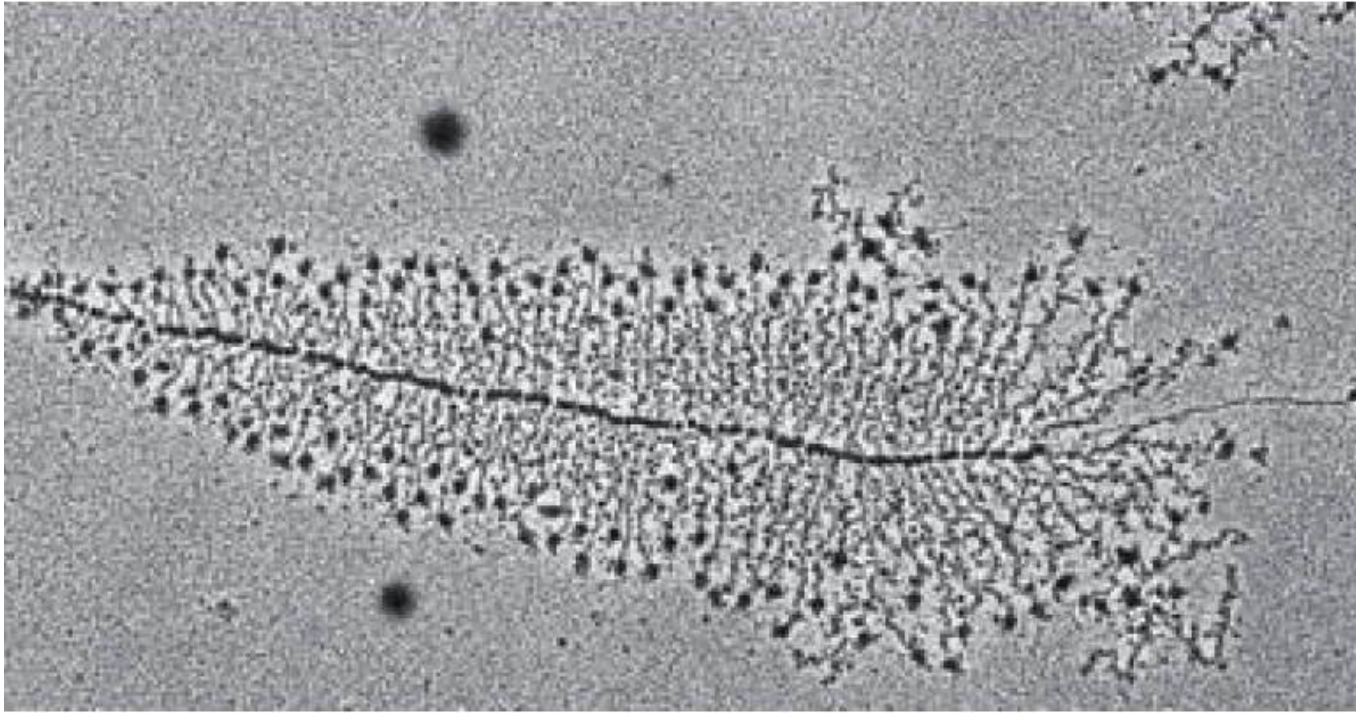


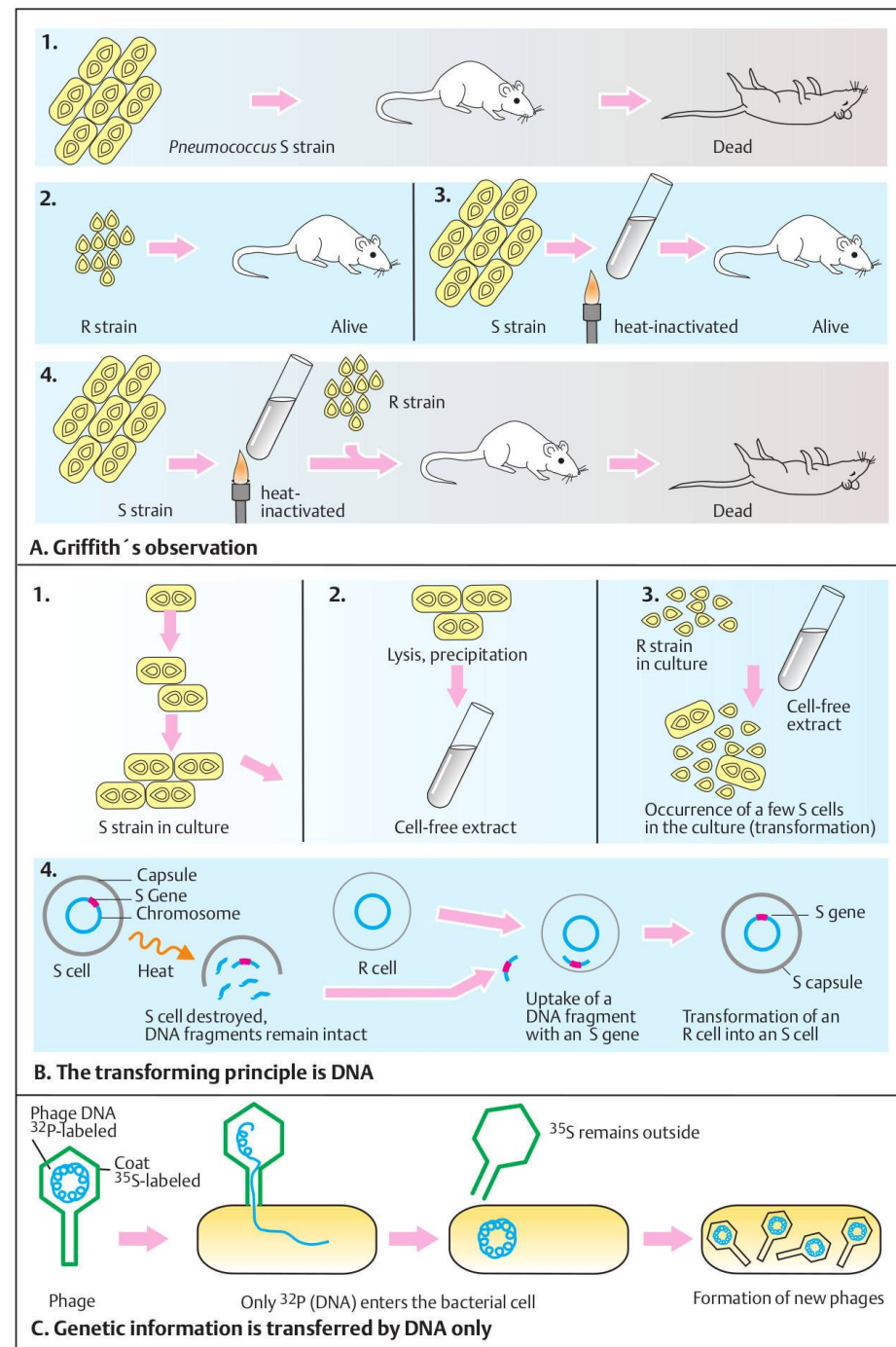
G 01 :

Organisation et expression du genome



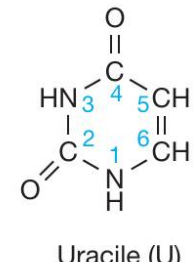
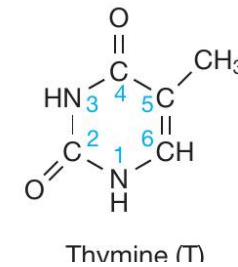
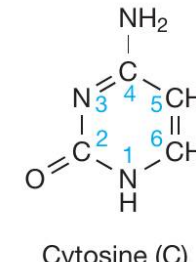
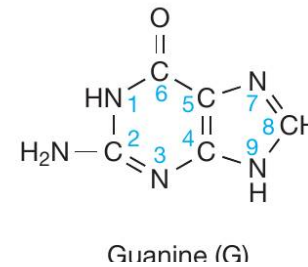
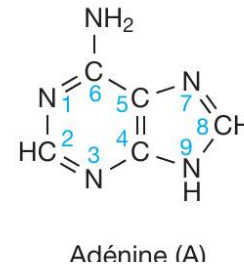
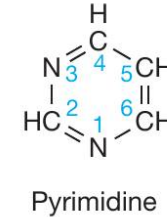
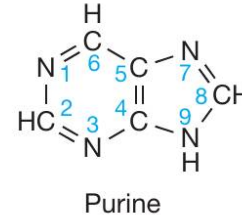
Introduction

- Définition genome
- Darwin
- Mendel
- Morgan & Sutton
- Griffith
- Avery, MacLeod & McCarthy
- Hershey & Chase



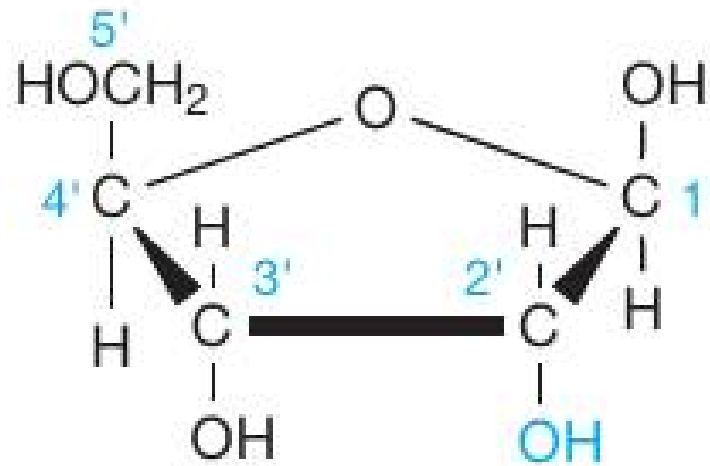
Bases azotées

- Pyrimidine
 - Cytosine, Thymine, Uracile
- Purine
 - Adénine, Guanine

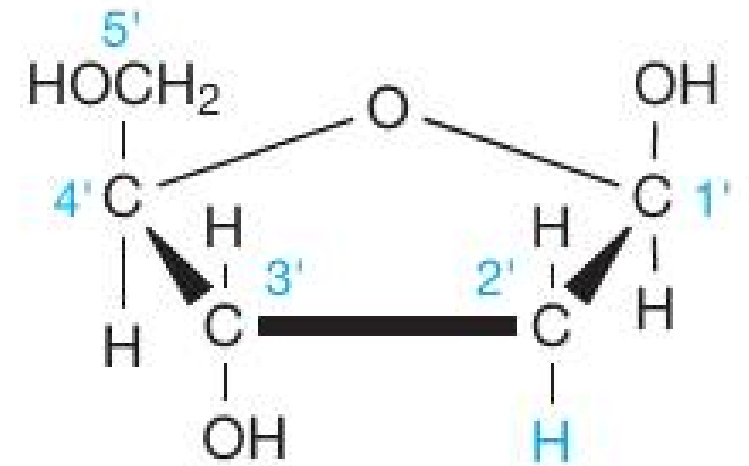


Sucre

- Pentose
- Ribose
- Desoxyribose
- Numérotation de carbones



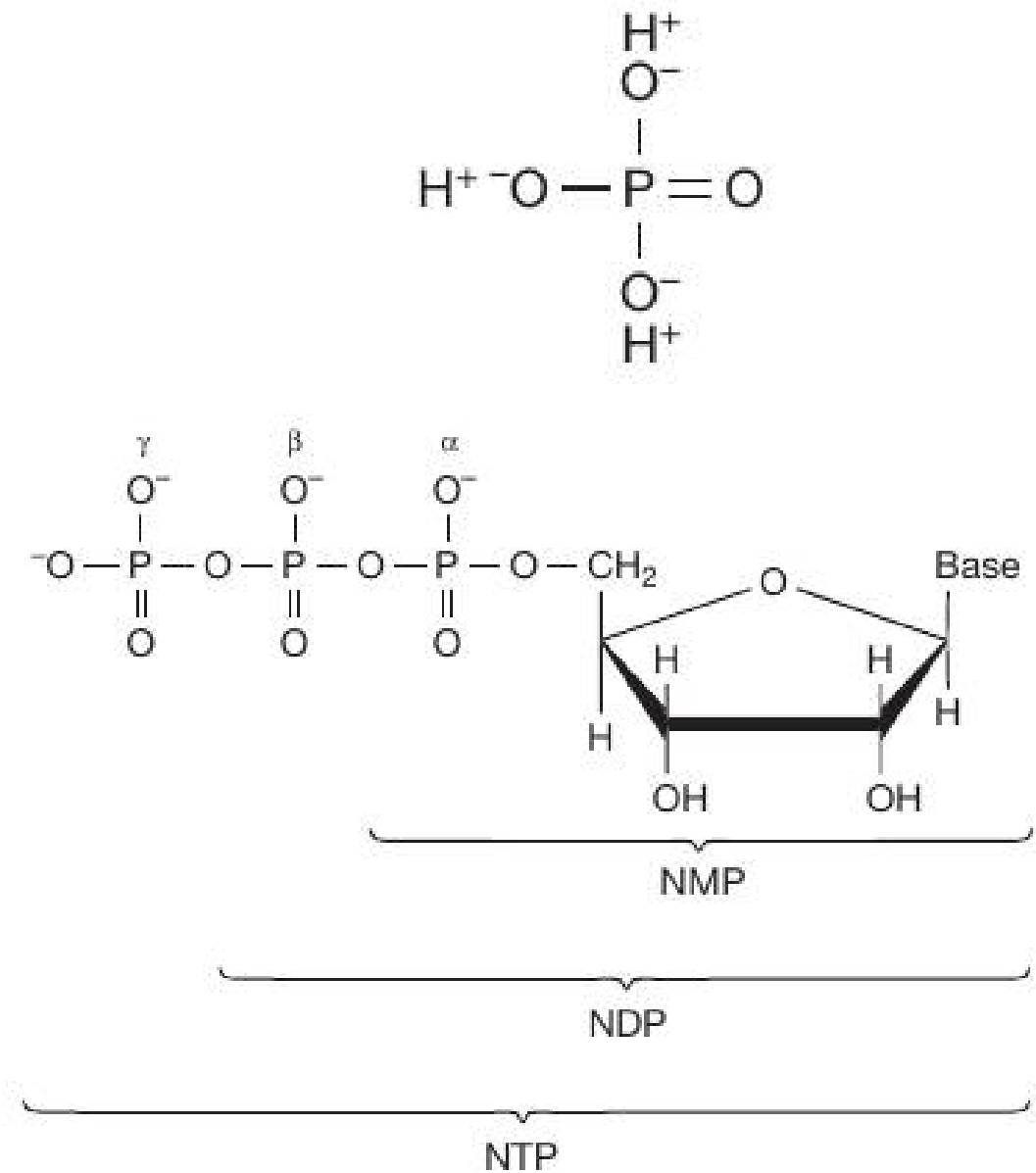
Ribose



Désoxyribose

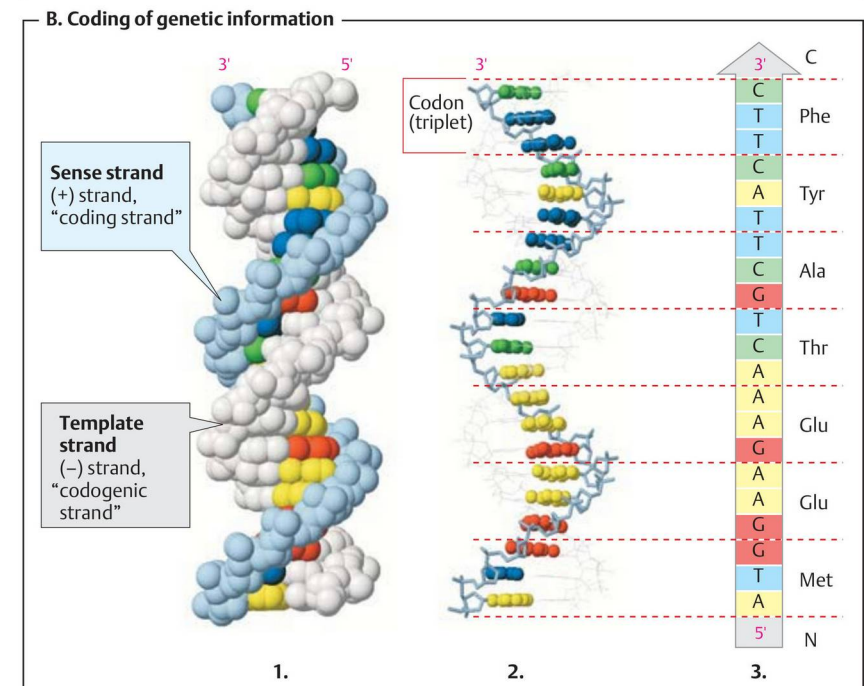
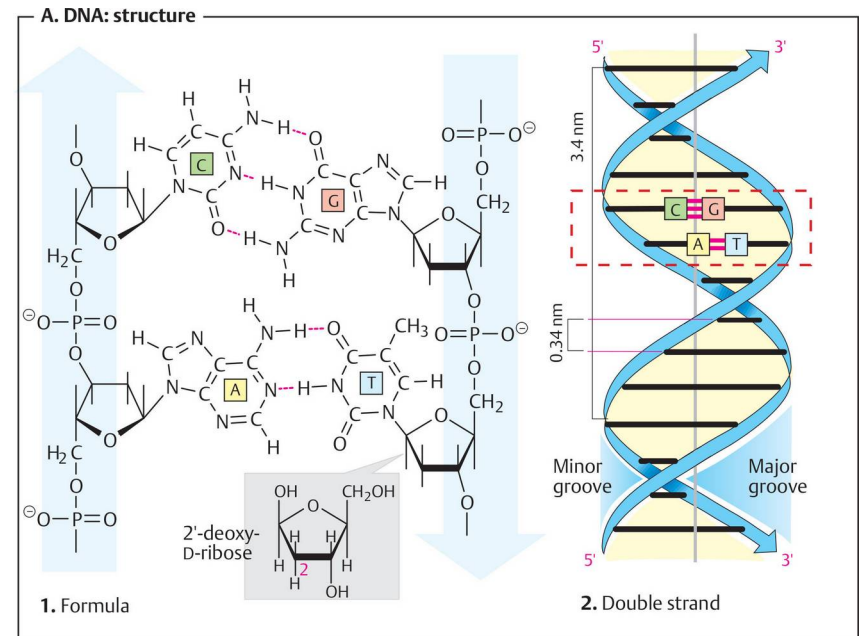
Phosphate

- Carbone 5'
- NMP
- NDP
- NTP
 - ATP, GTP



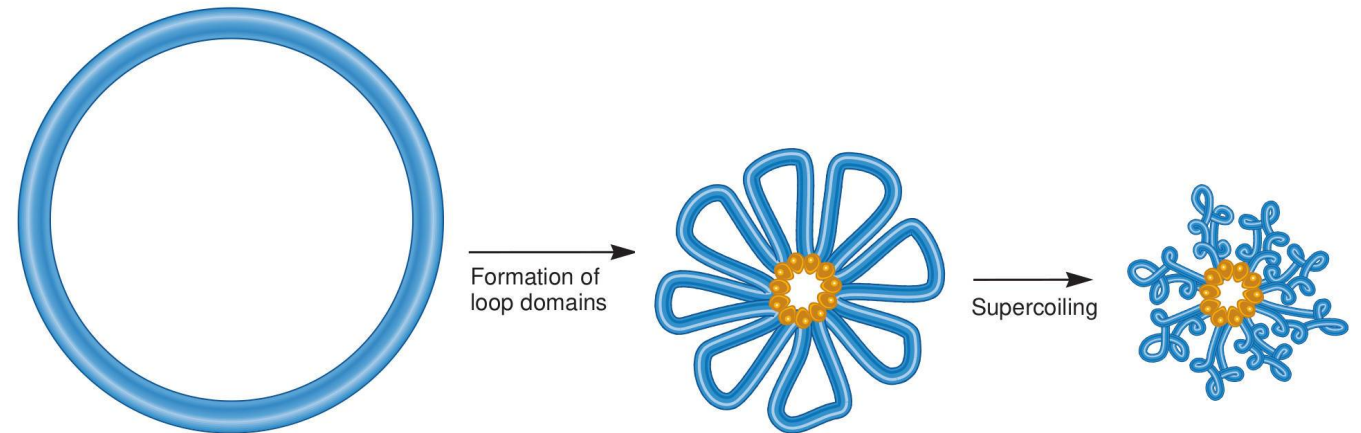
ADN

- Watson, Crick, Wilkins, Franklin
- Double hélice
- Brins antiparallèles
- Complémentarité des bases
- Montants et barreaux d'échelles



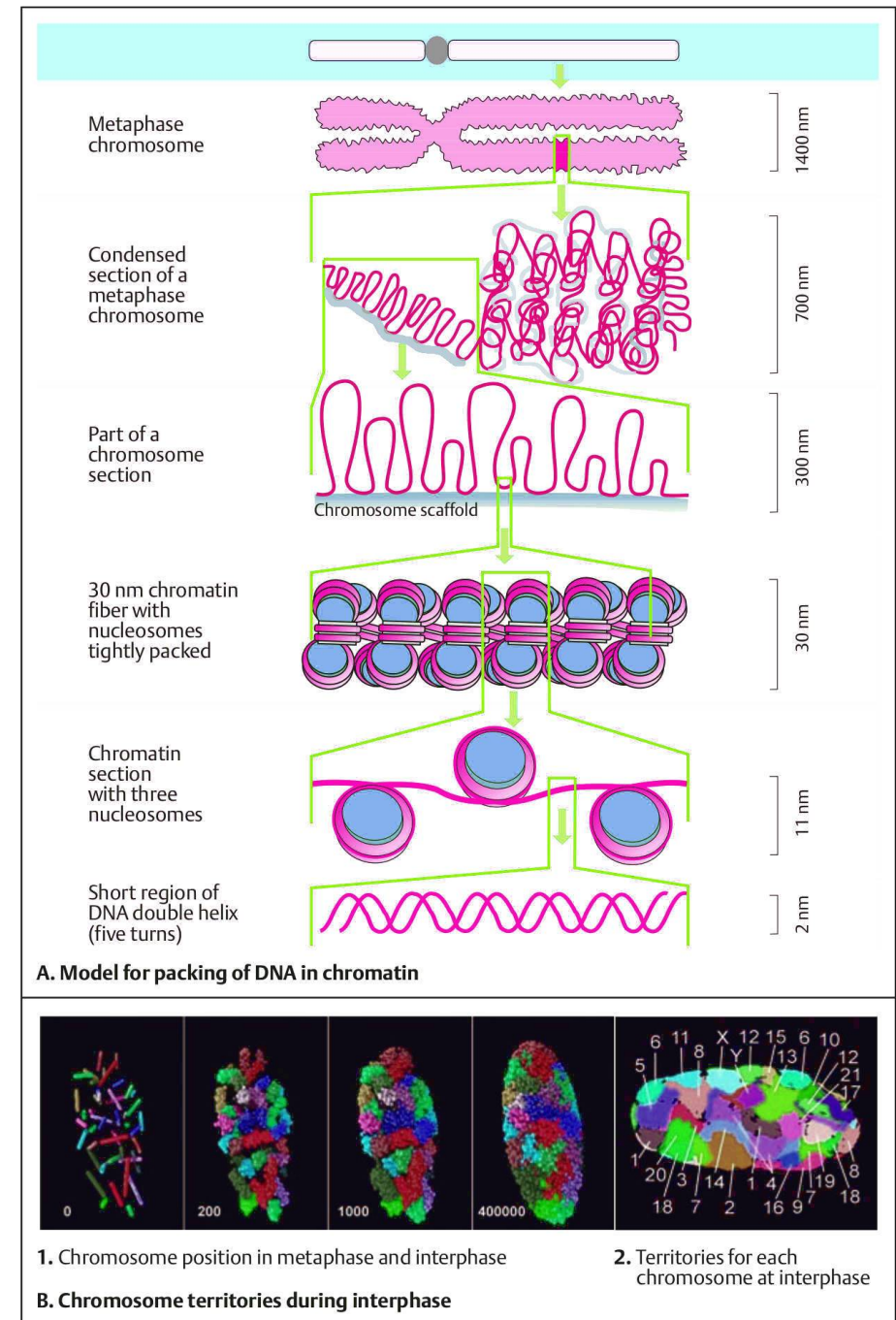
Chromosome bactérien

- Circulaire
- Double hélice
- Nucléotide
- Protéine H-NS



Chromosome eucaryote

- Chromosome interphasique
- Euchromatine VS Hétérochromatine
- Fibre nucléosomique
- Histones
- Fibre de 30nm
- Charpente protéique



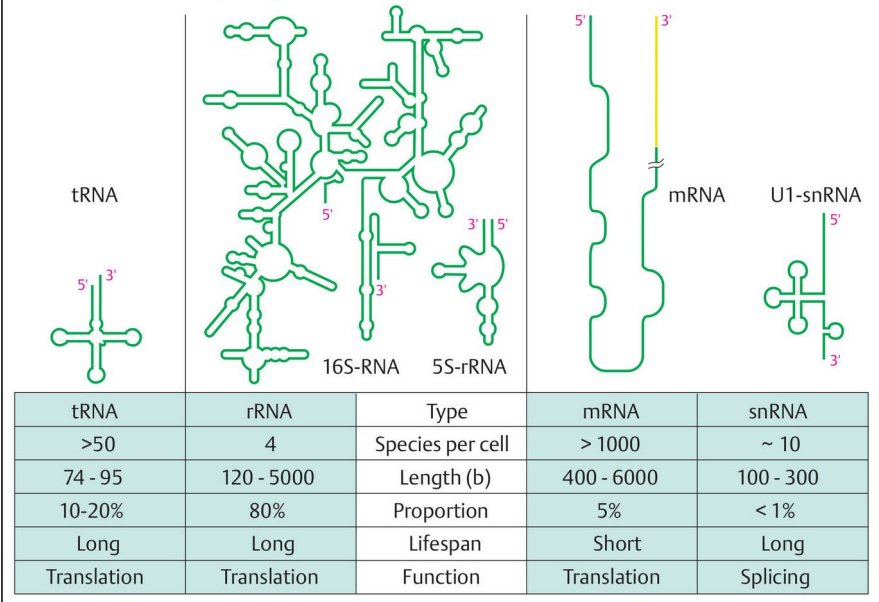
Localisation de l'ADN

- Bactéries
 - Chromosome bactérien
 - Plasmides
- Eucaryotes
 - Noyau
 - Organites semi-autonomes

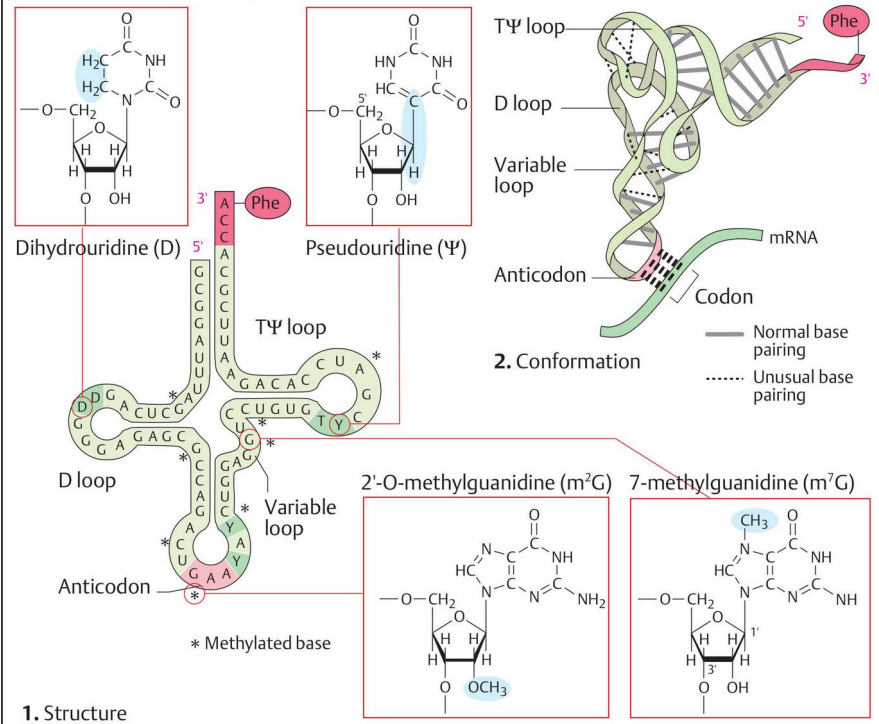
ARN

- ARNm
- ARNr
- ARNt

A. Ribonucleic acids (RNAs)



B. Transfer RNA (tRNA^{Phe})

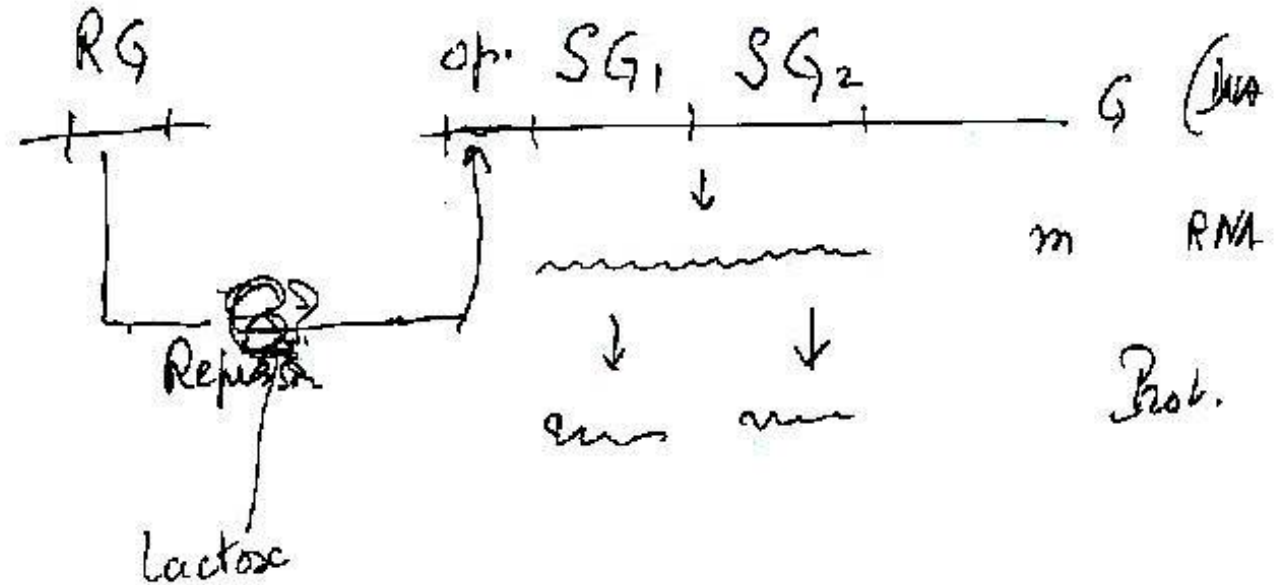


Gene

- Séquence codante = ORF
- Promoteur
- 5'UTR
- 3'UTR
- Séquences régulatrices
 - Enhancer
 - Silencer
 - Insulator

Operon

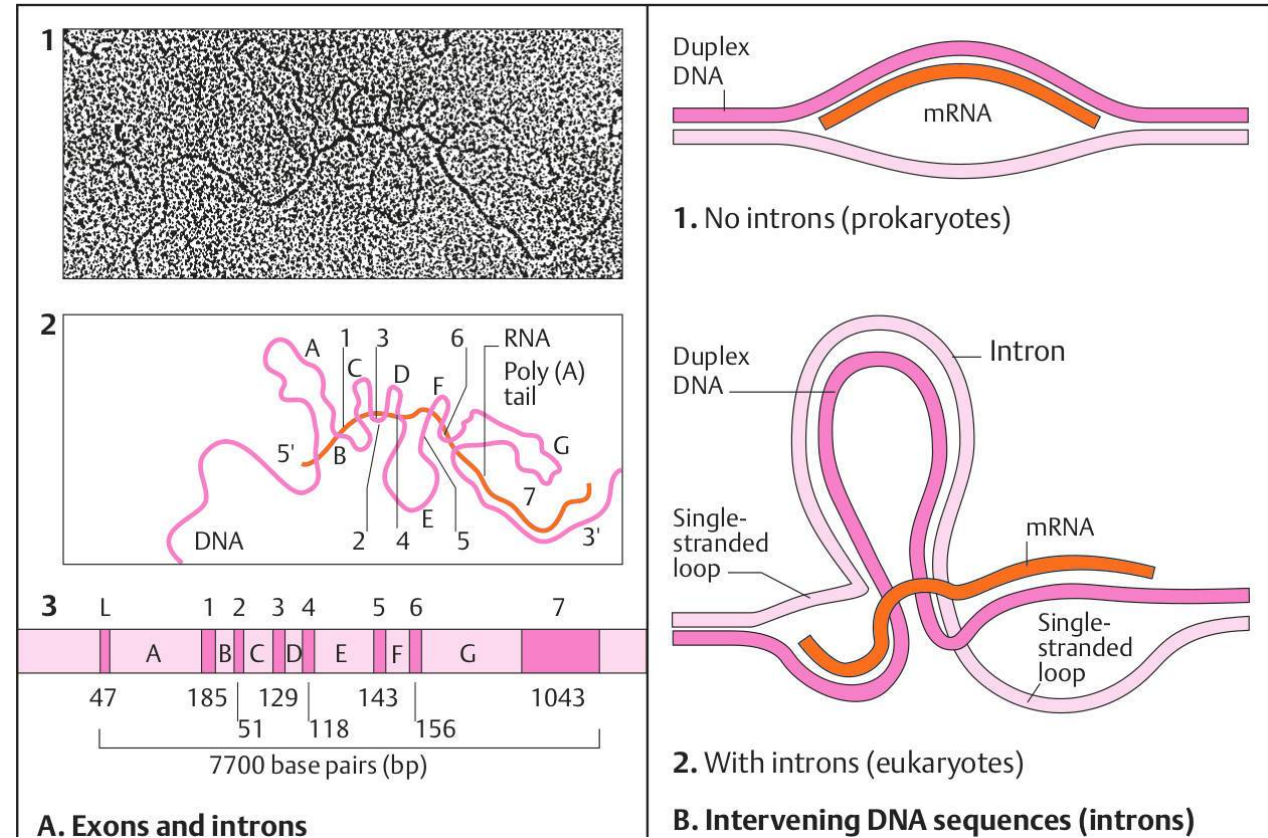
- Jacob & Monod
- Promoteur
- Operateur
- Plusieurs ORF
 - Exemple operon lactose
- ARN polycistronique



Francis Jacob

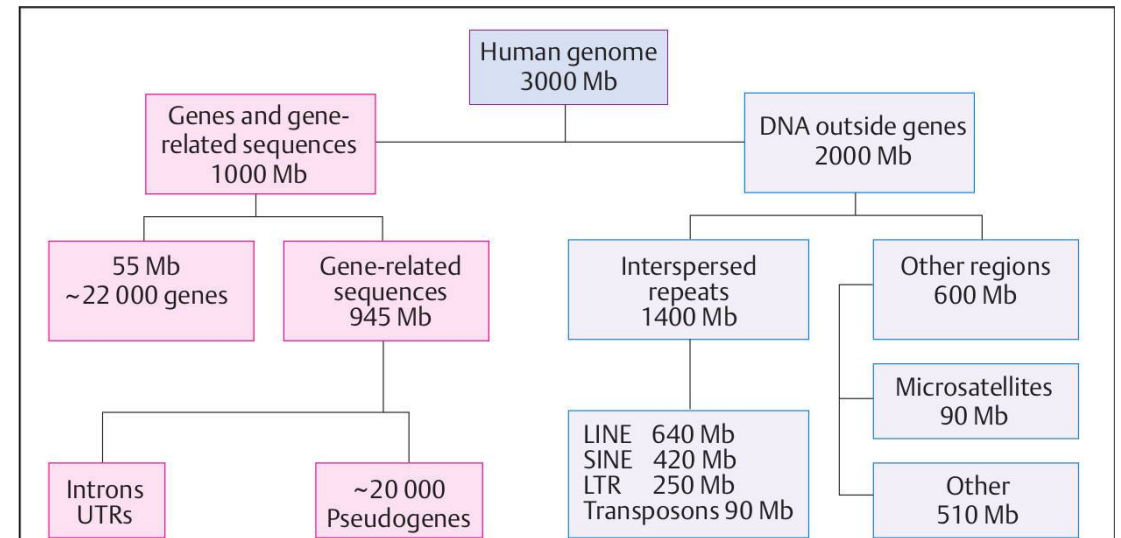
Genome eucaryote

- Gène eucaryote unique
 - Morcelé
 - Intron & Exon
- Séquences hautement répétées = satellite
- Séquence moyennement répétées
 - Gènes en tandem
 - Famille multigénique
 - Éléments génétiques mobiles



Genome eucaryote

- Gène eucaryote unique
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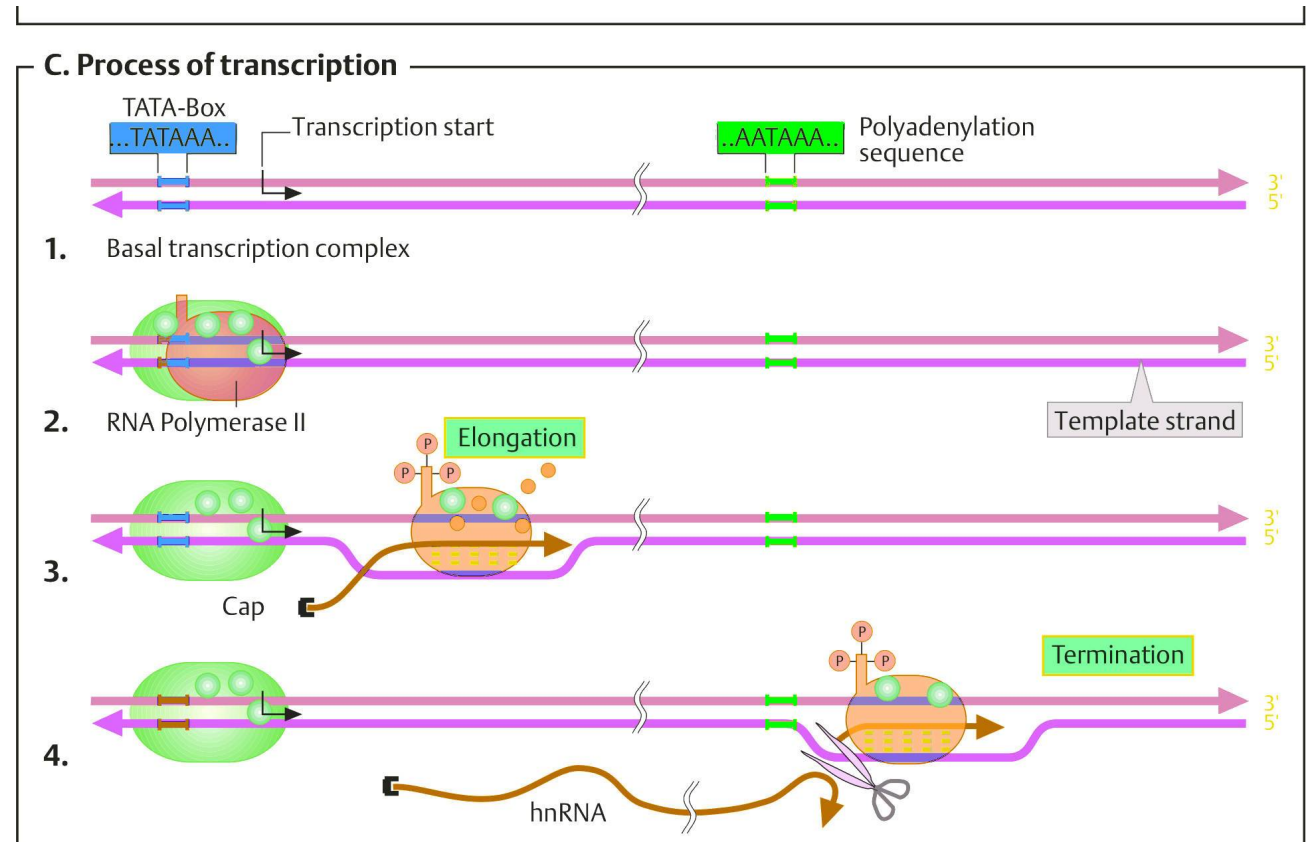
A. The types of sequence in the nuclear human genome

1. Long interspersed elements (LINE) (autonomous)		LINE-1 LINE-2 LINE-3	~600,000 copies ~370,000 copies ~44,000 copies
2. Short interspersed elements (SINE) (non-autonomous)		Alu family MIR MIR3	~1,200,000 copies ~450,000 copies ~85,000 copies
3. Retrovirus-like elements (LTR transposons)		Human endogenous retroviral sequences (HERV) (several classes; autonomous and non-autonomous)	~240,000 copies
4. DNA transposons		Several classes (autonomous and non-autonomous)	~300,000 copies

B. Major types of interspersed repetitive DNA in the human genome

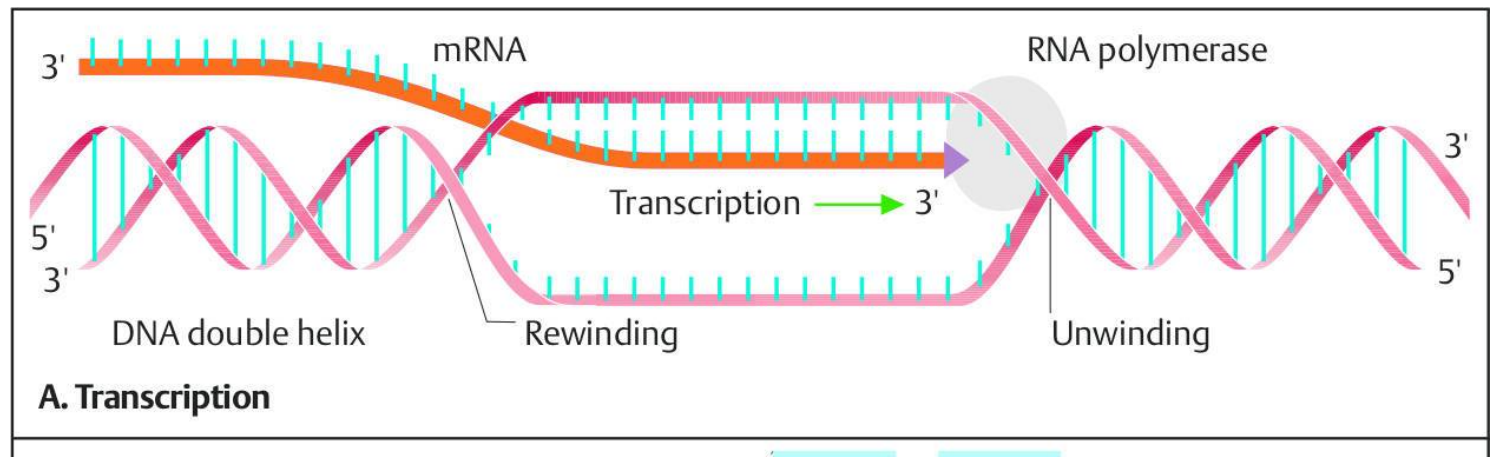
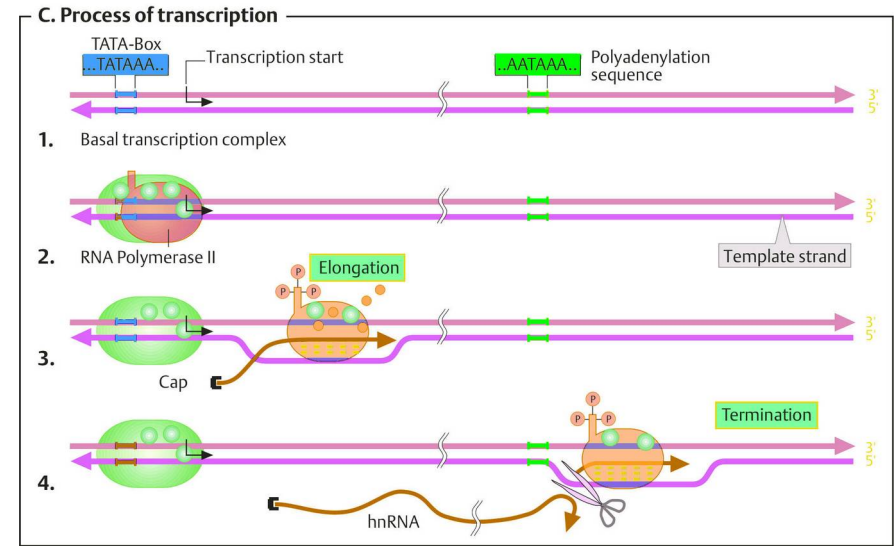
Transcription

- ARN Pol II
- Promoteur
 - TATA box
 - TBP
 - (CAAT)
 - TFII
 - Phosphorylation



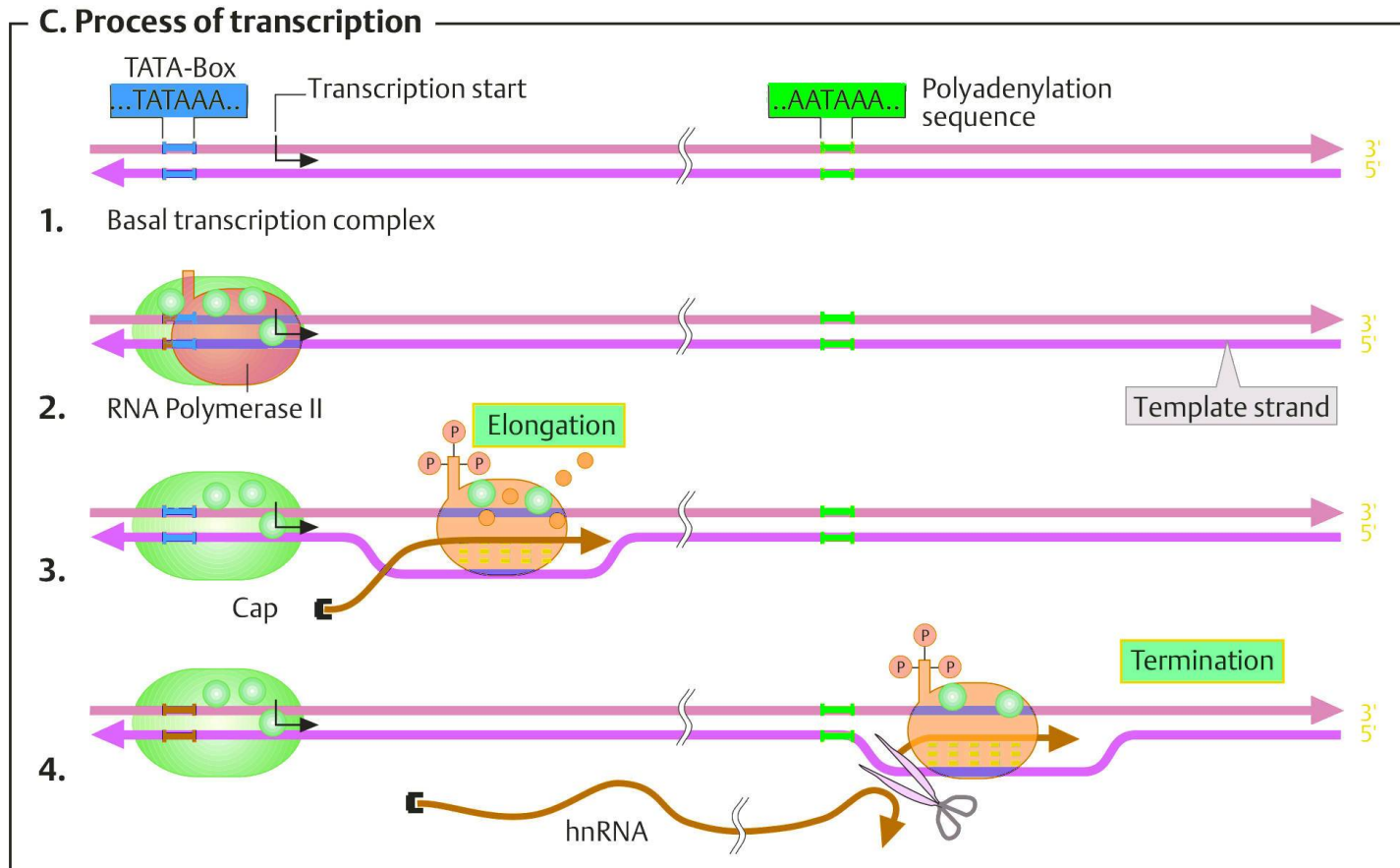
Elongation

- ARN Pol II
- Sens 5' → 3'
- 50nt/s
- 1 erreur/10 000nt
- Heteroduplex



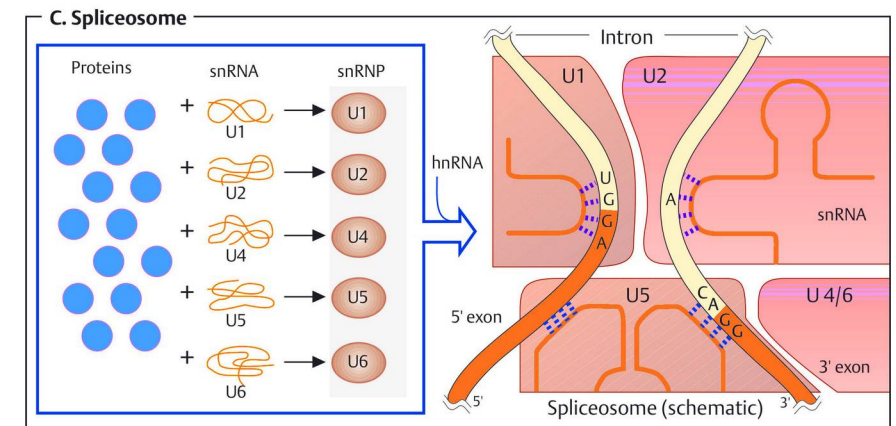
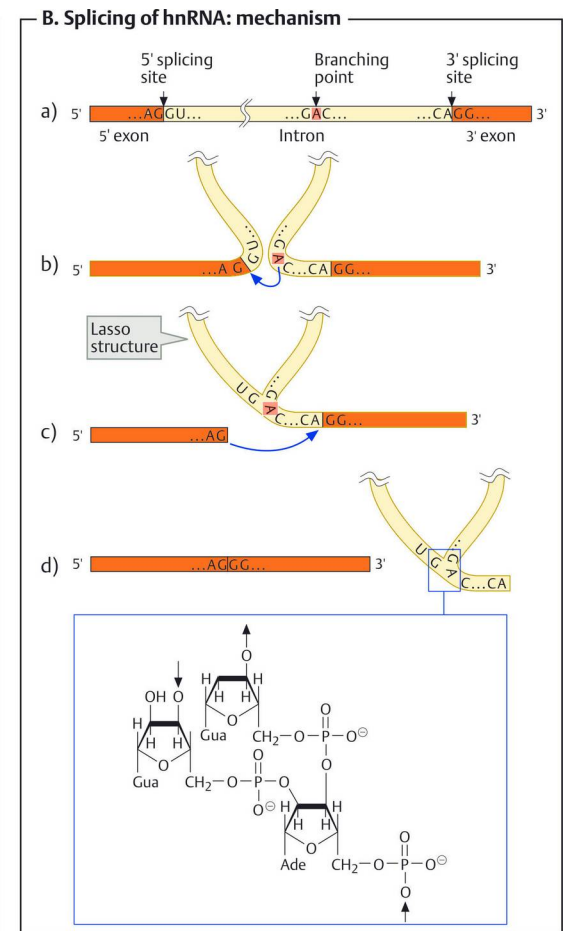
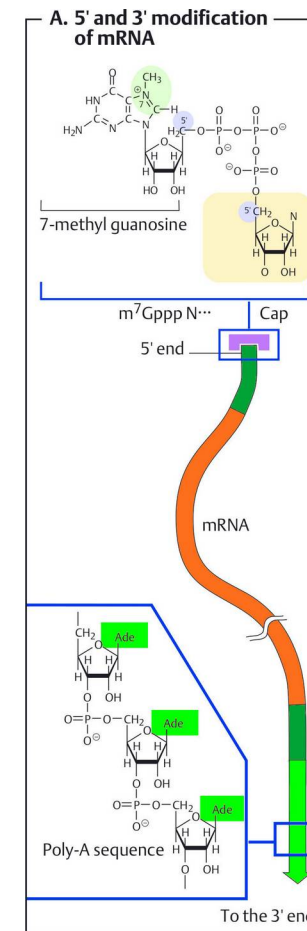
Terminaison

- Signal de polyadénylation
 - AAUAAA
 - CPSF
- Modèle torpille
- Modèle allostérique



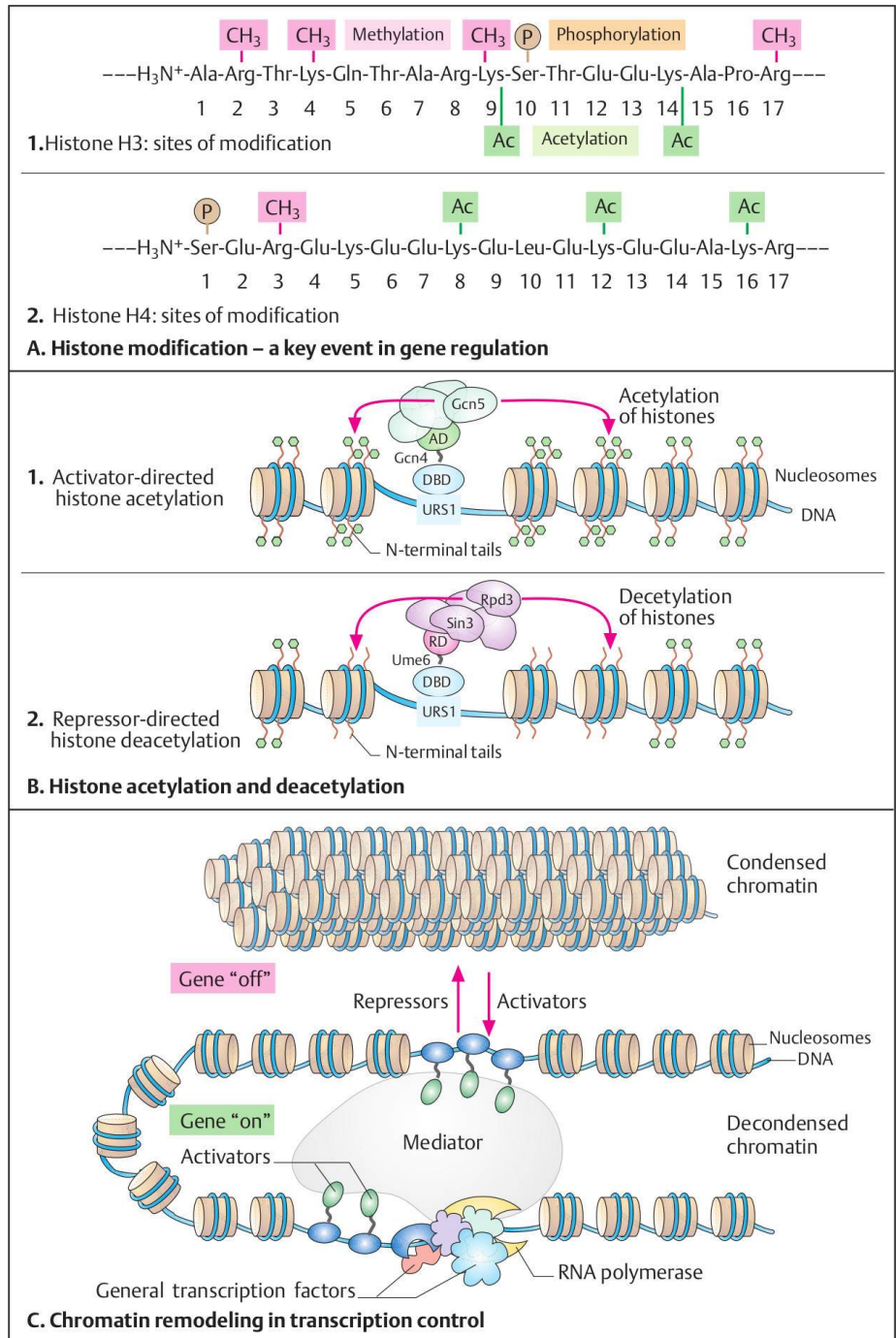
Maturation

- Coiffe de 7mG
- Polyadénylation
- Epissage



Controle

- Facteurs de transcription
- Hétérochromatine facultative
- Protéine de remodelage de la chromatine
- « Code histone »
- Méthylation de l'ADN



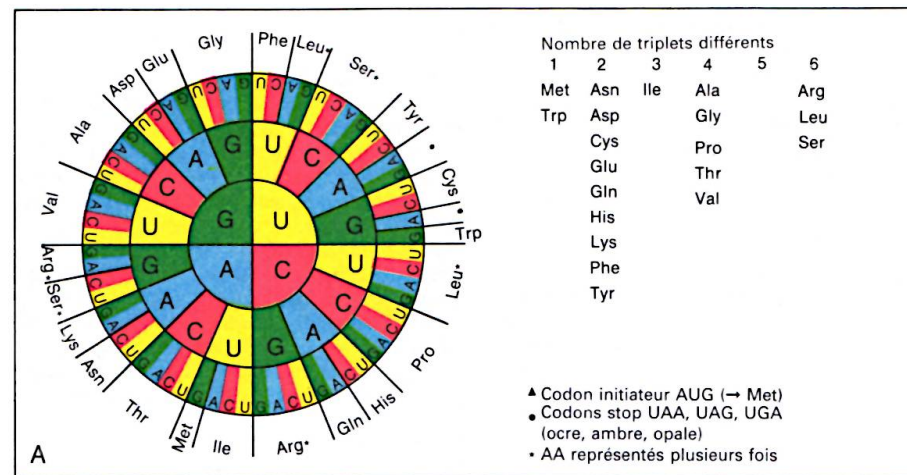
Code génétique

- 64 codons
- Codons stop
- Codon initiateur
- Code redondant ou dégénéré
- Code non-ambigu

nonpolar polar basic acidic (stop codon)

Standard genetic code

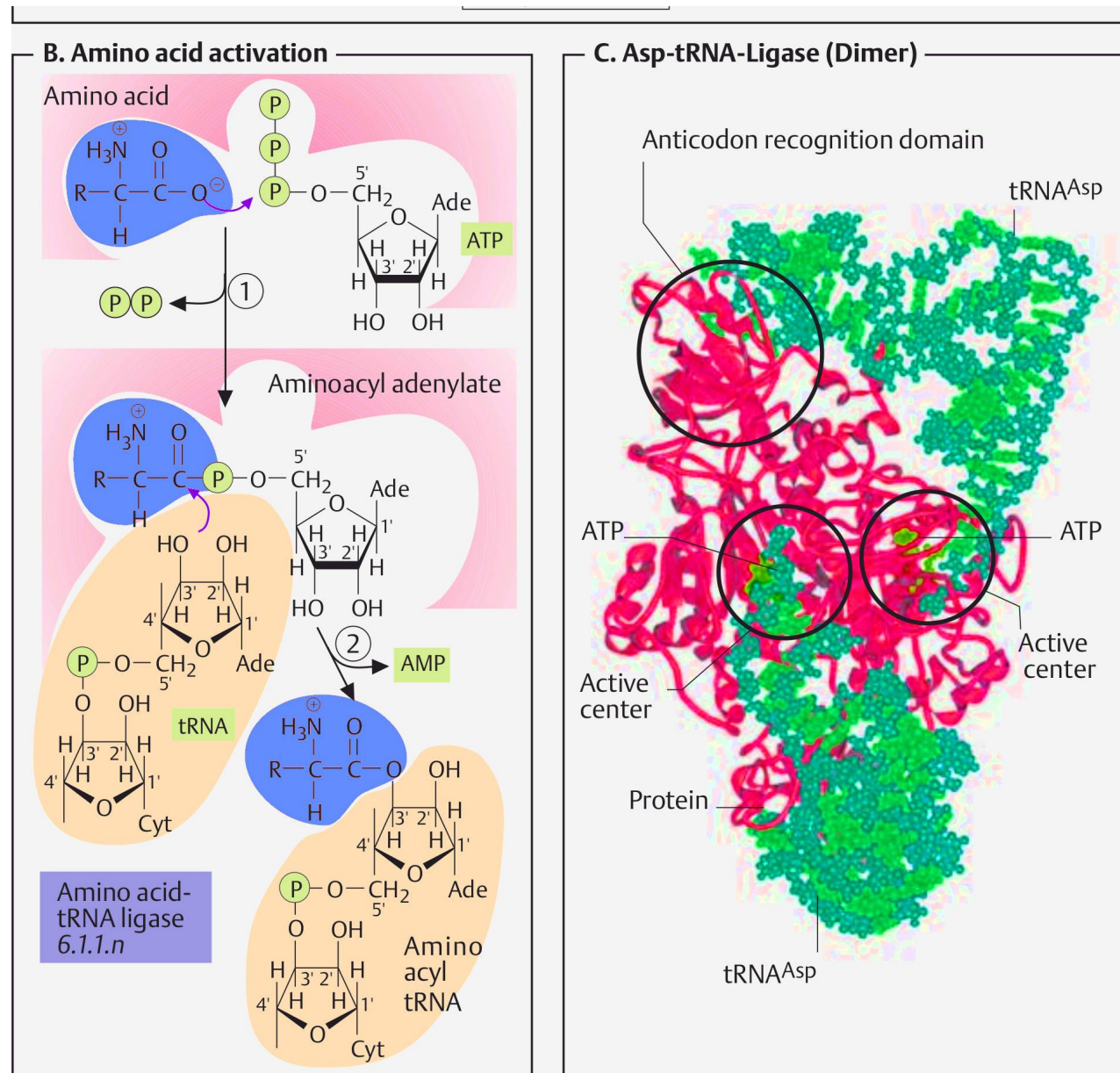
1st base	2nd base				3rd base
	U	C	A	G	
U	UUU (Phe/F) Phenylalanine	UCU (Ser/S) Serine	UAU (Tyr/Y) Tyrosine	UGU (Cys/C) Cysteine	U
	UUC	UCC	UAC	UGC	C
	UUA	UCA	UAA Stop (Ochre)	UGA Stop (Opal)	A
	UUG	UCG	UAG Stop (Amber)	UGG (Trp/W) Tryptophan	G
C	CUU (Leu/L) Leucine	CCU (Pro/P) Proline	CAU (His/H) Histidine	CGU (Arg/R) Arginine	U
	CUC	CCC	CAC	CGC	C
	CUA	CCA	CAA (Gln/Q) Glutamine	CGA	A
	CUG	CCG	CAG	CGG	G
A	AUU (Ile/I) Isoleucine	ACU (Thr/T) Threonine	AAU (Asn/N) Asparagine	AGU (Ser/S) Serine	U
	AUC	ACC	AAC	AGC	C
	AUA	ACA	AAA (Lys/K) Lysine	AGA (Arg/R) Arginine	A
	AUG ^[A] (Met/M) Methionine	ACG	AAG	AGG	G
G	GUU (Val/V) Valine	GCU (Ala/A) Alanine	GAU (Asp/D) Aspartic acid	GGU (Gly/G) Glycine	U
	GUC	GCC	GAC	GGC	C
	GUA	GCA	GAA (Glu/E) Glutamic acid	GGA	A
	GUG	GCG	GAG	GGG	G



Nomenclature des bases déterminant le code

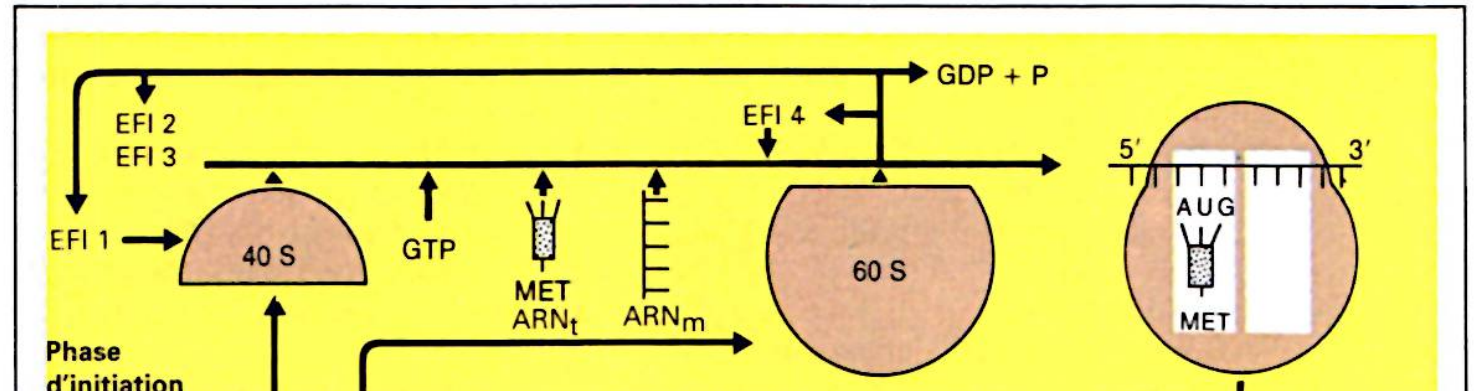
Charge de l'ARNt

- 2 étapes
- Wobble sur la 3ème base
- Double sélectivité



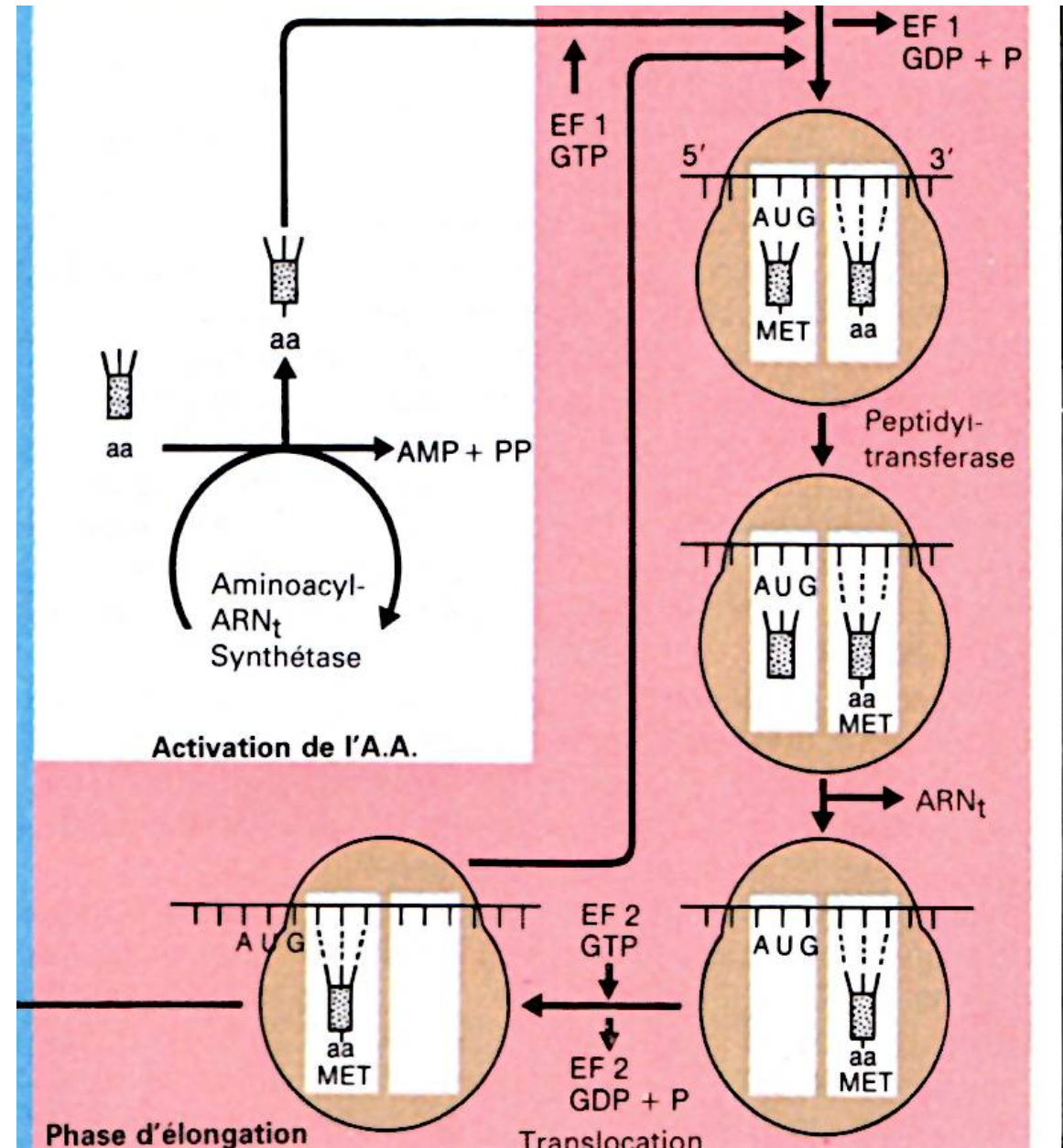
Initiation de la traduction

- IF-GTP+ARN_t-met
- Fixation à la 40S dans le site P
- Fixation à l'ARN_m
- Scan → AUG Kozak
- Fixation de la 60S



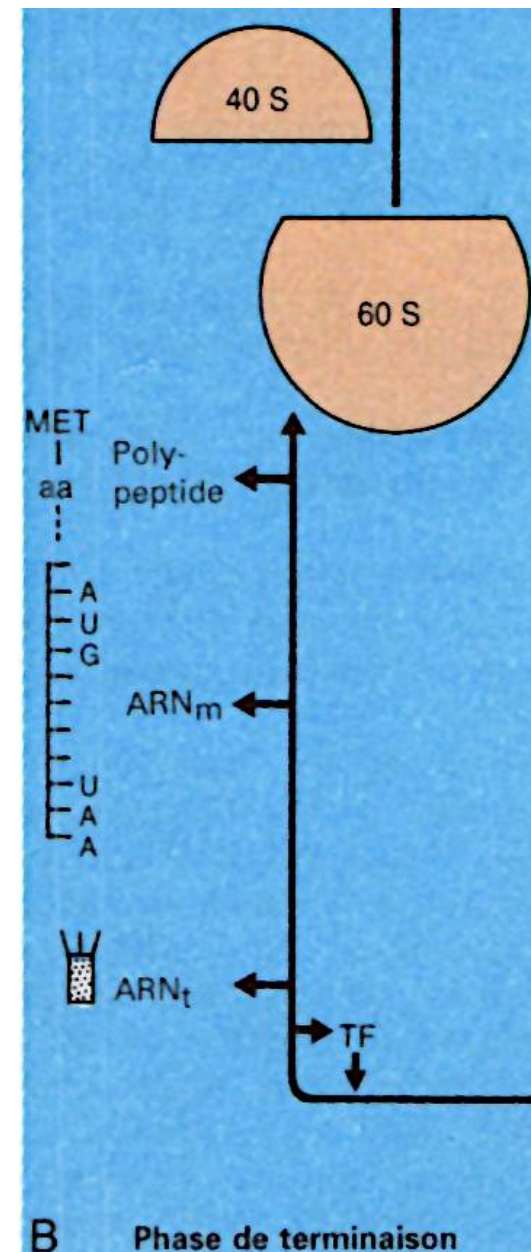
Elongation

- Arrivée de l'ARNt-AA dans le site A avec un EF-GTP
- Décodage : hydrolyse du GTP
- Activité peptidyl-transferase
- Un EF-GTP déplace le ribosome



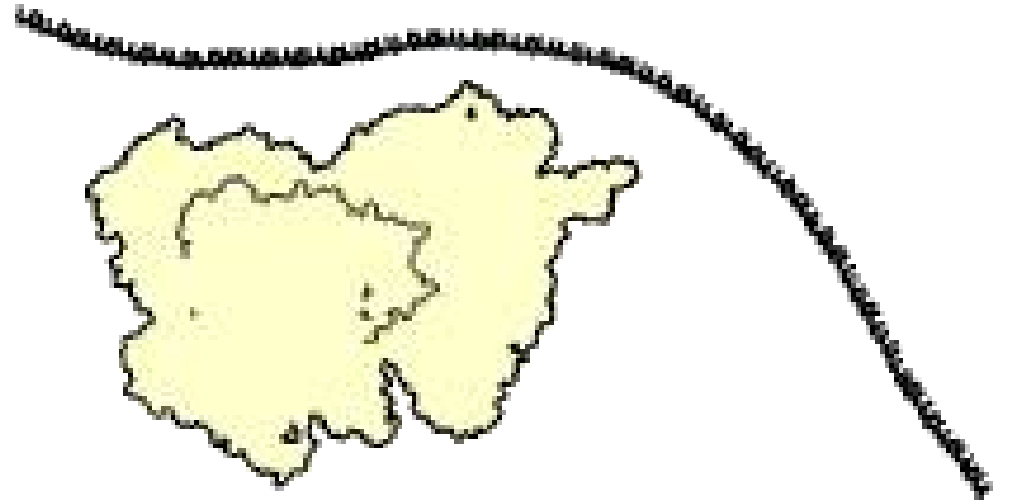
Terminaison

- Codon Stop
- Facteur de terminaison



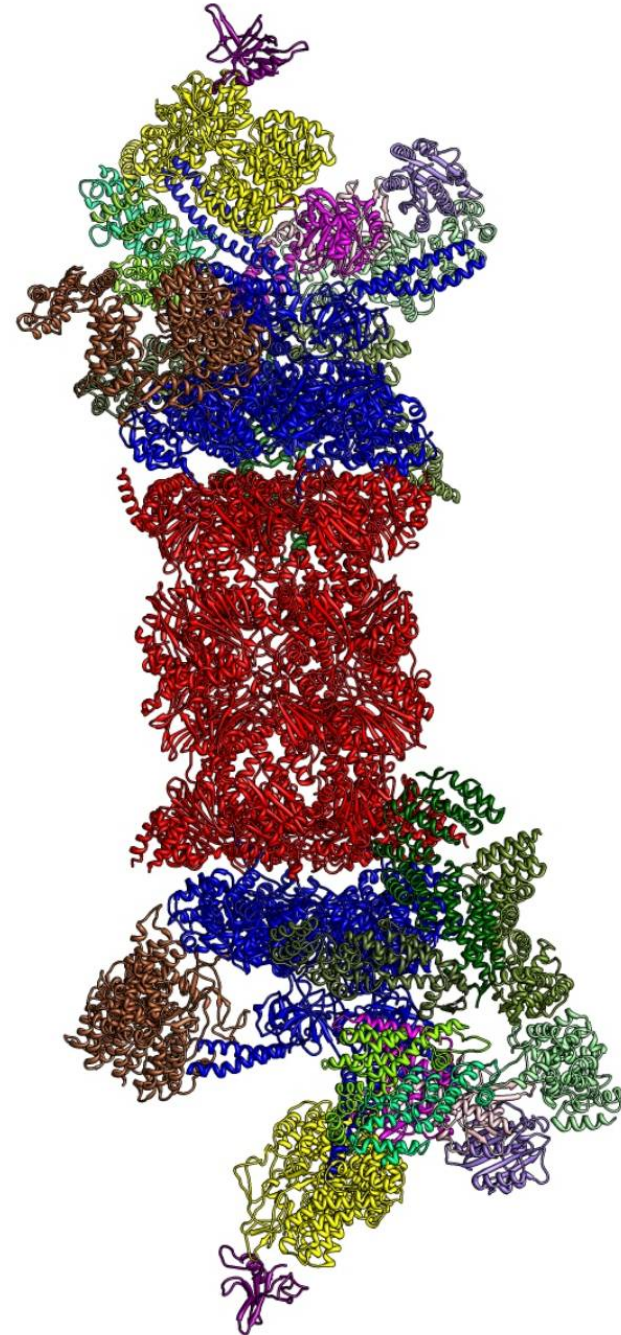
Adressage

- Co-traductionnel au REG
 - SRP
 - Translocon
- Post-traductionnel
 - Mitochondrie
 - Chloroplaste



Proteasome

- Ubiquitination
- Proteasome



Controle

- ARNi
- Drosha
- Dicer
- RISC

