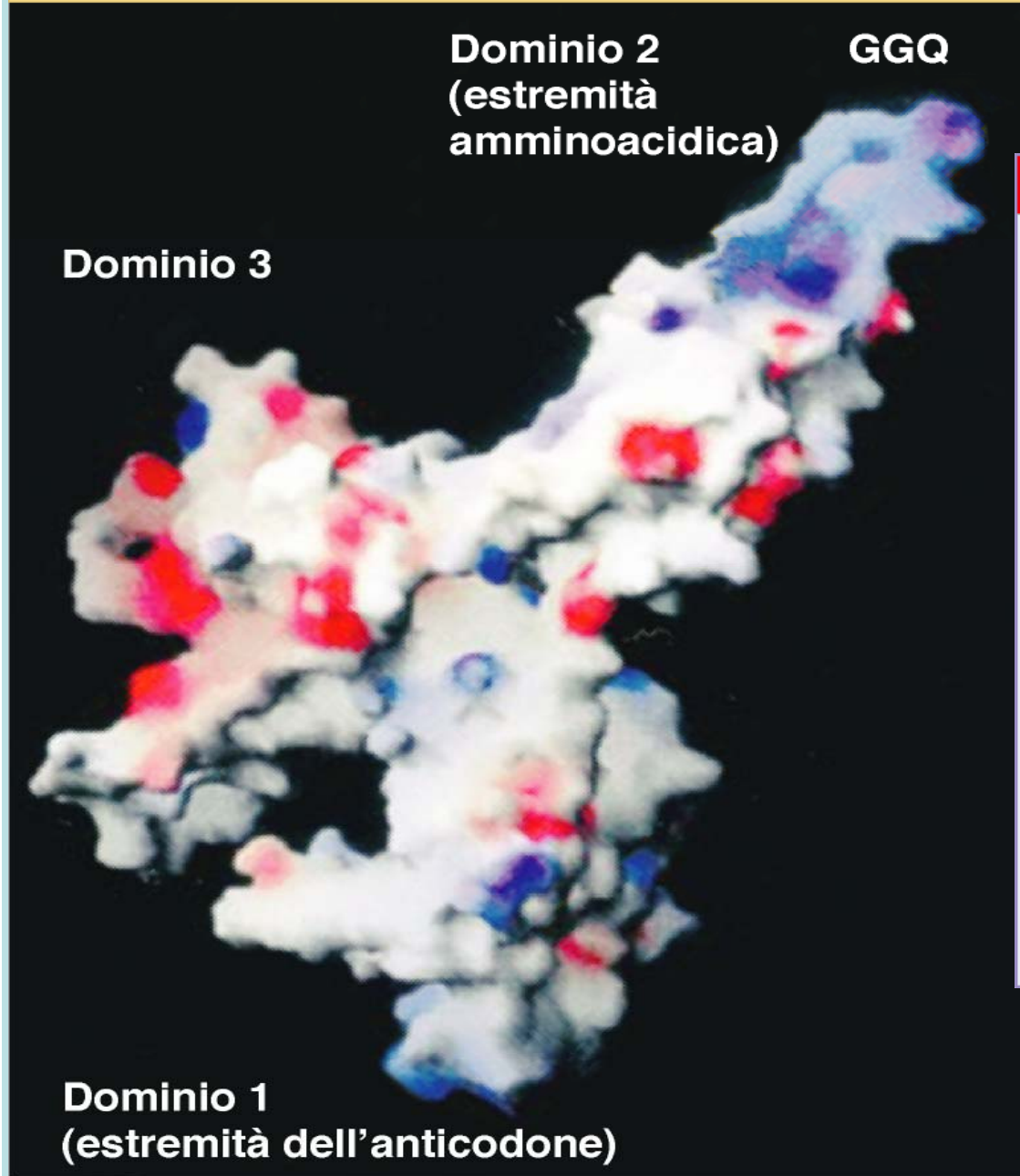
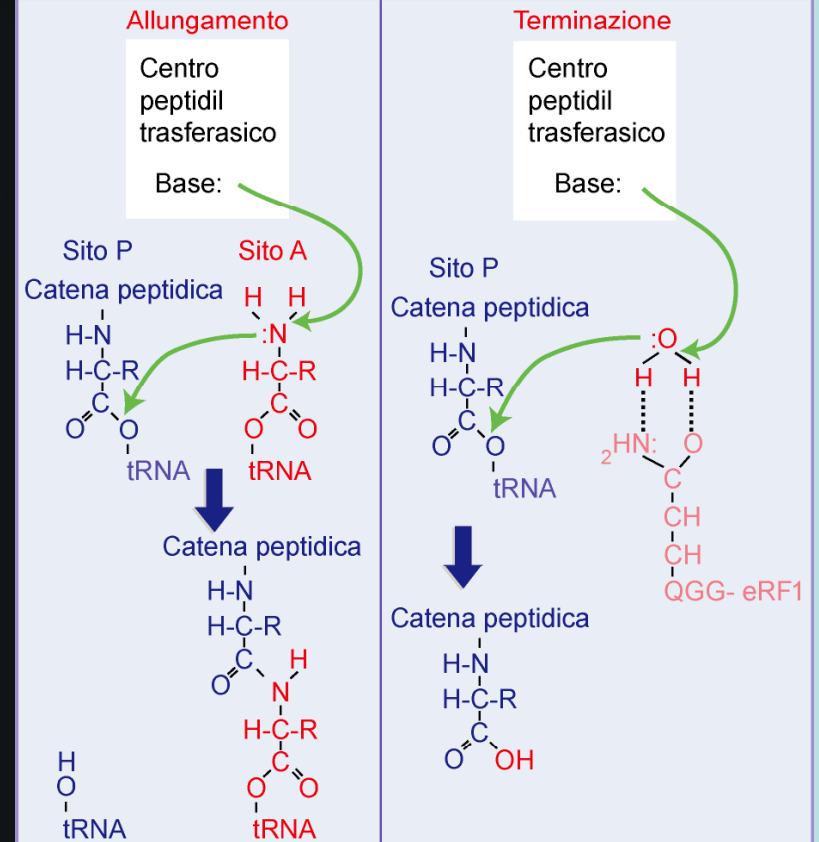


Figura 11.21 La terminazione della traduzione.

eRF1 imita il tRNA

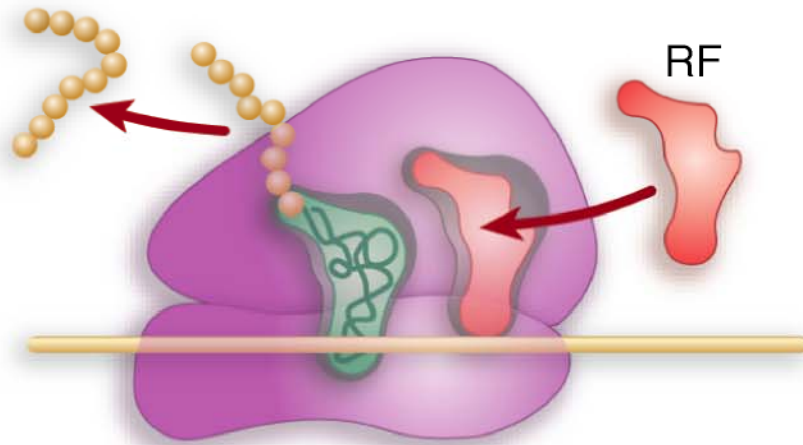


eRF1 presenta una molecola di acqua per idrolizzare il polipeptidil-tRNA

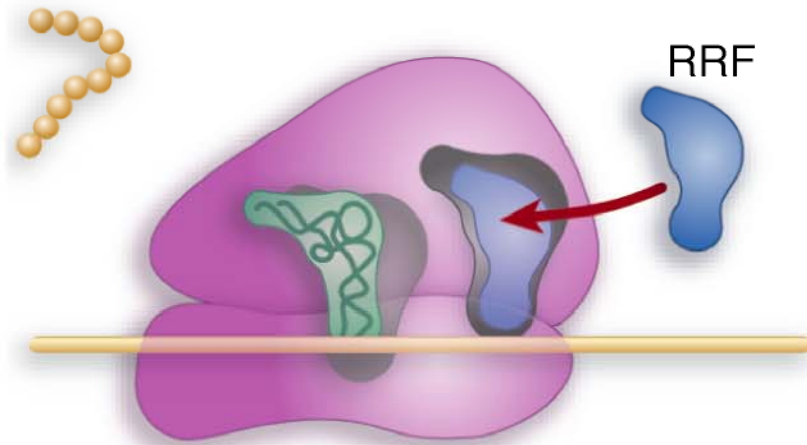


La terminazione richiede molteplici fattori proteici

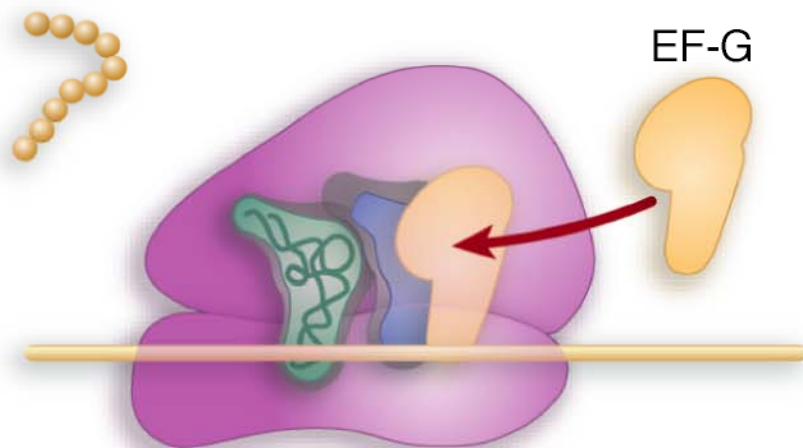
1. RF rilascia la catena proteica



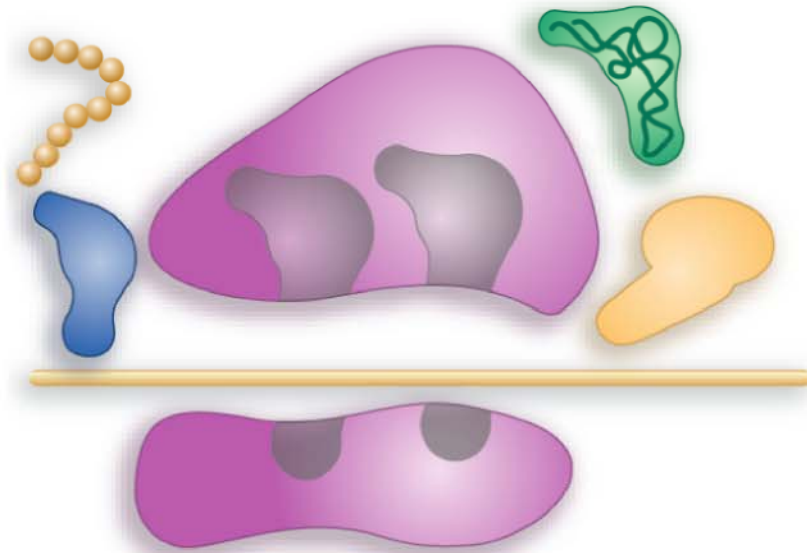
2. RRF entra nel sito A



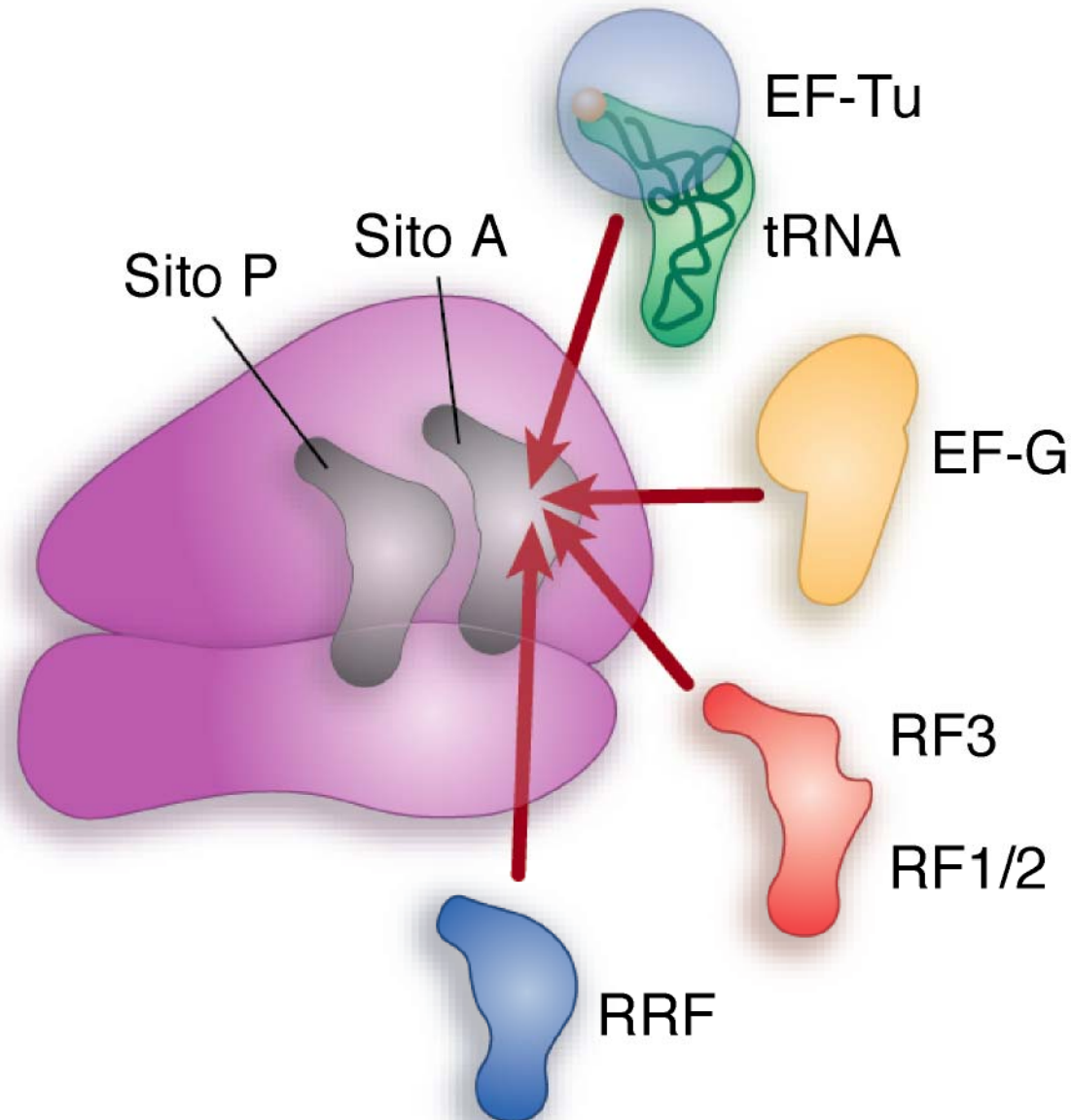
3. EF-G trasloca RRF



4. Il ribosoma si dissocia



Vari fattori sono simili nella forma


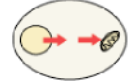





Inhibitor	Effect
Chloramphenicol	inhibits prokaryotic peptidyl transferase.
<u>Streptomycin</u> (Neomycin)	inhibits prokaryotic initiation, also induces mRNA misreading.
Tetracycline	inhibits prok. aminoacyl-tRNA binding to the ribosome small subunit.
Erythromycin	inhibits prokaryotic translocation through the ribosome large subunit.
<u>Fusidic acid</u>	similar to erythromycin only by preventing EF-G from dissociating from the large subunit.
<u>Puromycin</u>	resembles aa-tRNA, interferes with peptide transfer resulting in premature termination in prok. and euk.
Diphtheria toxin	catalyzes ADP-ribosylation of and inactivation of eEF-2.
Ricin	found in castor beans, catalyzes cleavage of the euk. 28S rRNA
Cycloheximide	inhibits eukaryotic peptidyltransferase.

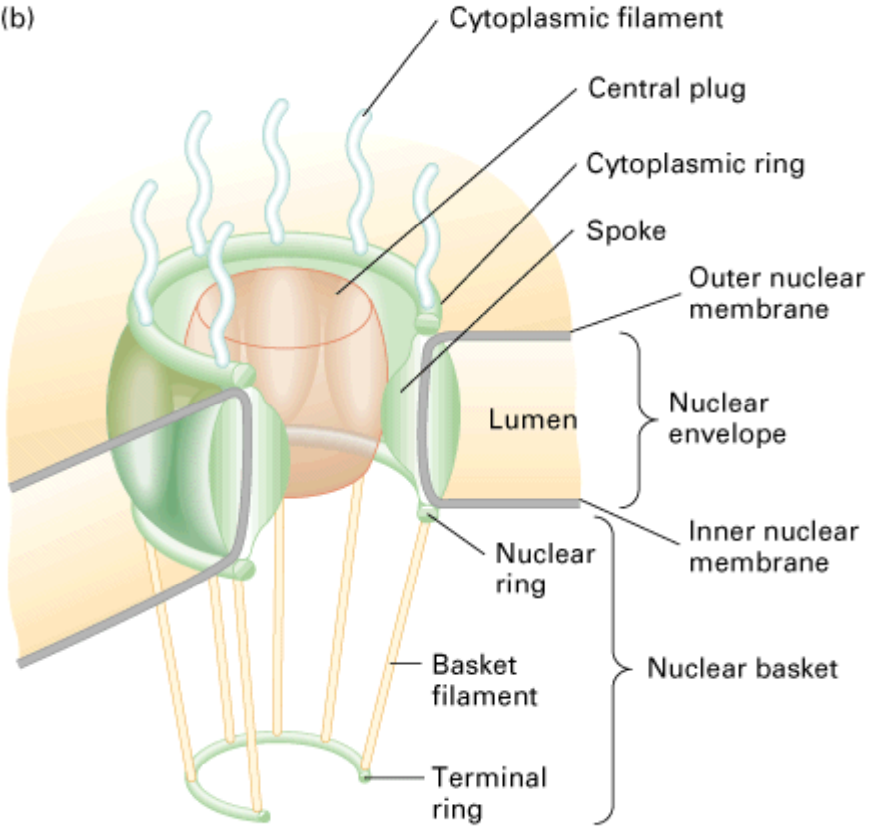
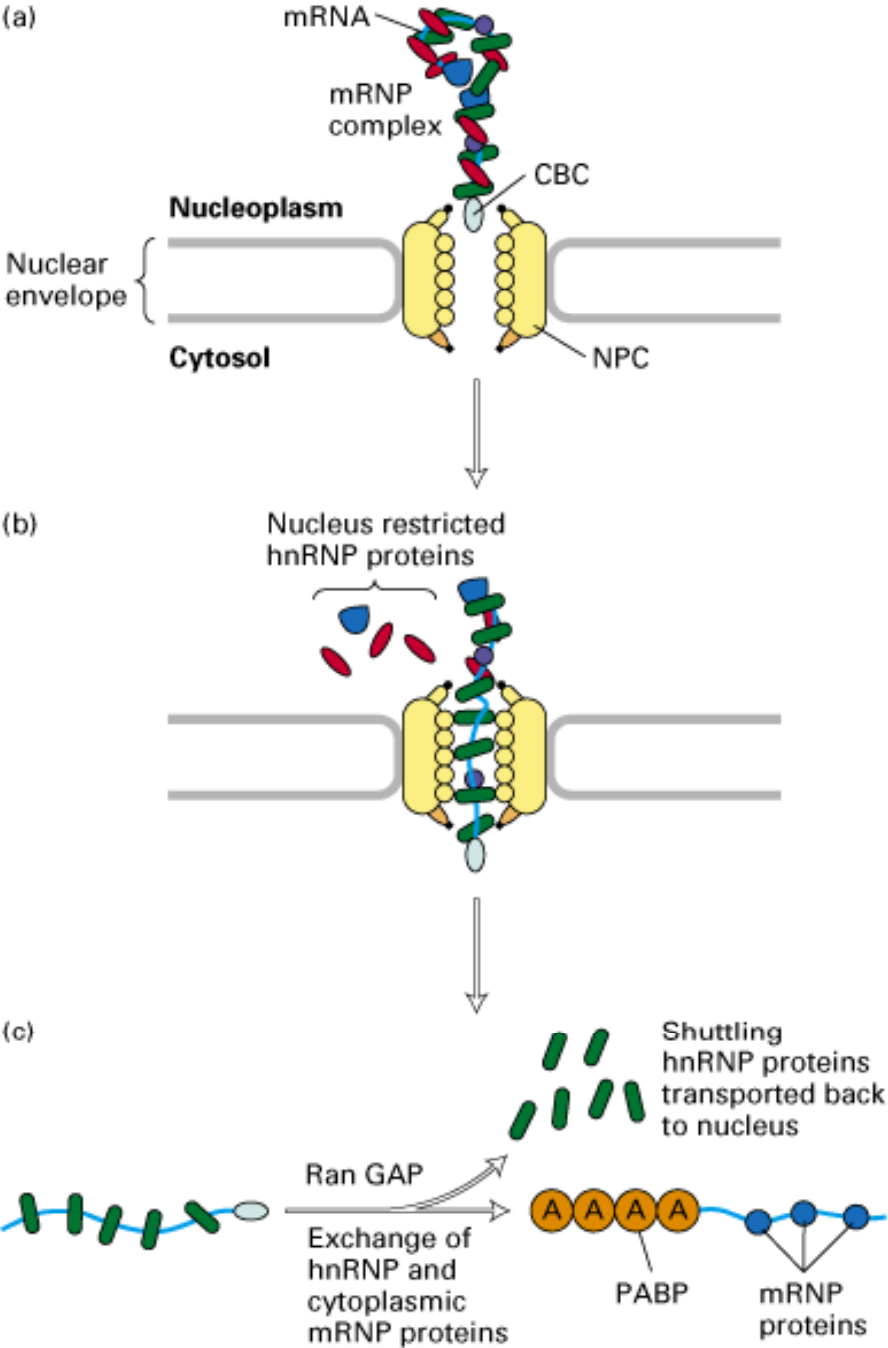
Trasporto e localizzazione degli mRNA

Negli eucarioti l'RNA viene traslocato e può essere localizzato

L'RNA può essere trasportato fra compartimenti cellulari

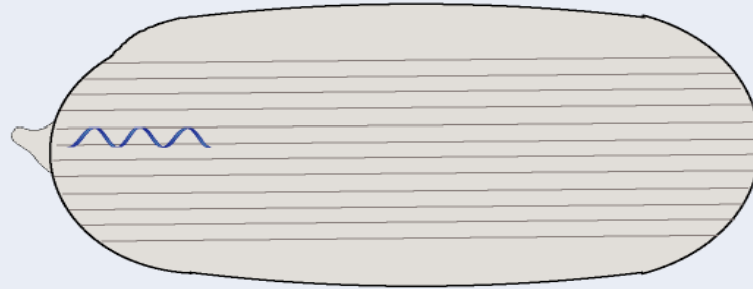
RNA	Trasporto	Localizzazione	
Tutti gli RNA	Nucleo → citoplasma	Tutte le cellule	
tRNA	Nucleo → mitocondrio	Molte cellule	
mRNA	Cellula nutrice → oocita	Embriogenesi di <i>Drosophila</i>	
mRNA	Polo anteriore → polo posteriore dell'oocita	Embriogenesi di <i>Drosophila</i>	
mRNA	Cellula → cellula	Floema delle piante	

Nuclear Pore Complex

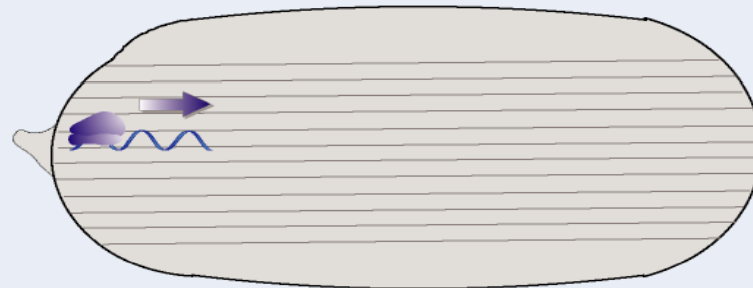


La localizzazione dell'mRNA genera un gradiente di proteina

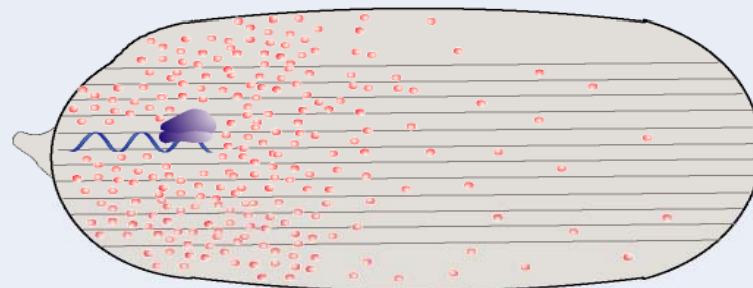
L'mRNA è localizzato all'estremità anteriore



L'mRNA è tradotto

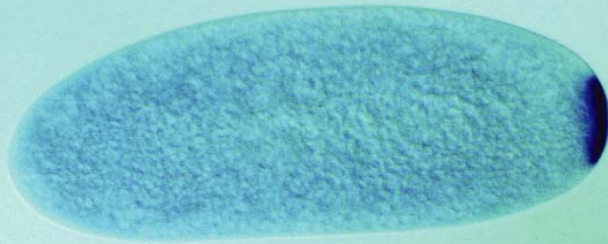


La proteina diffonde dal sito di sintesi

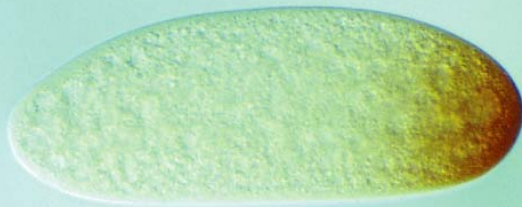


L'RNA e la proteina nanos sono posteriori

L'RNA *nanos* è localizzato all'estremità posteriore alla terza divisione

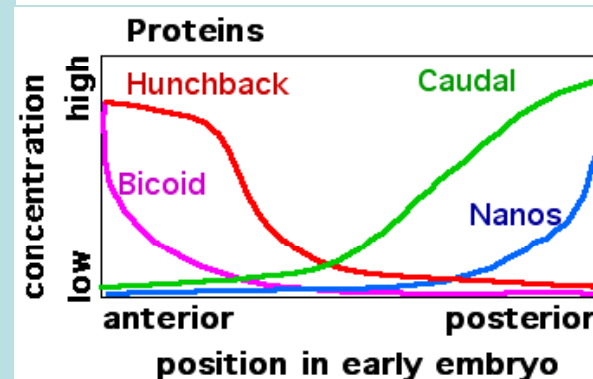
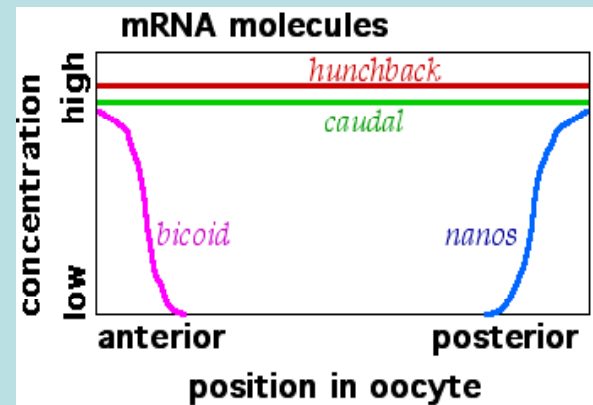


La proteina nanos diffonde dall'estremità posteriore all'ottava divisione



The developing egg (oocyte) is polarized by differentially localized mRNA molecules.

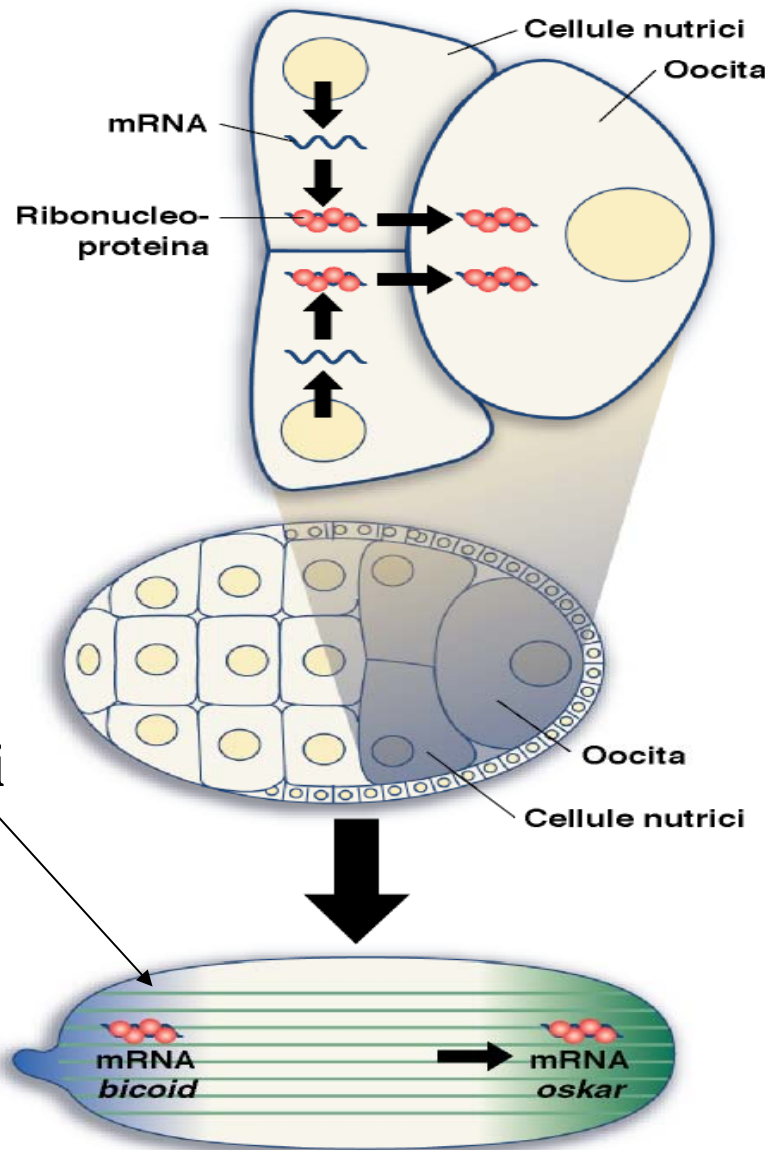
The genes that code for these mRNAs, called **maternal effect genes**, encode for proteins that get translated upon fertilization to establish concentration gradients that span the egg. *Bicoid* and *hunchback* are the maternal effect genes that are most important for patterning of anterior parts (head and thorax) of the *Drosophila* embryo. *Nanos* and *Caudal* are maternal effect genes that are important in the formation of more posterior abdominal segments of the *Drosophila* embryo.



Meccanismi di localizzazione

- **Trasporto specifico**
- **Degradazione selettiva**
- **Ancoraggio a siti specifici**

Gli mRNA possono essere trasportati nell'uovo



Microtubuli

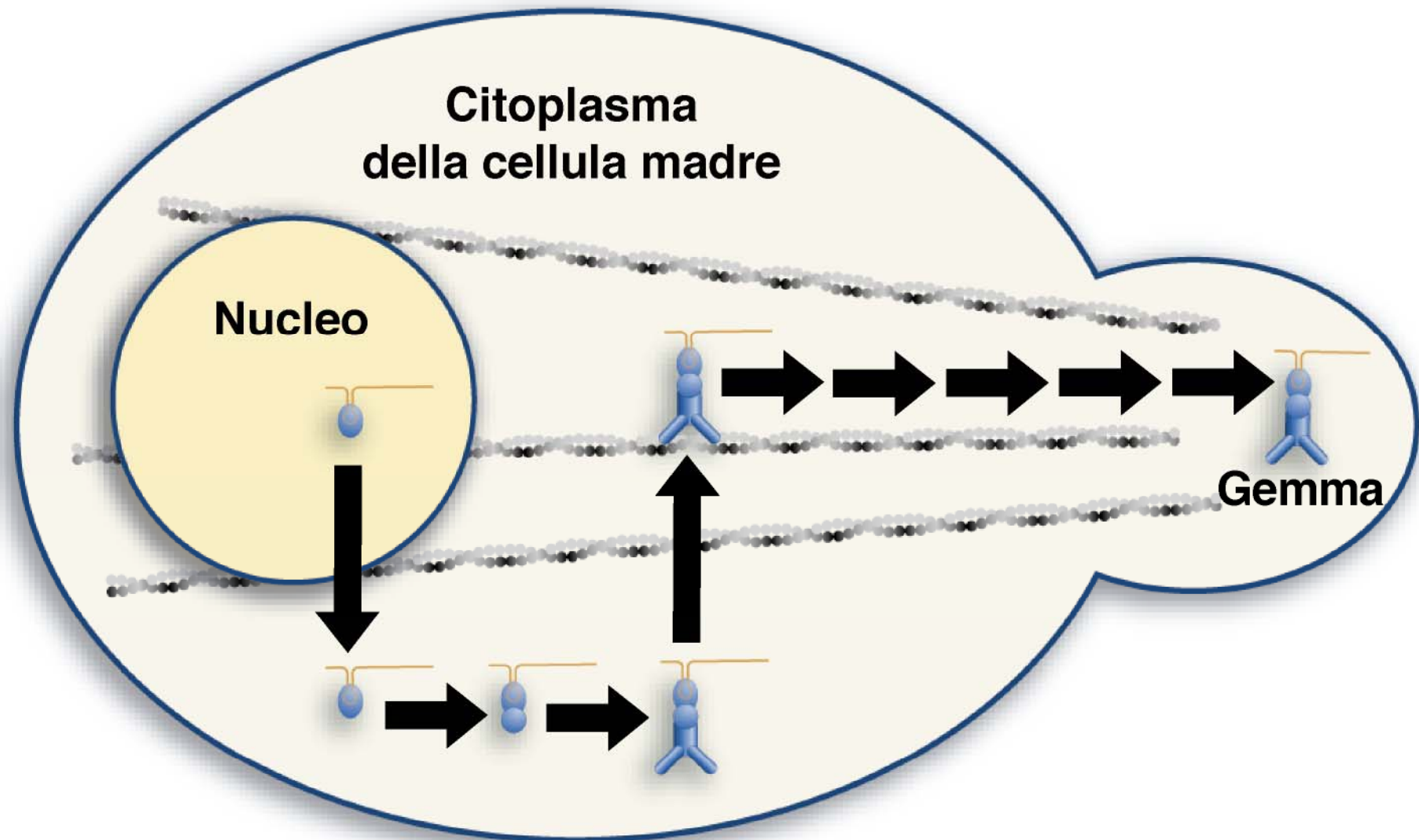
L'mRNA di *oskar* si sposta su binari fino all'estremità posteriore; l'mRNA di *bicoid* si localizza all'estremità anteriore

La localizzazione anteriore è determinata da particolari sequenze all'estremità 3' non tradotta (UTR) dell'mRNA

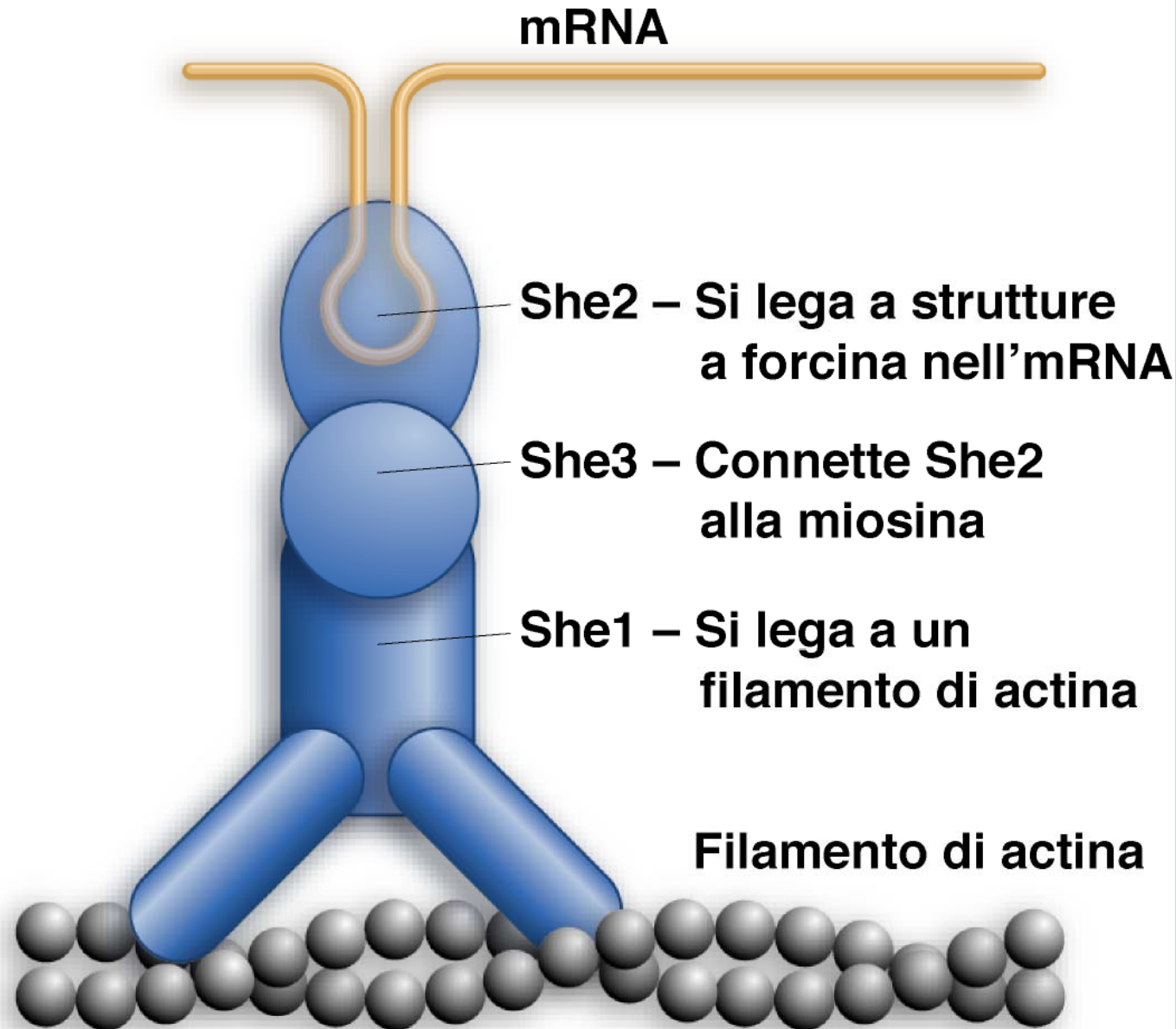
All'interno dell'oocita ci saranno concentrazioni diverse delle 2 proteine ai 2 poli della cellula → influenza sullo sviluppo dell'embrione

Nei lieviti alcuni mRNA si spostano dalla cellula madre alla gemma.

ASH1 si sposta lungo filamenti di actina fino alla gemma



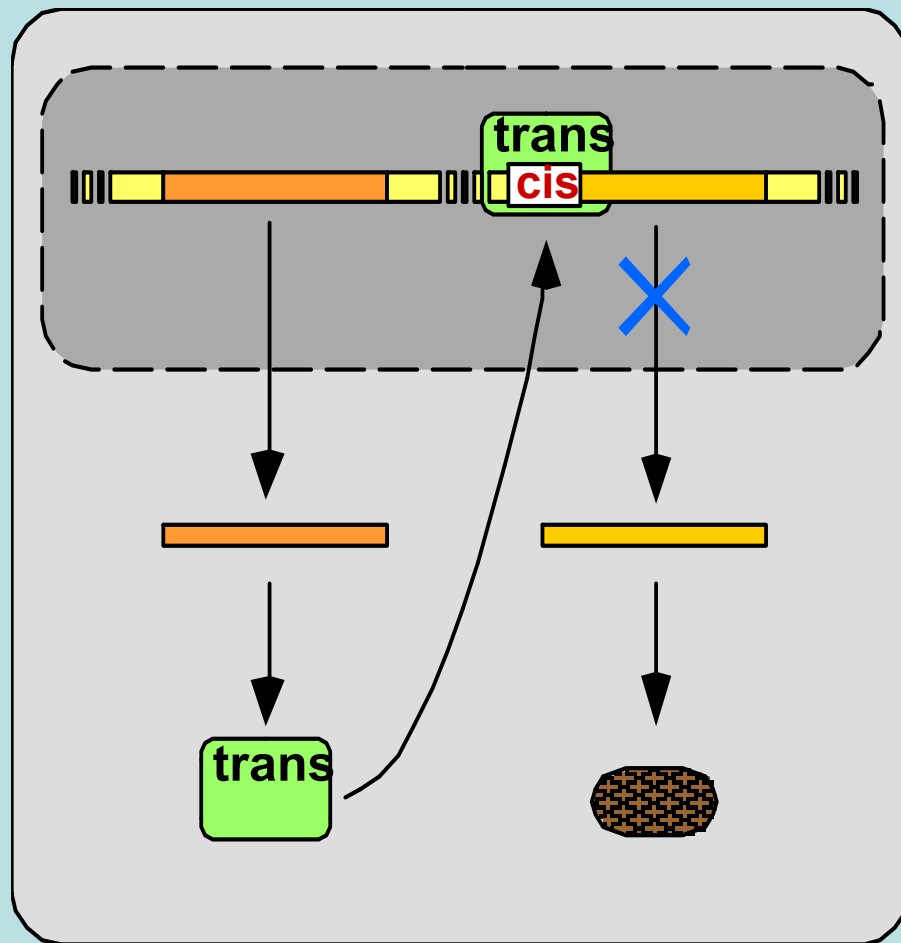
L'mRNA di ASH1 è legato a un motore



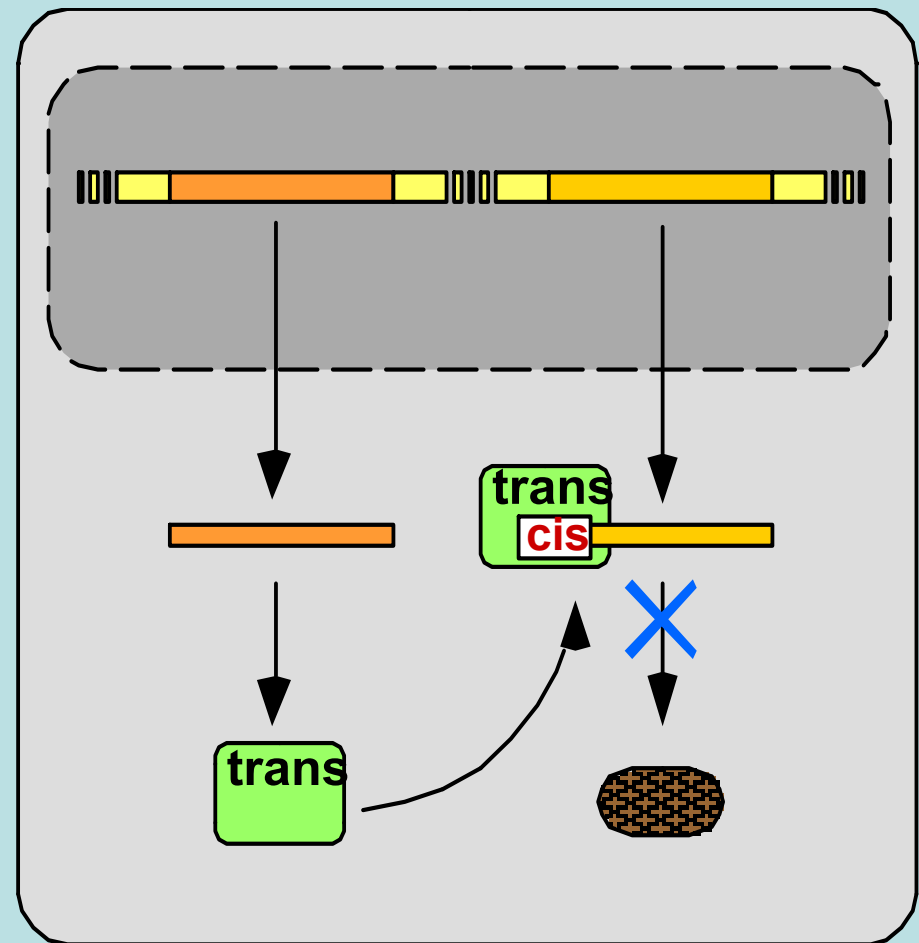
Regolazione della traduzione

- **Generale**
- **Specifica**

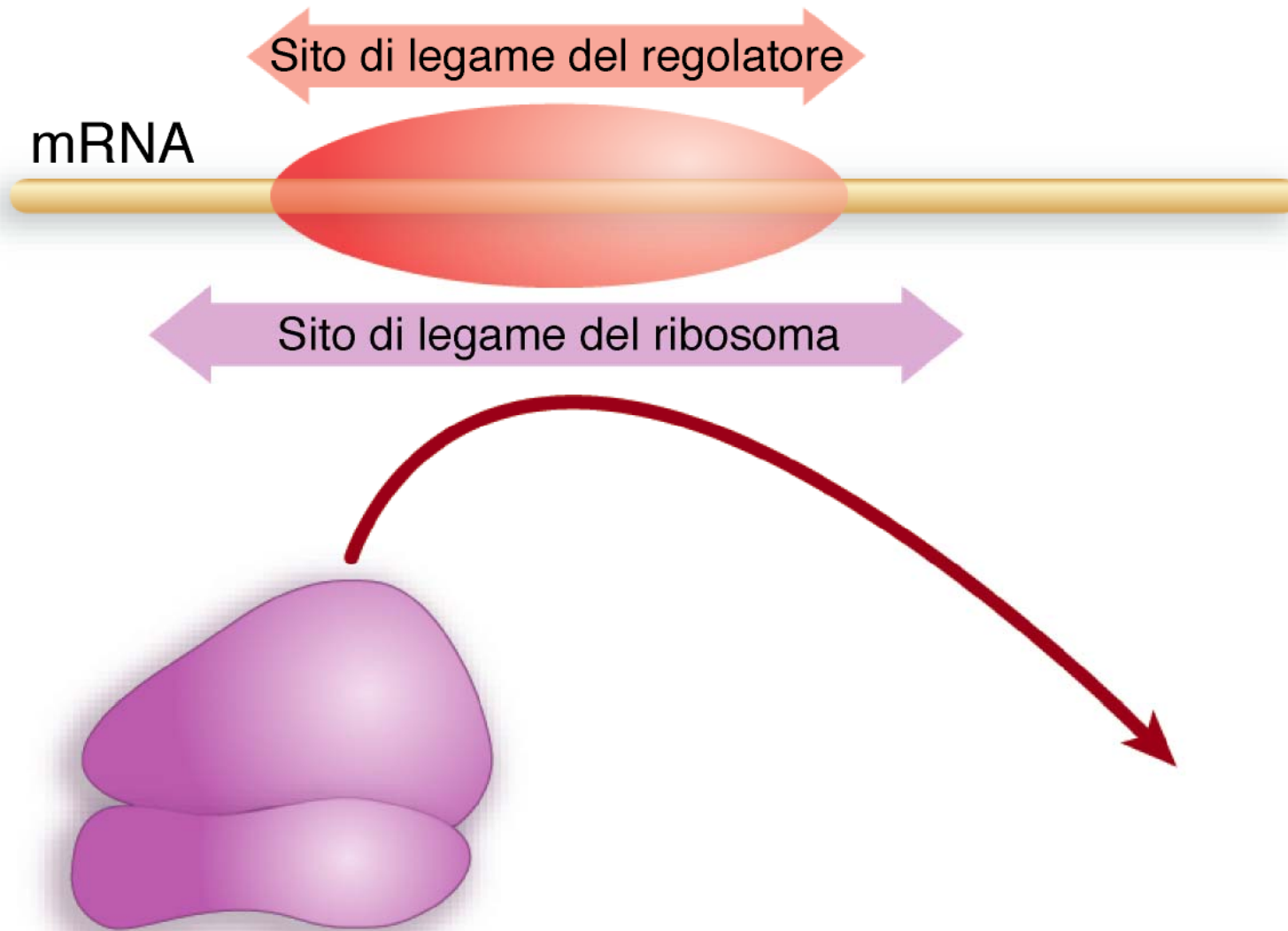
Regolazione trascrizionale



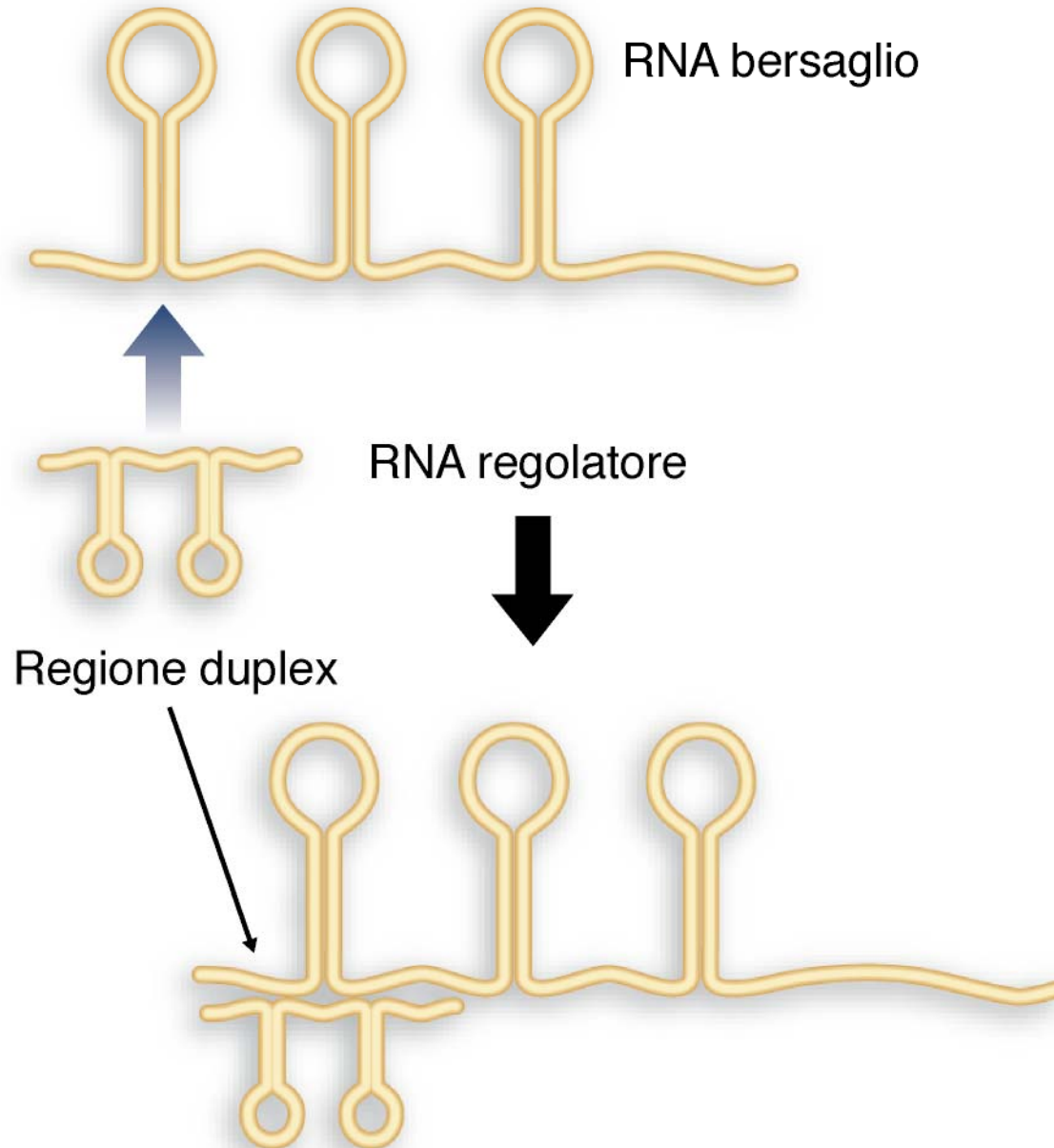
Regolazione traduzionale



Un regolatore può bloccare il legame del ribosoma

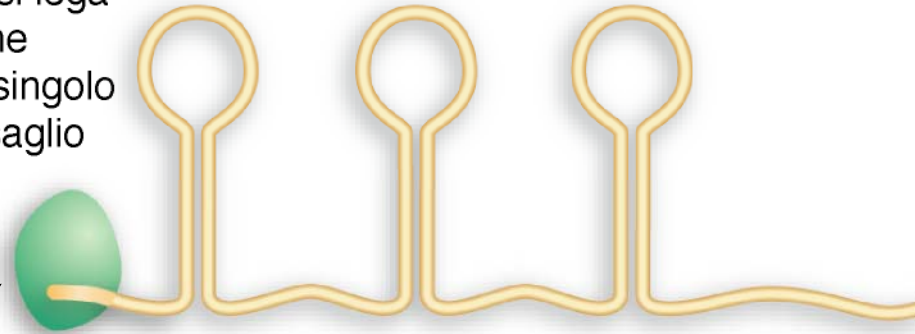


L'RNA regolatore si lega all'RNA bersaglio

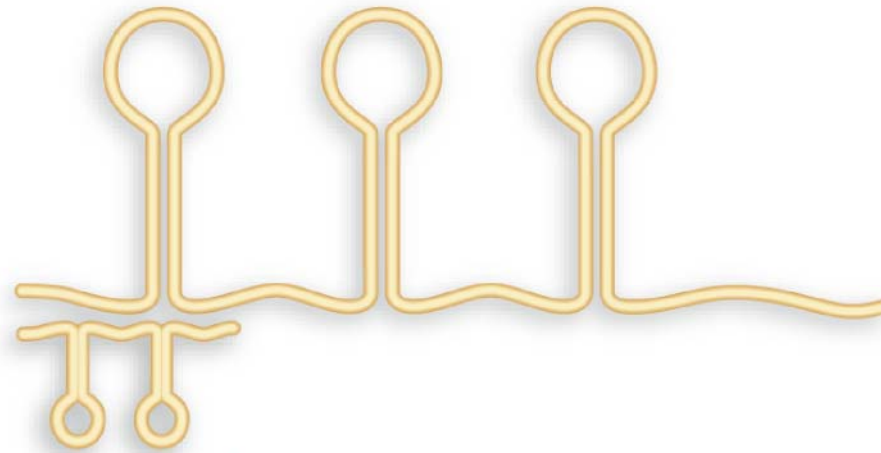


Il regolatore esclude il legame della proteina

La proteina si lega
a una regione
a filamento singolo
del suo bersaglio



Es. eIFs



La proteina non può legarsi
al bersaglio

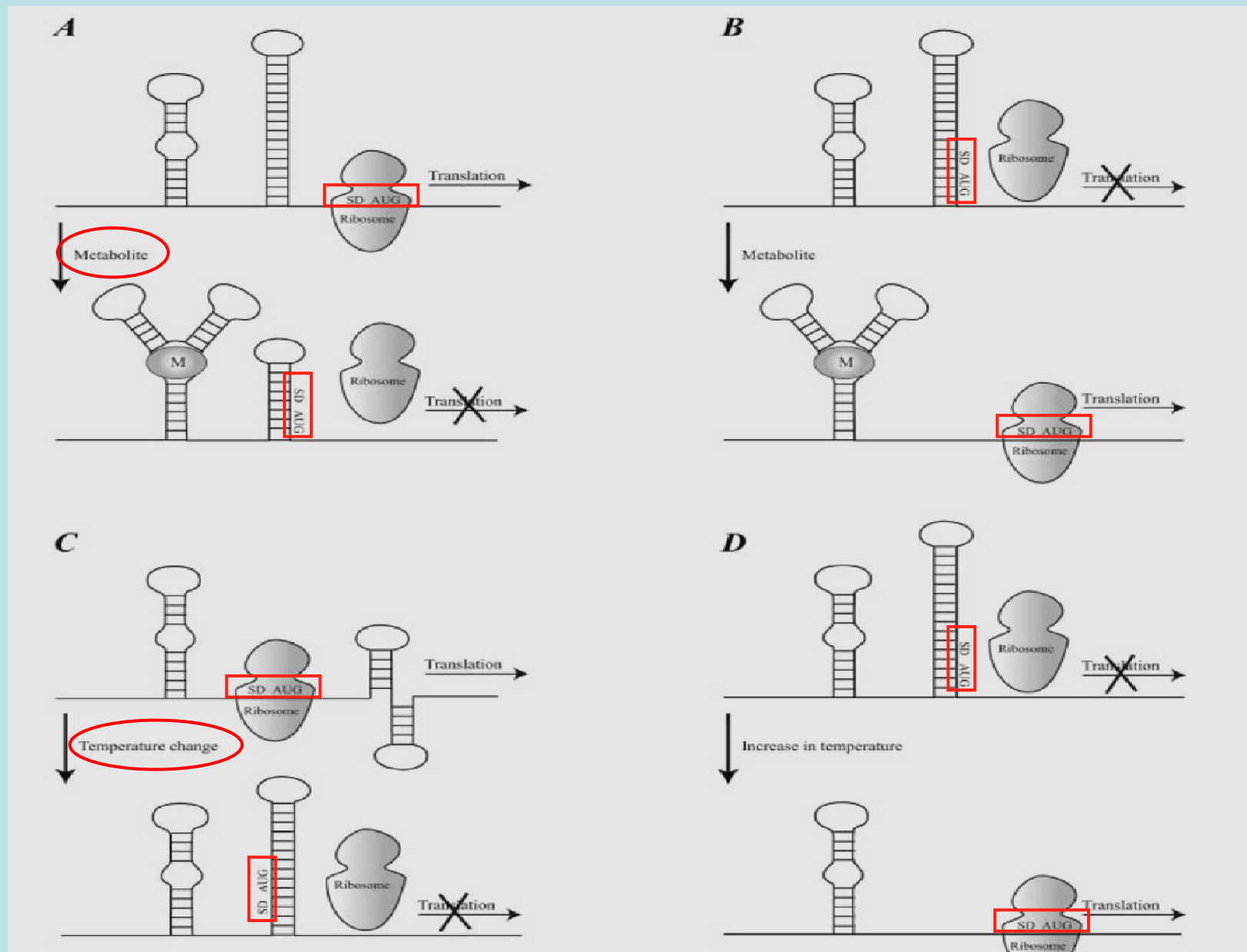


FIG. 13. Examples of translational regulation mechanisms. (A) Repression of translation by binding of a metabolite that stabilizes an alternative mRNA secondary structure and leaves the SD sequence and initiation codon (AUG) in a base-paired region. (B) Activation of translation by binding of a metabolite that stabilizes an alternative mRNA secondary structure and leaves the SD sequence and initiation codon (AUG) in an unpaired region, thus providing ribosomal access. (C) Repression of translation by the formation of an alternative mRNA secondary structure as a result of a change in temperature. (D) Activation of translation by an increase in temperature, causing a local melting of the mRNA secondary structure covering the SD and AUG region.

I repressori traduzionali si legano all'mRNA

Repressore	Gene bersaglio	Sito di azione
Proteina del rivestimento di R17	Replicasi di R17	Forcina che comprende il sito di legame del ribosoma
RegA di T4	mRNA precoci di T4	Varie sequenze che comprendono il codone di inizio
DNA polimerasi di T4	DNA polimerasi di T4	Sequenza Shine-Dalgarno
p32 di T4	Gene 32	Leader 5' a filamento singolo