

Insegnamento: Laboratorio Biologia Molecolare

Docenti:



Prof. SCHOEFTNER STEFAN, responsible corso – Lecture



Prof.ssa BANDIERAANTONELLA – Laboratory course

The course provides theoretical and practical training on techniques and experimental approaches in molecular biology.

- A focus will be set on the molecular biology and technologies related to of nucleic acids
- Basic techniques for DNA manipulation, gene study, gene cloning, gene expression analysis and recombinant DNA technology will be addressed; genome browser search.
- Laboratory exercises include the teaching of <u>laboratory safety standards</u> the handling of <u>laboratory instruments</u>, the <u>extraction of DNA</u> from bacteria and human cheak cells, <u>use of</u> <u>restirction enzymes</u>, <u>mapping of plasmids after digest by restriction digest</u>, gel <u>electrophoresis</u>, <u>amplificiation of nucleic acid sequences by PCR</u>, <u>mapping of polymorphisms in "student</u> <u>population"(Alu repeats, disease related SNPs)</u>, <u>analysis and interpretation of results in</u> <u>student polylation and overall population</u>.

Organization Course: Laboratorio Biologia Molecolare

Monday 30.09.2019 – 28.10.2019: 11 hours theroretical lections: Prof. Schoeftner (Technologies)

Monday 04.11.2019 – 16.12.2019: Theroretical lections: Prof. Schoeftner (Technologies) Laboratory course: Prof. Scaggiante

	lunedi 4 novembre	martedi 5 novembre	mercoledi 6 novembre	giovedi 7 novembre	venerdi 8 novembre	
9-10	Alimenti, nutrienti e salute	Alimenti, nutrienti e salute	Fisiologia	Lab. Biol. Bol. II TURNO - Bandiera	Lab. Biol. Bol. IV TURNO -	Fisiologia vegetale
10-11	Alimenti, nutrienti e salute	Alimenti, nutrienti e salute	Fisiologia	Lab. Biol. Bol. II TURNO - Bandiera	Lab. Biol. Bol. IV TURNO -	Fisiologia vegetale
11-12	Lab. di Biologia Molecolare (TEORIA)	Microbiologia	Farmacologia	Lab. Biol. Bol. II TURNO - Bandiera	Lab. Biol. Bol. IV TURNO -	Ecologia
12-13	Lab. di Biologia Molecolare (TEORIA)	Microbiologia	Farmacologia	Lab. Biol. Bol. II TURNO - Bandiera	Lab. Biol. Bol. IV TURNO -	Ecologia
13-14			Lab. Biochimica III e IV tumo			
14-15	Microbiologia	Farmacologia	Lab. Biol. Bol. I TURNO - Bandiera	Lab. Biol. Bol. III TURNO - Ecologia		
15-16	Microbiologia	Farmacologia	Lab. Biol. Bol. I TURNO - Bandiera	Lab. Biol. Bol. III TURNO - Ecologia		
16-17	Fisiologia	Ecologia	Lab. Biol. Bol. I TURNO - Bandiera	Lab. Biol. Bol. III TURNO - vegetale		
17-18	Fisiologia	Fisiologia vegetale	Lab. Biol. Bol. I TURNO - Bandiera	Lab. Biol. Bol. Fisiologia III TURNO - vegetale		

Monday Lecture: ca. 1,5 hours: Theoretical lecture (Schoeftner) – related to topic in weekly laboratory exercise

ca. 0,5 hours: Prof. Scaggiante – practical background to laboratory background

Labortory exercises: Turno I – IV; all "turni will do the same porgram in the respective week Prof. Bandiera + 2 experienced tutors (20-25 stundents per turno; 4-5 students per work station

Contents of Theoretical Lecture (Prof. Schoeftner)

1. Anatomy of the cell, biomolecules, concept of preparation of RNA/Protein/DNA.

2. Recombinant DNA techniques, Cloning vectors, endonucleases, artificial chromosomes, recombinant protein expression, introduction of genes into host-organisms.

3. DNA sequencing, bacterial immunity, manipulation of the genome content of pro- and eukaryotic organisms, siRNA/shRNA mediated knock-down approaches.

4. Hybridization related techniques (RNA-FISH, DNA-FISH, Southern blot, Northern blot), Electrophoresis, methods to study DNA:protein interaction (band shift, DNA footprinting, chromatin immunoprecipitation)

5. PCR technologies: standard PCR, RT—PCR and variants

6. Gene expression analysis: array technology and high content sequencing, determination of 3' and 5' ends of RNA, single molecule transcript analysis

7. Exercise session: Students get introduction into the use of the ENSEMBL genome browser and primer construction. Students will do primer design for practical part as homework.

Contents of Practical Course (Prof. Bandiera)

Application of molecular biology techniques for the diagnosis and **monitoring of specific genetic conditions (allelic variants) and genetic variation of Alu repeat in students of the course**.

- 1. THE MOLECULAR BIOLOGY LABORATORY: Rule of conduct and safety, hazardous reagents and material safety data sheet; equipment and lab instrumentation. The use of automatic lab pipettes for small volume manipulation.
- 2. PLASMIDS: Plasmid DNA extraction by a commercial kit, evaluation of extraction yield, preparation of samples for electrophoretic analysis. Plasmids will be subjected to control digest using restriction enzymes and will serve as positive and negative control for subsequent PCR.
- 3. PREPARATION OF GENOMIC DNA:
- 4. Anonymized preparation of genomic DNA from cheek cells of students and determination of concentration.
- 5. PCR AMPLIFICATION OF SITE OF GENETIC ALU REPEAT VARIANT: Alu repeats number variation on a locus of chromosome 16 will be determined by specific PCR. Agarose Gel electrophoresis will be used to monitor differences in Alu repeat number.

6. PCR AMPLIFICATION OF SITE OF ALLELIC G6PD VARIANTS:

A locus harboring allelic variants of the G6PD gene will be amplified by PCR. PCR products will be purified after gel electrophoresis and subjected to digest using restriction enzymes. Digested DNA will be separated by gel electrophoresis.

8. DATA ANALYSIS AND DISCUSSION:

Chi-square analysis will be used to to compare the Alu genotype frequencies within the class population with those predicted by the Hardy-Weinberg equation. The genotypic frequencies of the class population can also be compared with the genotypic frequencies of another population in the database.

Exam

\rightarrow 2 written exams:

Exam 1:

Reports on lab work at the end of each lab practice (Prof. Bandiera).

Reports will be evaluated assessing:

-diligence, attendance, presentation accuracy

-personal skills, synthesis, description and clarity in presentation, technical terms knowledge

-understanding degree, explanation and discussion skills, presence of conceptual errors.

 \rightarrow A total of 15 points can be reached.

 \rightarrow A minimum of 7,5 points is necessary to participate in the second part of the exam2

Exam 2

Learning progress on the theoretical lectures (Prof. Schoeftner) will be monitored in a written exam. Total points: 16.

Exam 2 consists of 12 multiple choice questions (0,5 points per question) and 2 "open questions" (5 points per question, max 1 page answer to question) on broader topics addressed during the theoretical lectures and virtual lab.

The final mark of the course results from the sum of both exams. Maximum points: 31 A minimum of 18 points is required to pass the exam "Laboratorio Biologia Molecolare".

CONTENTS LABORATORY COURSE

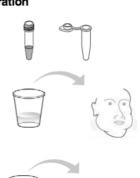
Turni: 5 Inscription in turni via Moodle federato in ca. 2 weeks!

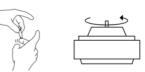
1. Preparation of Genomic DNA

Quick Guide

Lesson 1 Cheek Cell DNA Template Preparation

- Label one 1.5 ml micro test tube with your initials. Label one screwcap tube containing 200 µl of InstaGene matrix with your initials.
- Obtain a cup containing saline solution from your instructor. Pour the saline into your mouth and rinse vigorously for 30 seconds. Expel the saline back into the cup.
- Transfer 1 ml of your saline rinse into the micro test tube (NOT the screwcap tube) with your initials. If a P-1000 micropipet is not available, carefully pour ~1 ml of your saline rinse into your micro test tube (use the graduations on the side of the micro test tube to estimate 1 ml).
- 4. Spin your tube in a balanced centrifuge at full speed for 2 minutes. When the centrifuge has completely stopped, remove your tube. You should see a match-head sized pellet of whitish cells at the bottom of the tube. If you don't see a pellet of this size, decant the saline, refill your tube with more of your oral rinse, and repeat the spin.
- 5. After pelleting your cells, pour off the saline. Being careful not to lose your pellet, blot your tube briefly on a paper towel or tissue. It's OK for a small amount of saline (< 50 µl, about the same size as your pellet) to remain in the bottom of the tube.
- 6. Resuspend the pellet by vortexing or flicking the tube so that no clumps of cells remain.
- 7. Using a 2–20 μ l adjustable-volume micropipet set to 20 μ l, transfer all of your resuspended cells to the screwcap tube containing InstaGene.
- 8. Screw the cap tightly on the tube. Shake or vortex to mix the tube contents.



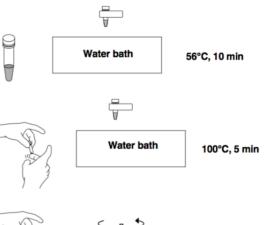


2-0

톺

Centrifuge

- 9. When all members of your team have collected their samples, place the tubes in the foam micro test-tube holder, and incubate at 56°C for 10 minutes in a water bath. At the halfway point (5 minutes), shake or vortex the tubes gently, then place back in the 56°C water bath for the remaining 5 minutes.
- Remove the tubes, shake or vortex, and place the tubes in a boiling water bath (100°C). Incubate at 100°C for 5 minutes.
- 11. Remove the tubes from the boiling water bath and shake or vortex the contents to resuspend. Pellet the matrix by spinning at 6,000 x g for 5 minutes (or 2,000 x g for 10 minutes) in a centrifuge.
- 12. Store your screwcap tube in the refrigerator until the next laboratory period (or proceed to step 2 of Lesson 2).

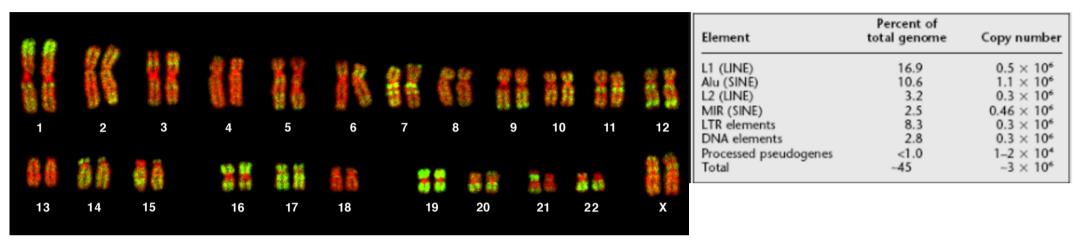




Centrifuge

2. Determination of presence or absence of Alu insert within the PV92 locus

Alu repeats in humans

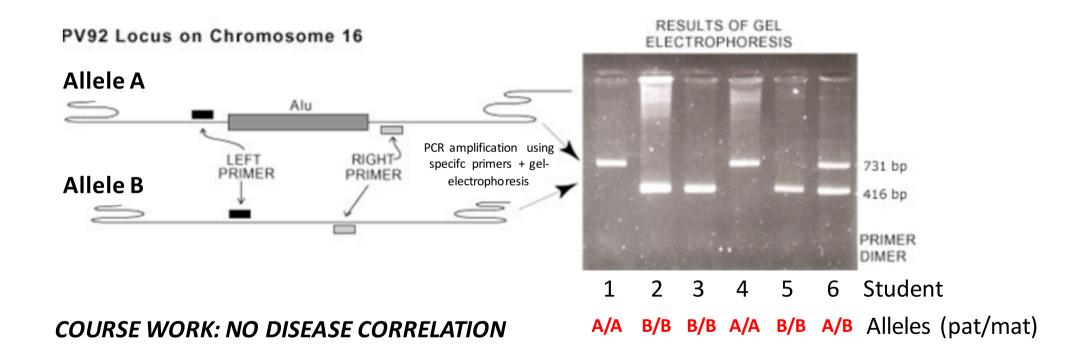


Karyotype from a female human lymphocyte (46, XX). Chromosomes were hybridized with a probe for Alu elements (green) and counterstained with TOPR (red). Alu elements were used as a marker for chromosomes and chromosome bands rich in genes.

Throughout evolution, intron sequences have been the target of random insertions by short repetitive interspersed elements, also known as SINEs.7 SINEs have become randomly inserted within our introns over millions of years. One such repetitive element is called the Alu sequence7 (Figure 2). This is a DNA sequence about 300 base pairs long that is repeated, one copy at a time, almost 500,000 times within the human genome.8 The origin and function of such randomly repeated sequences is not yet known. The Alu name comes from the Alu I restriction enzyme recognition site that is found in this sequence.

2. Determination of presence or absence of Alu insert within the PV92 locus

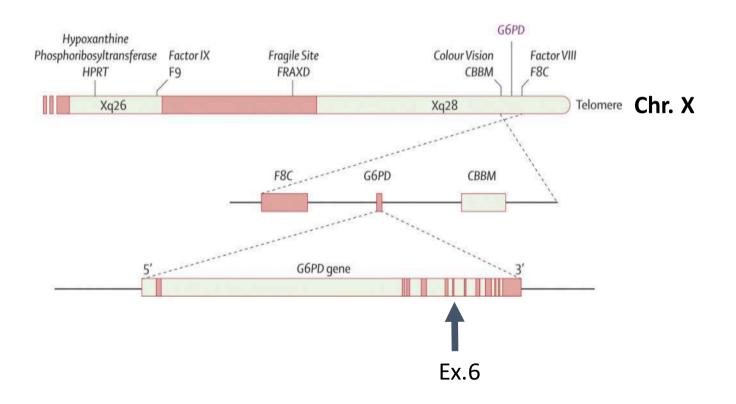
Alu repeats: Throughout evolution, intron sequences have been the target of random insertions by short repetitive interspersed elements, also known as SINEs.7 SINEs have become randomly inserted within our introns over millions of years. One such repetitive element is called the Alu sequence7 (Figure 2). This is a DNA sequence about 300 base pairs long that is repeated, one copy at a time, almost 500,000 times within the human genome.8 The origin and function of such randomly repeated sequences is not yet known. The Alu name comes from the Alu I restriction enzyme recognition site that is found in this sequence.



Students will perform a bioinformatics exercise to investigate the genotypic frequencies for the Alu polymorphism in their class population and compare them with the genotypic frequencies of other populations.

3. Determination of a G6PD variant by RFLP

Glucose-6-phosphate dehydrogenase deficiency (G6PDD) is an inborn error of metabolism that predisposes to red blood cell breakdown. Most of the time, those who are affected have no symptoms. Following a specific trigger, symptoms such as yellowish skin, dark urine, shortness of breath, and feeling tired may develop. Complications can include anemia and newborn jaundice (yellow pigmentation). Some people never have symptoms

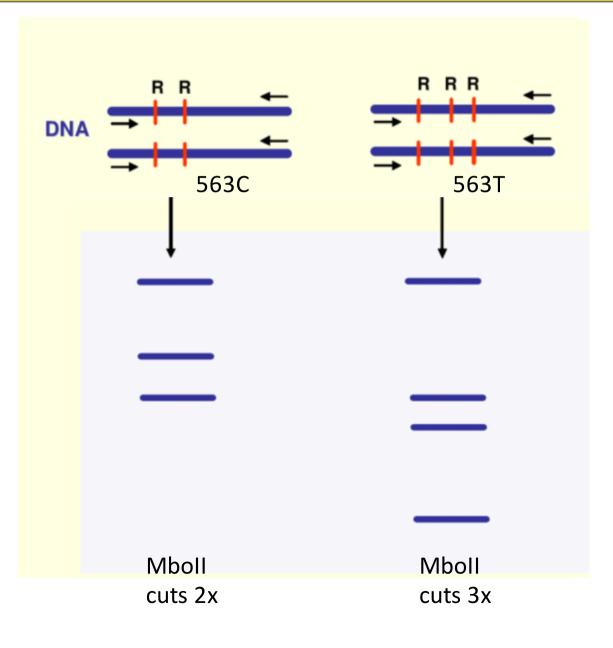


It is an X-linked recessive disorder that results in **defective glucose-6-phosphate dehydrogenase enzyme**. Red blood cell breakdown may be triggered by infections, certain medication, stress, or foods such as fava beans. Depending on the specific mutation the severity of the condition may vary. Diagnosis is based on symptoms and supported by blood tests and **genetic testing (PCR-RFLP)**

Many variants have been determined (>190)

				1	Descriptive	mutations	5			
Mutation			Gene		Protein					
Designation	Short name	Isoform G6PD- Protein	OMIM-Code	Туре	Subtype	Position	Position	Structure change	Function change	
G6PD-A(+)	Gd-A(+)	G6PD A	+305900.0001	Polymorphism nucleotide	A→G	376 (Exon 5)	126	Asparagine→Aspartic acid (ASN126ASP)	No enzyme defect (variant)	
G6PD-A(-)	Gd-A(-)	G6PD A	+305900.0002	Substitution nucleotide	G→A	376 (Exon 5) and 202	68 and 126	Valine→Methionine (VAL68MET) Asparagine→Aspartic acid (ASN126ASP)		
G6PD- Mediterranean	Gd-Med	G6PD B	+305900.0006	Substitution nucleotide	C→T	563 (Exon 6)	188	Serine→Phenylalanine (SER188PHE)	Class II	<
G6PD-Canton	Gd- Canton	G6PD B	+305900.0021	Substitution nucleotide	G→T	1376	459	Arginine→Leucine (ARG459LEU)	Class II	
G6PD- Chatham	Gd- Chatham	G6PD	+305900.0003	Substitution nucleotide	G→A	1003	335	Alanine→Threonine (ALA335THR)	Class II	
G6PD- Cosenza	Gd- Cosenza	G6PD B	+305900.0059	Substitution nucleotide	G→C	1376	459	Arginine→Proline (ARG459PRO)	G6PD-activity <10%, thus high portion of patients.	
G6PD-Mahidol	Gd- Mahidol	G6PD	+305900.0005	Substitution nucleotide	G→A	487 (Exon 6)	163	Glycine→Serine (GLY163SER)	Class III	
G6PD-Orissa	Gd- Orissa	G6PD	+305900.0047	Substitution nucleotide	C→G	131	44	Alanine→Glycine (ALA44GLY)	NADP-binding place affected. Higher stability than other variants.	
G6PD-Asahi	Gd- Asahi	G6PD A-	+305900.0054	Substitution nucleotide (several)	A→G ± G→A	376 (Exon 5) 202	126 68	Asparagine → Aspartic acid (ASN126ASP) Valine → Methionine (VAL68MET)	Class III.	

3. DETECTION OF G6PD 563 C→T VARIANT



G6PD 563 C \rightarrow T VARIANT IN EXON 6

- 1. G6PD Exon 6 specific primers
- 2. PCR amplify specific region of students
- 3. Purify PCR product
- 4. Digest purified DNA using Mboll
- 5. Run agarose gel
- 6. 563C→T variants results a new Mboll site in the PCR fragment
- 7. Additional band appears in gel

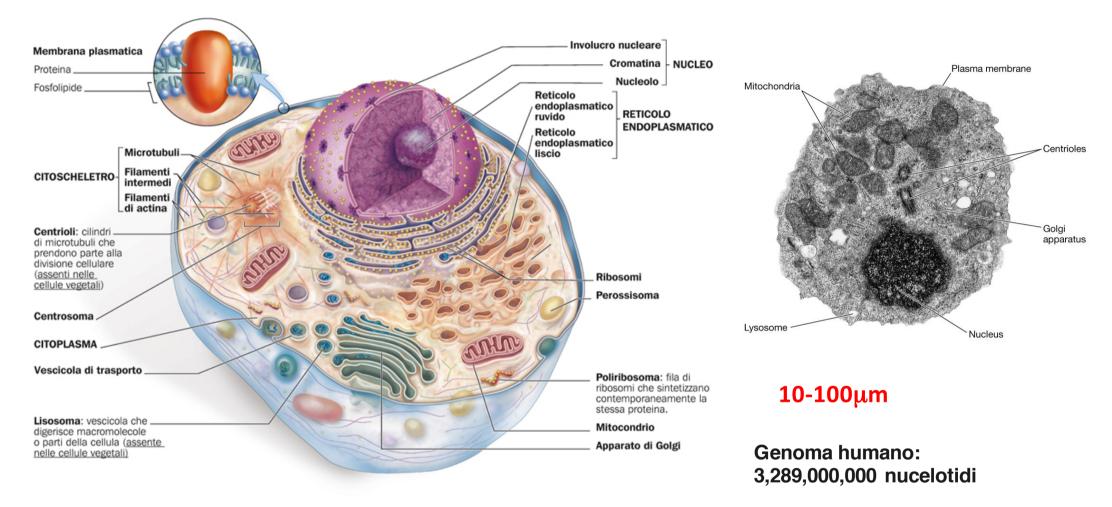
R = sito per enzima di restrizione Mboll

Materiale Didattica

Messaggi UNIVERSITÀ DEGLI STUDI DI TRIEST	Cerca corsi	STEFAN SCHOEFTNER V								
Moodle@UniTs Corsi Supporto	🚳 Dashboard 🛛 🛗	Eventi 🚔 I miei corsi	🎝 Questo corso	► ⊡ ⊀						
Corsi Dipartimento di Scienze della Vita	Laurea triennale (DM270)	SM51 - SCIENZE E TECNOLOG	GIE BIOLOGICHE A.A. 2018 - 2019							
210SM - LABORATORIO DI BIOLOGIA MOLECOLARE 2018										
Ricerca nei forum Vai Ricerca avanzata ?	Annunci									
<u> </u>	RNI DI LABORATO	RIO								
Annunci recenti (Nessuna news è stata ancora spedita)	TURNO I TURNO II									
Prossimi eventi	21 TURNO III 21 TURNO IV									
Non ci sono eventi prossimi										
Vai al calendario Nuovo evento PR	OGRAMMA ESER	CITAZIONI aa 2018-19								
Δttività										

1. RECOMBINANT DNA TECHNIQUES

La cellula eucariote



- Dimensioni: circa dieci volte piu' grandi delle cellule procariotiche (10-100 µm)
- La membrana plasmatica racchiude il materiale cellulare, lo separa dall'ambiente e regola il passaggio di sostanze cellula/esterno
- **Compartimentazione interna**: all'interno della membrana si trova il **citoplasma**, l'insieme del contenuto cellulare, comprendente il **citosol** (soluzione acquosa di piccole e grandi molecole) ed una serie di **organuli**, compartimenti funzionalmente specializzati delimitati da membrana o comunque strutturalmente separati (Apparato di Goghi; Mitocondrio; Reticilo endoplasmatico)

Genomes



Sex-reversal, autosomal Hyperglycinemia, nonketotic appression of tumorigenicity, pancreas Diaphyseal medullary stenosis Melanoma Trichoepithelioma, multiple familial Immotile cilia syndrome Cartilage-hair hypoplasia X-ray repair Fanconi anemia, complementation group G Sialuria Hyperoxaluria, primary, type II Cardiomyopathy Deafness autosomal recessive Choreoacanthocytosis Prostate-specific gene Bamforth-Lazarus syndrome Tyrosine kinase-like orphan receptor Brachydactyly, type B1 Nephronophthisis (infantile) Neuropathy, sensory and autonomic, type 1 Fructose intolerance Basal cell carcinoma, sporadic Muscular dystrophy, Fukuyama congenital Basal cell nevus syndrome Dysautonomia (Riley-Day syndrome) Esophageal cancer Endotoxin hyporesponsiveness Amyotrophic lateral sclerosis, juvenile dominant Berardinelli-Seip congenital lipodystrophy Dystonia, torsion, autosomal dominant Lethal congenital contracture syndrome Leukemia, acute undifferentiated Tuberous sclerosis Hemolytic anemia Telangiectasia, hereditary hemorrhagic Ehlers-Danlos syndrome, types I and II Joubert syndrome Leukemia, T-cell acute lymphoblastic

136 million base pairs

Ovarian cancer Albinism, brown and rufous Interferon, alpha, deficiency Loukomia Cyclin-dependent kinase inhibitor Venous malformations, multiple cutaneous and mucosal Arthrogryposis multiplex congenita, distal, type 1 Galactosemia Acromesomelic dysplasia, Maroteaux type Myopathy, inclusion body, autosomal recessive Hypomagnesemia with secondary hypocalcemia Friedreich ataxia Geniospasm Bleeding diathesis Hemophagocytic lymphohistiocytosis, familial Chondrosarcoma, extraskeletal myxoid Pseudohermaphroditism, male, with gynecomastia Tangier disease HDL deficiency, familial Fanconi anemia type C Xeroderma pigmentosum Epithelioma, self-healing, squamous Leukemia, T-cell acute lymphoblastic Muscular dystrophy, limb-girdle, type 2H Bladder cancer Sex reversal, XY, with adrenal failure Leukemia transcription factor, pre-B-cell Porphyria, acute hepatic Lead poisoning, susceptibility to Citrullinemia Dopamine-beta-hydroxylase deficiency Amyloidosis, Finnish type Microcephaly, primary autosomal recessive Leigh syndrome Leukemia Nail-patella syndrome Prostaglandin D2 synthase (brain) Pituitary hormone deficiency

Genoma umano aploide: 3.2 x 10⁹ bp (3200000000 bp)

→ 22 autosomi

→ eterocromosomi (X ed Y)

→ 23000 geni

Dimensione dei cromosomi: 45-275 Mb;

- \rightarrow 2.9 x 109 bp: eucromatina = attivo
- → Genoma noto: >90% dell'eucromatina.

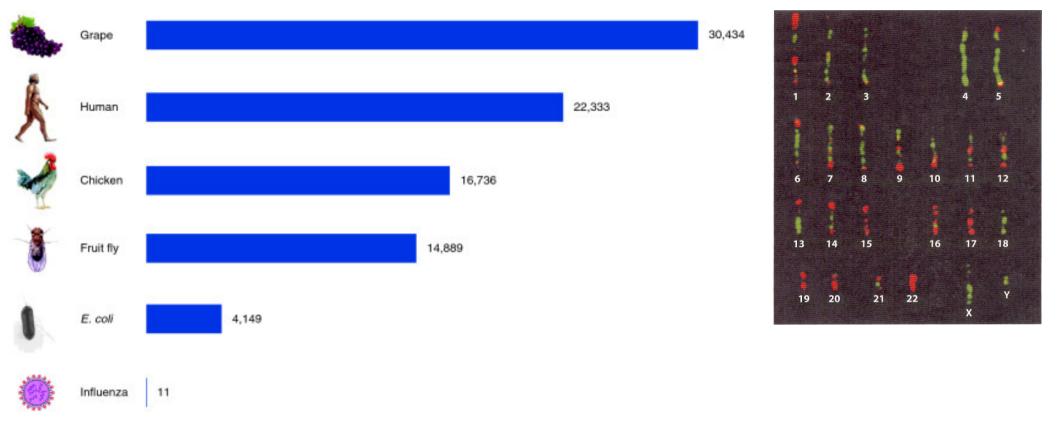
L'utillizo della infromazione genetica:

5.000-10.000 geni espressi da ogni cellula

- **100.000** specie proteiche diverse per modificazioni posttraduzionali
- **10**⁸ specie proteiche diverse nel genere umano (plasma: *proteoma di proteomi*)

ENORME COMPLESSITA

Gene numbers in different organisms



Mappa genica umana

Le regioni in rosso indicano porzioni dei cromosomi ad <u>alta densità genica</u> (ad esempio i cromosomi 15, 16, 17, 19, 20 e 22).

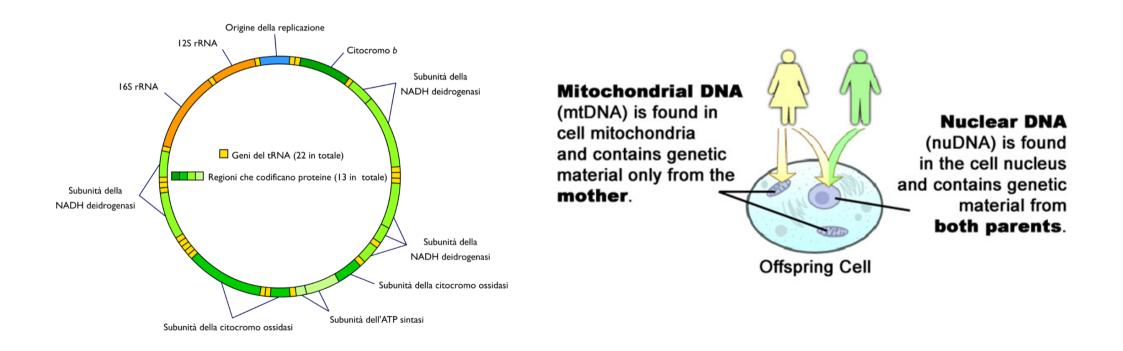
Altri cromosomi come 4, 18, X e Y mostrano una <u>colorazione rossa molto debole e sono poveri di</u> <u>geni.</u>

MITOCHONDRIAL DNA

DNA mitocondriale dell'uomo:

16569 paia di basi e 37 geni (codificano per 13 polipeptidi sintetizzati dal ribosoma mitocondriale

22 tRNA e 2 rRNA), coinvolti nella produzione di proteine necessarie alla respirazione cellulare.



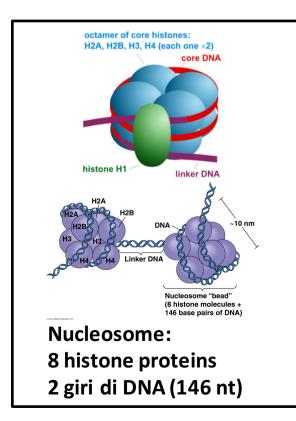
LA CROMATINA

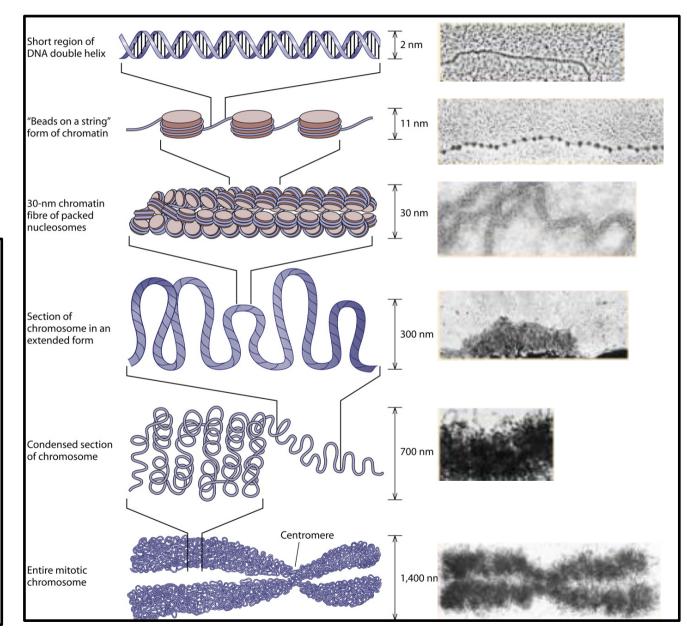
La **cromatina** è la forma in cui gli acidi nucleici si trovano nella cellula.

Funzione:

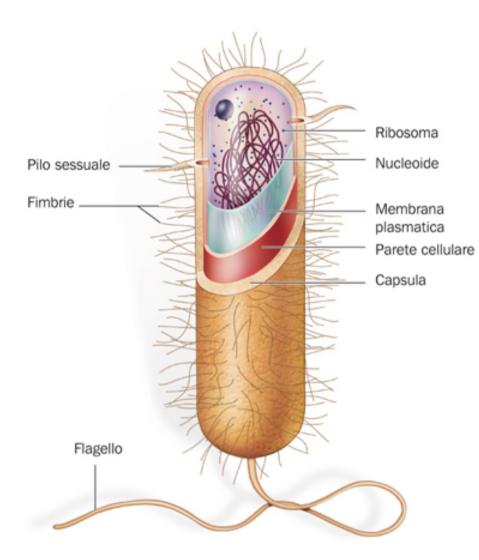
- impacchettamento del DNA

- -rafforzare il DNA per permettere la mitosi
- prevenire danni al DNA
- -controllare la replicazione del DNA
- e l'espressione (attivita) del gene





PROCARIOTI



Le cellule procariotiche (da *pro*, prima e *karyon*, nucleo) sono **prive di un nucleo** racchiuso da una membrana.

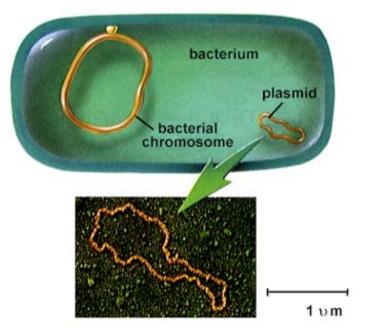
Gli organismi unicellulari costituiti da cellule procariotiche, i **procarioti**, sono classificati in due domini:

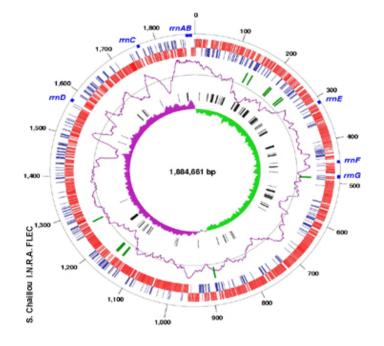
Archaea (archei);*Bacteria* (batteri).

PROCARIOTI - EUCARIOTI

Il materiale genetico, il DNA, e' organizzato in un **singolo cromosoma circolare,** localizzato nell'area nucleare o **nucleoide**, una regione della cellula non delimitata da membrana.

1-2 μ m (1.000.000 μ m = 1m)





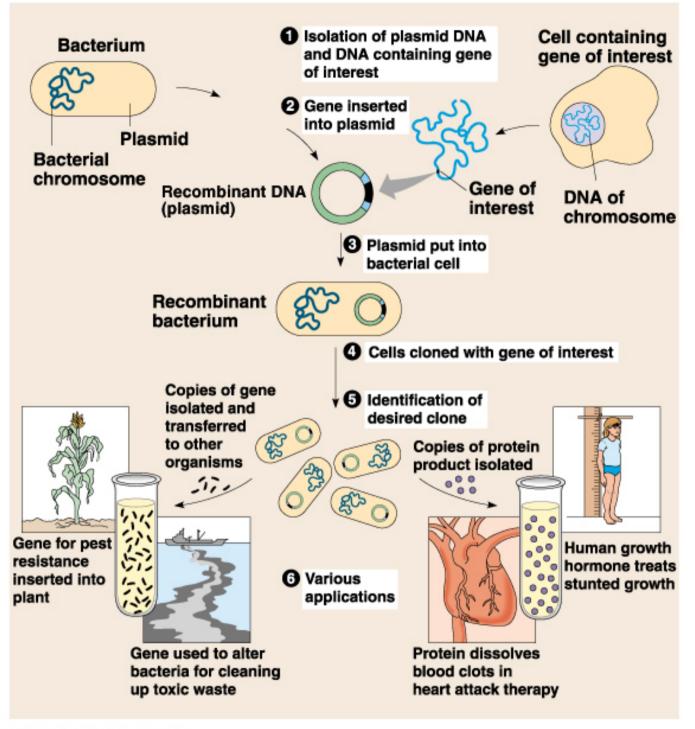
- In aggiunta al DNA principale i batteri possono contenere piccole molecole di DNA circolare, dette **plasmidi**, che codificano per enzimi catabolici, per la resistenza ad antibiotici o legati a meccanismi per lo scambio di materiale genetico tra organismi.
- Genoma: 130.000 14.000.000 nucleotidi

DNA RICOMBINANTE

tecnica che permette di

- ottenere brevi segmenti di DNA clonati e di studiarne la sequenza nucleotidica
- * di trasferirli nel genoma di altre cellule
- di controllare l'incorporazione e l'espressione del DNA clonato
- di introdurre mutazioni nel DNA e di studiarne gli effetti

A General Strategy to study or use recomobinant DNA



©1999 Addison Wesley Longman, Inc.