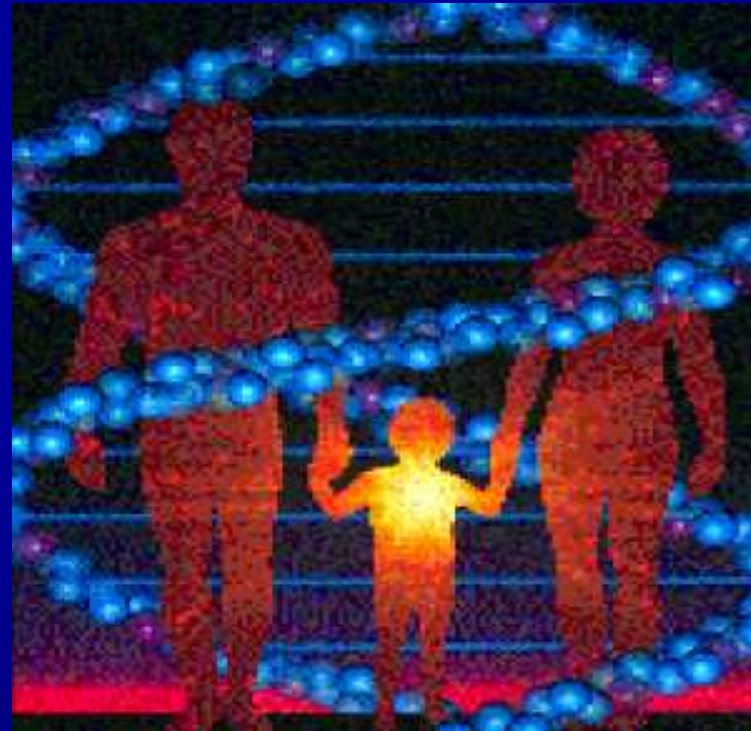


# GENETIC COUNSELLING

Process involving the individual  
and extending to him/her family



# GENETIC COUNSELLING

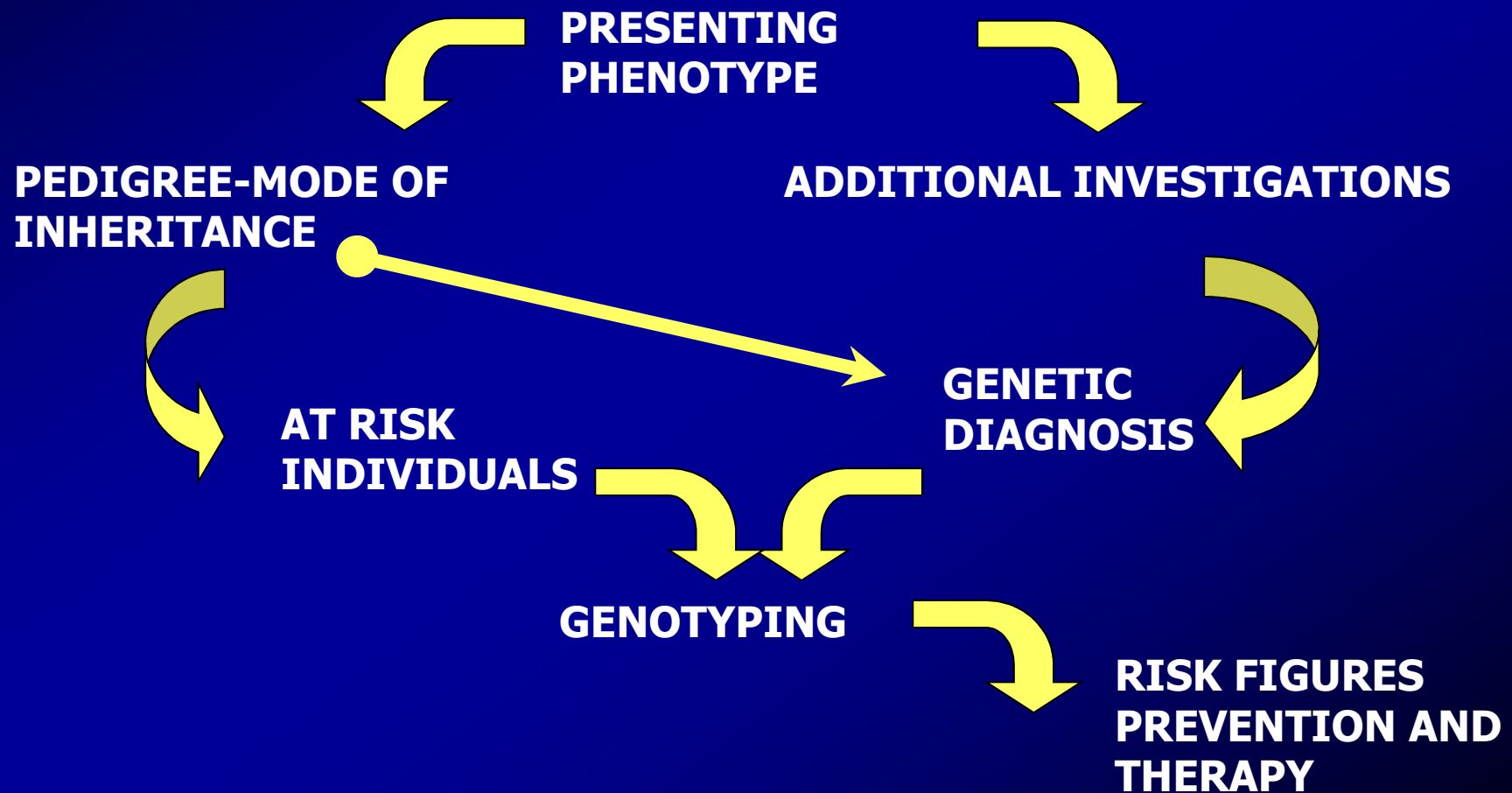
A non directive process by which patients or relatives at risk of a disorder that may be hereditary are advised of the consequences of the disorder, the probability of developing and transmitting it and the ways in which this may be prevented or ameliorated (Harper).

# GENETIC COUNSELLING

The genetic counselling has the goal to offer in a collaborative and integrated perspective possible options regarding:

- the mode of inheritance and related risk figures
- the cohort of genetic testing available
- the cohort of reproduction monitoring
- support in the autonomous decision process

# DIAGNOSIS: FUNDAMENTAL ELEMENTS OF GENETIC COUNSELLING



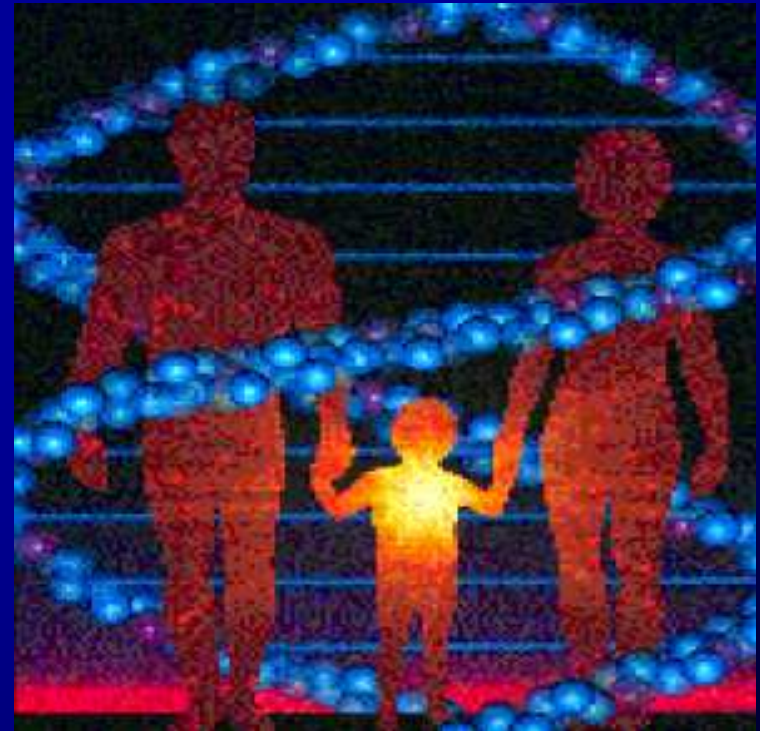
# **BASIC METHODOLOGY IN GENETIC COUNSELLING: an integrated view**

- Collection of the family history and pedigree (mode of inheritance)
- Detailed clinical information regarding the patient and relatives
- Proposing and organising novel clinical or instrumental investigations
- Collecting informed consents and ethical issues
- Addressing laboratory testing
- Interpreting the genetic testing outcome

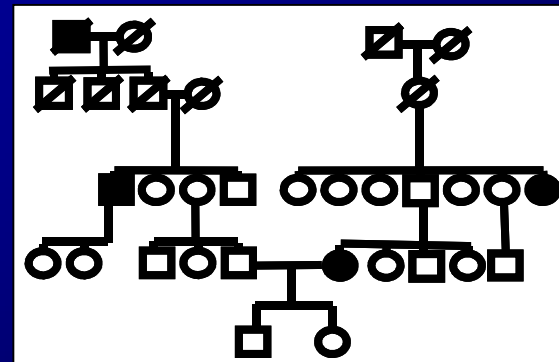
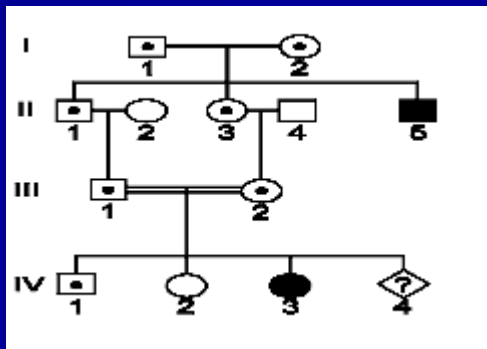
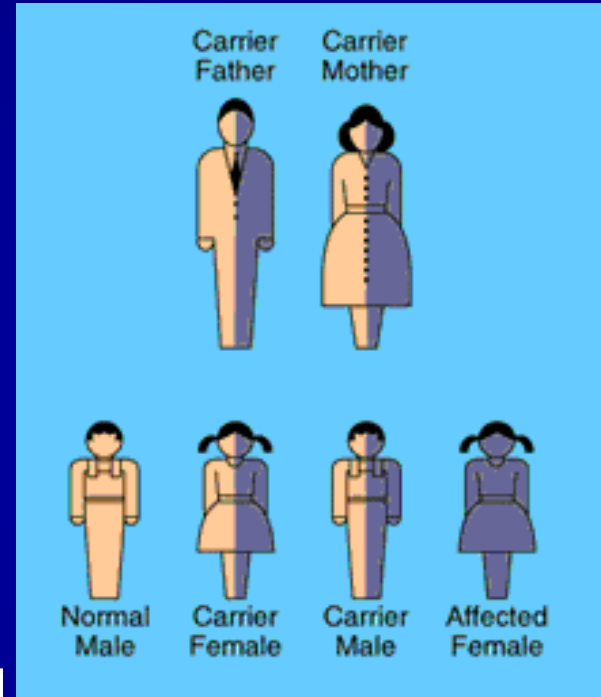
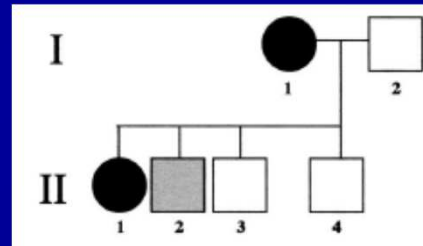
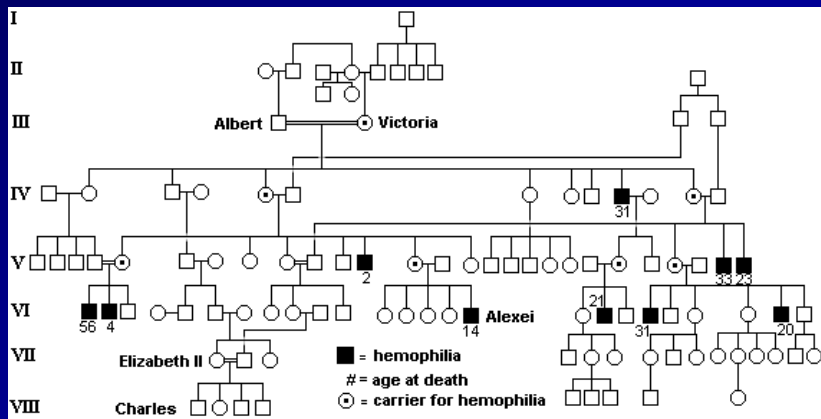
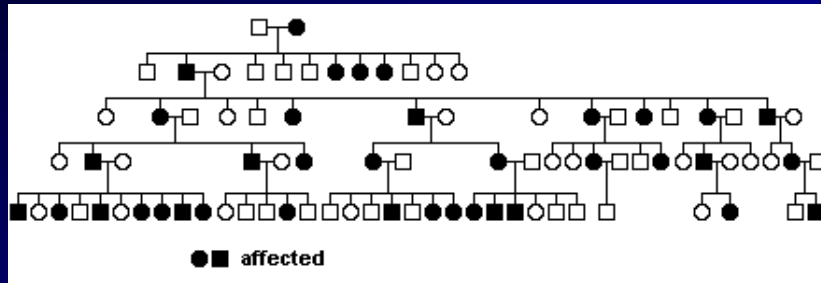
# OBJECTIVES OF GENETIC COUNSELLING

- To discriminate between environmental and genetic factors in the presenting phenotype
- To address a better phenotype definition
- To identify the possible mode of inheritance of the condition
- To identify the causative genotype
- To identify the at risk individuals
- To provide information regarding the quality and quantity of risk
- To provide support in the autonomous decision process in prevention and therapeutic strategies, taking into account the ethical issues

Collection of the  
family history and  
pedigree (mode of  
inheritance)

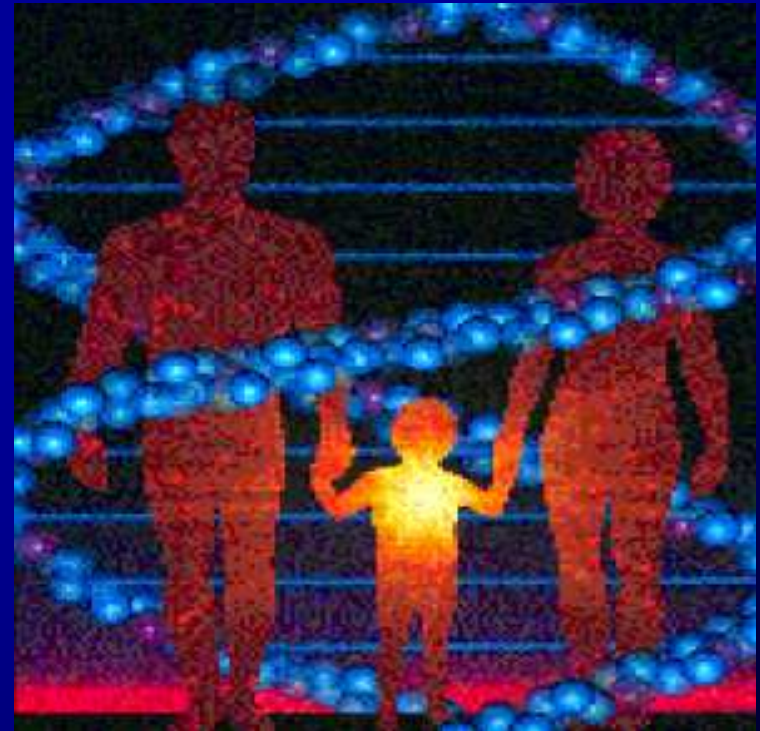


# Collection of the family history and pedigree (mode of inheritance)





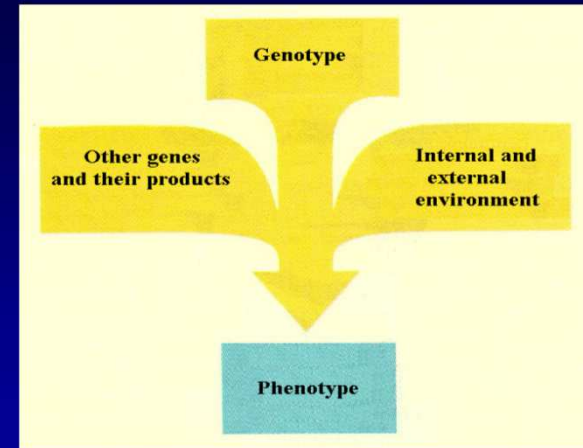
Detailed clinical  
information regarding  
the patient and  
relatives



# PHENOTYPE DIAGNOSIS (PHENOMICS)

CLINICAL  
INSTRUMENTAL  
BIOCHEMICAL  
IMMUNOCYTOCHEMICAL

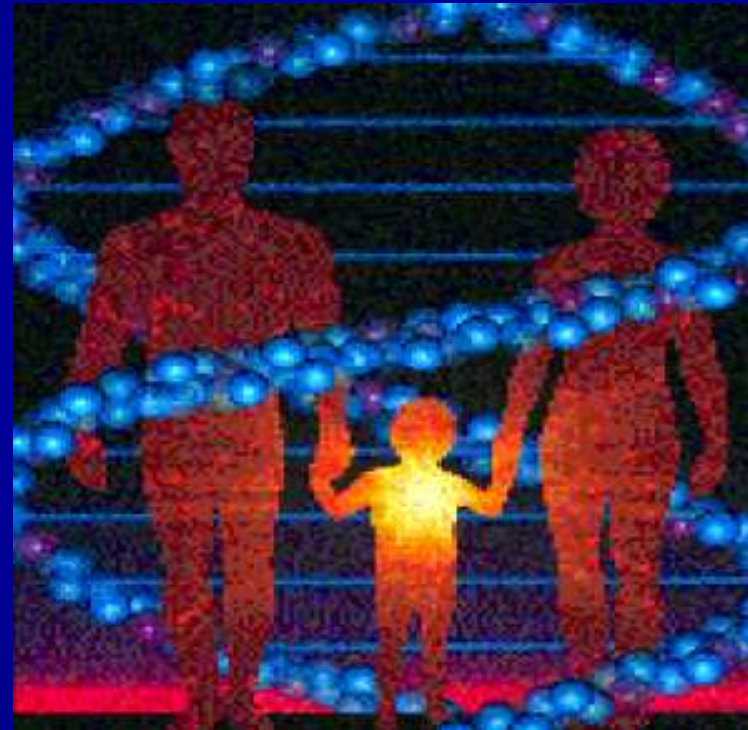
CLOSE INTERACTIONS  
BETWEEN MEDICAL  
GENETICIST AND OTHER  
SPECIALISTS



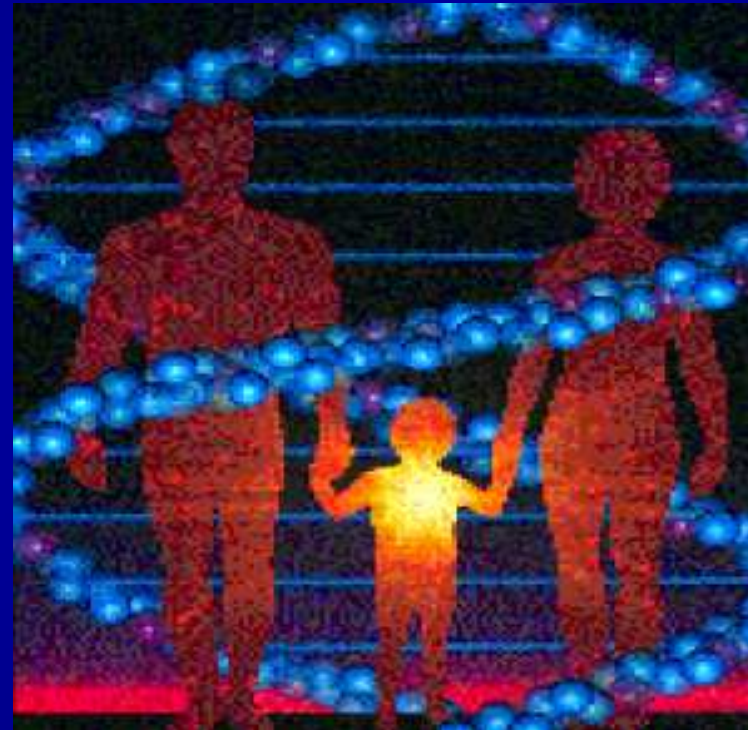
# Common pedigree problems

- **Different mutations can result in the same phenotype**
  - **Locus heterogeneity - mutations in different genes eg. Alpha and beta thalassemia**
  - **Allelic heterogeneity - different mutations in same gene e.g. Hb S and C**

- Proposing and organising novel clinical or instrumental investigations



- Collecting informed consents and ethical issues



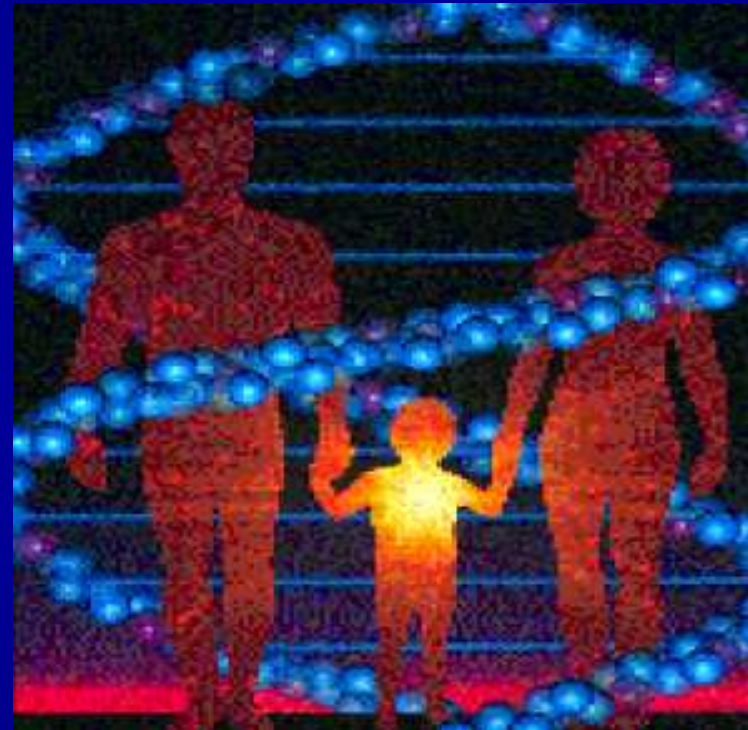
# Linee guida etiche per test genetici (WHO)

- \* they should be encouraged when the information might be used for prevention and therapy ( *health benefit* )
- \* they should be voluntary and based on the informed consent ( *autonomy* )
- \* they should be offered to adult individuals, even if a therapy or prevention are not available ( *autonomy* )
- \* the genetic tests on asymptomatic children should be avoided if a clear impact on prevention or therapy does not exist ( *health benefit* )
- \* Schools, education and governative programs, insurance and job providers must not have access to genetic testing results

The informed consent must always be collected informing the individuals about:

- significance of the test
- limits of the test
- possible results
- impact of the test on the reproductive life

- Addressing and interpreting laboratory testing



# GENETIC TESTS

Genetic tests  
are the most important  
product,  
in terms of clinical impact, of  
applied research



- Definition

The analysis of human DNA, RNA, chromosomes in order to detect heritable disease-related genotypes/mutations for clinical purposes

(US Task force on genetic testing 1999)

# GENETIC TESTS



## Investigations carried out

- Prenatal
  - Perinatal
  - Postnatal
- \* The typing of DNA/RNA and chromosomal regions will allow the identification of the genetic constitution of each individual

# GENETIC TESTS

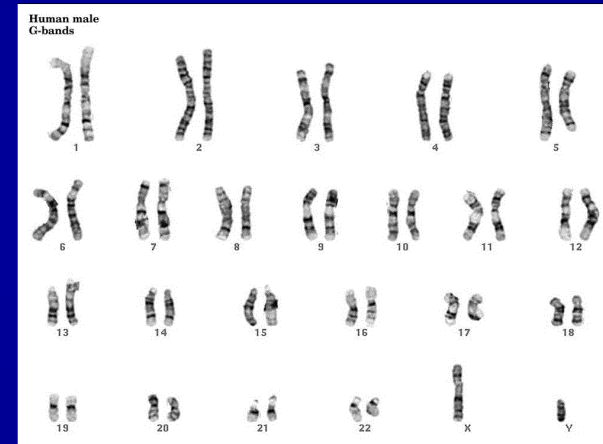


**Tests done on the genome as a whole**  
**cytogenetics**  
**gene profiling (arrays)**

**Test done on specific genes/regions**  
**molecular genetics**  
**molecular cytogenetics**

# GENETIC TESTS: cytogenetics

- Can review the whole genome
- Used for gross rearrangement abnormalities



## Characteristics:

- low resolution (standard karyotype)
- intermediate resolution (FISH)
- high resolution (CGH)

# **GENETIC TESTS: molecular genetics**

**-Can identify both small mutations  
and gross rearrangements**

**-Characteristics:**

**define the genotype  
are gene-specific**

**Limitations:**

**time consuming**

**expensive**

**requires high specialisation**

**mutations screening often difficult to be set up**

**(allele heterogeneity, private mutations, atypical mutations)**



# SCOPE OF GENETIC TESTS

- ✦ Diagnostic Test
- ✦ Carrier identification test
- ✦ Presymptomatic (or preclinical) test
- ✦ Predisposition test
- ✦ Screening test
- ✦ Medical/legal test

La bioinformatica come tool per  
la analisi di geni malattia

# Gene cards

**National Center for Biotechnology Information**

National Library of Medicine

National Institutes of Health

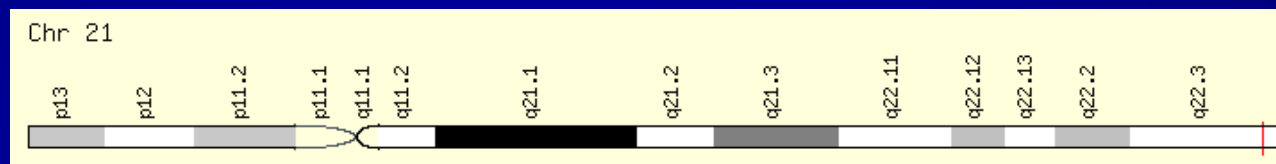
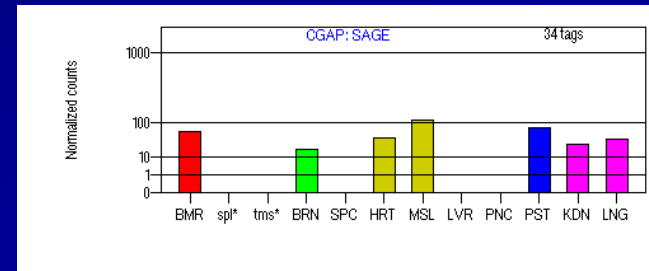
**The AceView genes**

*(last updated September 13th, 2005)*

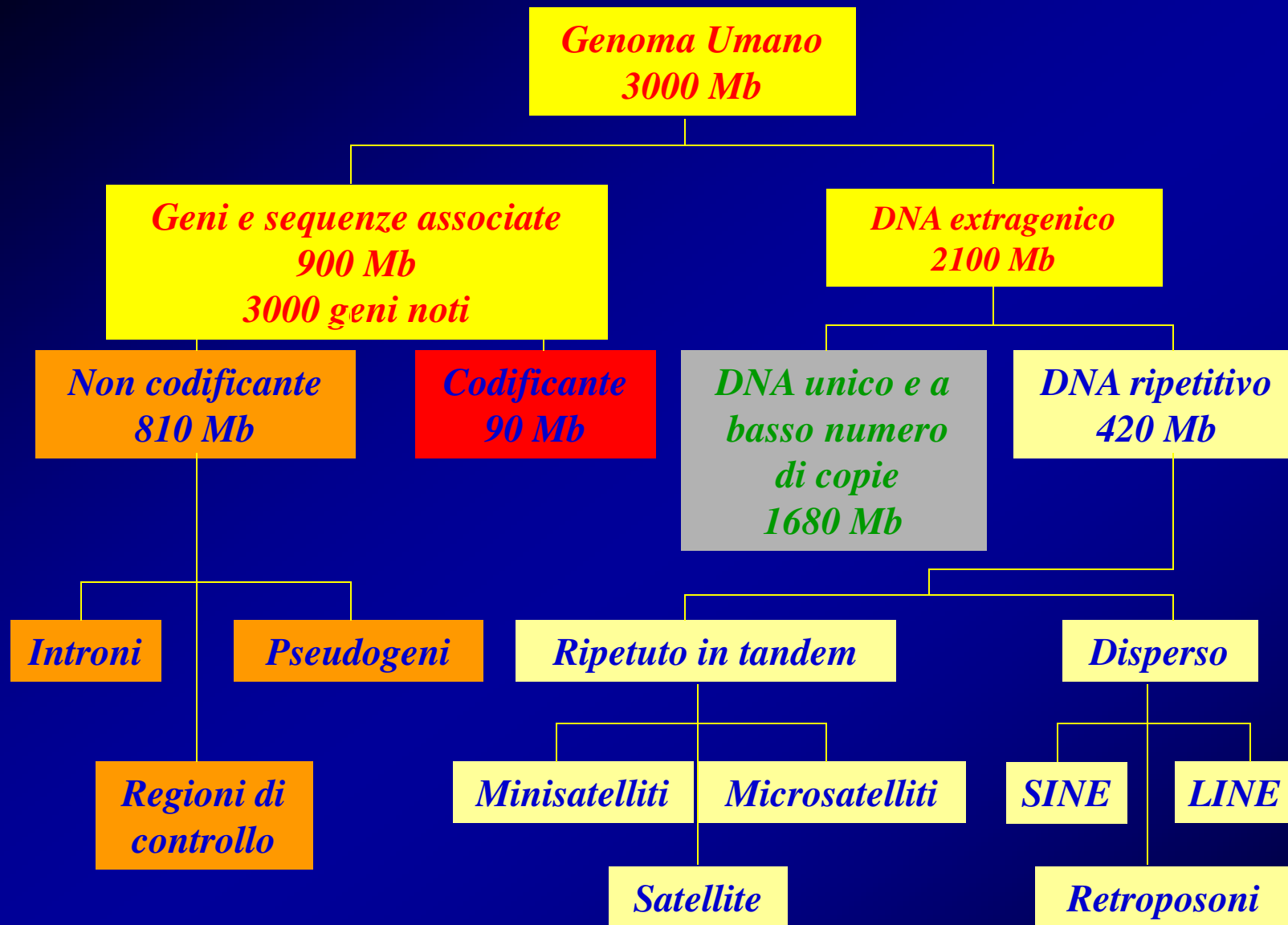




gcgccctgccatgggggaggggtgccaggggagagggcactgggggtgtctgagcgac  
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 CAGGAGGTCATCTCGCCGGACACTACCGAGAGAAACAACAACCTGCCAGGtgcca  
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 tcgggggaggtgaggtcaaaccacaacacagggcaggggcccaggaaacggggctc  
 caacagcagtcctcctgaggtggctcgtgacaggtcctgtgccccacaAGAAG  
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 CTTACCGACTGCGCGCTGGCCAACATGACGGAGCAGATCCGGCAGGACCGCAGC  
 AAGGGCACCCTCCACTTCGCCGTGGTCATCACCGACGGCCAGTCACCGGCAGCC  
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 CTTCGCCGTGGCCCCAACCAGAACCTGAAGGAGCAGGGCCTGCGGGACATCGCC  
 AGCAGCCGACGAGCTCTACCGCAACGACTACGCCACCATGCTGCCCGACTCCA  
 CGGAGATCGACCAGGACACCATCAACCGCATCATCAAGGTCATGgcegtccacce  
 actcggggcctcactttaccocctctgtgagtgcgaggccc



# Gene card for COL6alpha1 (congenital muscular dystrophy Ullrich type)



**Il DNA spazzatura è veramente tale?**

## LINK AL GENE

*Vediamo ora più in dettaglio la parte di record relativa al gene*

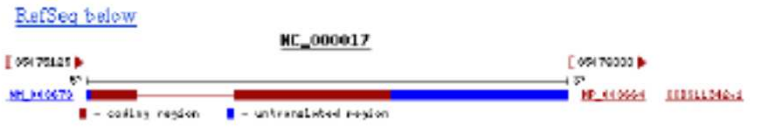
FEATURES	Location/Qualifiers	
source	<a href="#">1..963</a> /organism="Homo sapiens" /mol_type="mRNA" /db_xref="taxon:9606" /chromosome="17" /map="17q12"	tipo molecola (DNA, mRNA...) cromosoma e posizione di mappa
<a href="#">gene</a>	1..963 /gene="TCAP" /note="synonyms: TELE, CMD1N, T-cap, LGMD2G, telethonin" /db_xref="GeneID: <a href="#">8557</a> " /db_xref="LocusID: <a href="#">8557</a> " /db_xref="MIM: <a href="#">604488</a> "	nome ufficiale del gene e sinonimi link al database dei geni Entrez Gene link a malattie genetiche

**1: TCAP *titin-cap (telethonin)* [*Homo sapiens*]**  
 GeneID: 8557    Primary source: [HGNC:11610](#)    updated 03-Feb-2006

Summary ? ↑

**Official Symbol:** TCAP and **Name:** *titin-cap (telethonin)* provided by [HUGO Gene Nomenclature Committee](#)  
**See related:** [HPRD:05133](#), [MIM:604498](#)  
**Gene type:** protein coding  
**Gene name:** TCAP  
**Gene description:** *titin-cap (telethonin)*  
**RefSeq status:** Reviewed  
**Organism:** [Homo sapiens](#)  
**Lineage:** *Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominoidea; Homo*  
**Gene aliases:** TELE; CMD1N; T-cap; LGMD2G; telethonin  
**Summary:** Sarcomere assembly is regulated by the muscle protein titin. Titin is a giant elastic protein with kinase activity that extends half the length of a sarcomere. It serves as a scaffold to which myofibrils and other muscle related proteins are attached. This gene encodes a protein found in striated and cardiac muscle that binds to the titin Z1-Z2 domains and is a substrate of titin kinase, interactions thought to be critical to sarcomere assembly. Mutations in this gene are associated with limb-girdle muscular dystrophy type 2G.

Genomic regions, transcripts, and products ? ↑



struttura gene, parte trascritta, parte tradotta, esoni/introni, UTR

Genomic context [See TCAP in MapViewer](#)    ? ↑



Contesto genomico

# RICERCA DI GENI

Proviamo a fare una ricerca complessa usando gli operatori booleani

## [Getting started](#)

Look for genes by name part and multiple species

Look for genes by chromosome and symbol

## Sample queries

[transporter AND \("Drosophila melanogaster"\[orgn\] OR "Mus musculus"\[orgn\]\)](#) [more...](#)

[\(II\[chr\] OR 2\[chr\]\) AND adh\\*\[sym\]](#) [more...](#)

**Cerchiamo uno dei geni della rubisco**

[http://www.rcsb.org/pdb/molecules/pdb11\\_1.html](http://www.rcsb.org/pdb/molecules/pdb11_1.html)

**in Arabidopsis thaliana**

The screenshot shows the NCBI Entrez search interface. At the top, there are navigation tabs for Entrez, Pubmed, Nucleotide, Protein, Genome, Structure, PDB, Taxonomy, Books, and OMIM. The search bar contains the text "Gene" in a dropdown menu, followed by "for rubisco AND Arabidopsis thaliana [organism]". There are "Go" and "Clear" buttons. To the right, there is a checkbox for "current records only" which is checked. Below the search bar, there are links for "Limits", "Preview/Index", "History", "Clipboard", and "Details". On the left side, there is a sidebar with "Entrez" and "SITE MAP" links. Below the search bar, there is a "Display" dropdown menu set to "Summary", a "Show:" dropdown menu set to "5", and a "Send to" dropdown menu set to "Text". At the bottom, it shows "Items 1-5 of 13" and a pagination control showing "Page 1 of 3" with a "Next" link.

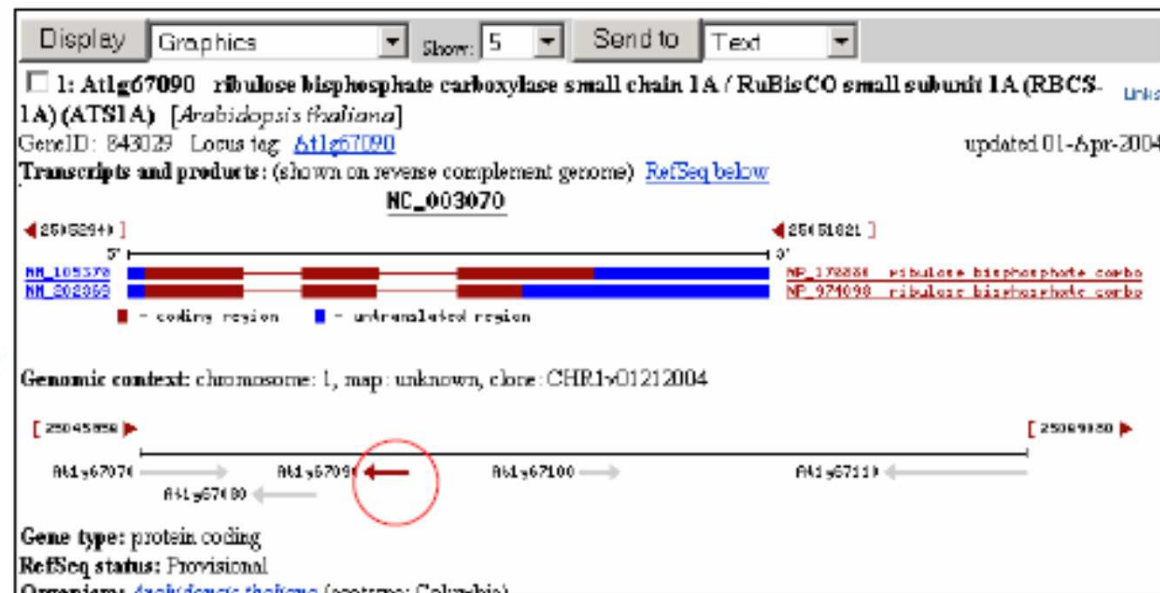
# RICERCA DI GENI

•Siamo interessati al gene “subunità 1A” di rubisco.

Database	From	Genome	Structure	PubMed	Taxonomy	Books	ORF
	for	rubisco AND 1A AND Arabidopsis thaliana [org]	<input type="button" value="Go"/>	<input type="button" value="Clear"/>	<input checked="" type="checkbox"/>	current records only	
<a href="#">Limits</a>	<a href="#">Preview/Index</a>	<a href="#">History</a>	<a href="#">Clipboard</a>	<a href="#">Details</a>			
Display	Summary	Show: 5	Send to	Text			

- 1: [At1g67090](#) Link  
ribulose biphosphate carboxylase small chain 1A / RuBisCO small subunit 1A (RBCS-1A) (ATS1A) [*Arabidopsis thaliana*]  
**Other Aliases:** At1g67090, F1019.14  
**Chromosome:** 1  
**GeneID:** 843029

struttura gene, parte trascritta (elica complementare!), parte tradotta, esoni/introni, UTR



## LINK ALLE MALATTIE GENETICHE

*Si possono esaminare le malattie genetiche associate a quella posizione di mappa (gene)*

```
FEATURES          Location/Qualifiers
  source           1..963
                  /organism="Homo sapiens"
                  /mol_type="mRNA"
                  /db_xref="taxon:9606"
                  /chromosome="17"
                  /map="17q12"
  gene            1..963
                  /gene="TCAP"
                  /note="synonyms: TELE, CMD1N, T-cap, LGMD2G, telethonin"
                  /db_xref="GeneID:8557"
                  /db_xref="LocusID:8557"
                  /db_xref="MIM:604488"
```

Mutazioni (alterazioni della sequenza nucleotidica di un gene) possono riflettersi in alterazioni della funzionalità della proteina da esso codificata. Queste mutazioni possono causare le cosiddette **malattie genetiche**.

Esempio: una mutazione a carico del gene della  $\beta$  globina fa sì che una particolare base del gene venga sostituita con un'altra, ciò altera il codone e nella proteina ciò si riflette nella sostituzione di un glutamato con una valina e in una ridotta funzionalità della proteina che causa una malattia genetica detta anemia a cellule falciformi (anemia falciforme).

Mutazioni a carico di geni differenti causano molte malattie genetiche diverse per questo è stato costituito il database OMIM.





## Database di malattie genetiche (umane)



**NCBI**  
OMIM  
Online Mendelian Inheritance in Man  
Johns Hopkins University

Search OMIM for [Go] [Clear]

Display: Detailed Show 20 Send to

Alt: 1

**\*604488**  
**TITIN-CAP: TCAP**  
Alternative titles: symbols  
**TELETHONIN**  
Gene map locus: [11q12](#)

**TEXT**

**DESCRIPTION**  
Telethonin is a sarcomeric protein of 19 kD found cultured myocytes. Telethonin is a substrate of the other sarcomeric proteins. After activation by phorbol myristate.

**MAPPING**  
[Yaffe et al. \(1997\)](#) mapped the telethonin gene to

**.0001 MUSCULAR DYSTROPHY, LIMB-GIRDLE, TYPE 2G [TCAP, GLN53TER]**  
[Moxima et al. \(2000\)](#) observed 2 different mutations in the telethonin gene in 3 LGMD2G (601954) families. A C57C-to-T transition in exon 2 created a premature stop codon (glu53 to ter) and affected patients from 2 kindreds were homozygous for this mutation, whereas patients from a third kindred were heterozygous. The second mutation in the latter patient was a deletion of 2 guanine nucleotides within 4 guanines at the junction of exon 1 and intron 1 (604488.0002).

**.0002 MUSCULAR DYSTROPHY, LIMB-GIRDLE, TYPE 2G [TCAP, 2-BP DEL, 637GG]**  
In a family with limb-girdle muscular dystrophy type 2G (601954), [Moxima et al. \(2000\)](#) found that affected members were compound heterozygotes for the Q53X mutation (604488.0001) and for deletion of 2 guanine nucleotides within a 4 guanine run (nucleotides 637-640 in the genomic sequence) at the junction of exon 1 and intron 1.

**.0003 CARDIOMYOPATHY, DILATED, 1N [TCAP, ARG87GLN]**  
[Knoll et al. \(2002\)](#) screened for mutations in the TCAP gene in 380 dilated cardiomyopathy patients (DCM) with 100 controls from the collected population of the University Hospital Benjamin Franklin in Berlin and identified a mutation, arg87 to glu (R87Q), in 1 patient. The resulting form of DCM was designated CMD1N (607487). The R87Q mutation was not found in any individual of the control population or in any of 400 control individuals in Japan.

**REFERENCES**  
1. Knoll, R.; Hoshijima, M.; Hoffman, H. M.; Person, V.; Lorenzen-Schmidt, J.; Bang, M.-L.; Hayashi, T.; Sligo, N.;

Anche qui  
possiamo  
fare ricerche  
complesse

Esempio di una query sul database OMIM: da notare l'estensiva descrizione di quanto noto sulla/e malattia/e determinate da mutazioni a carico del gene in esame

## LA REGIONE CODIFICANTE (CDS)

Consideriamo ora solo la parte codificante (tradotta in aminoacidi) della sequenza di RNA messaggero

```
gene      1..963      ← Il trascritto è lungo 963 nucleotidi
/gene="TCAD"
/note="synonyms: TELE, CMD1N, T-cap, LGMD2G, telethonin"
/db_xref="GeneID:8557"
/db_xref="LocusID:8557"
/db_xref="MIM:604488"

CDS       15..518    ← CDS: La parte tradotta va dalla base 15 alle 518
/gene="TCAD"
/note="19 kDa sarcomeric protein;"
go_component: cytoplasm [goid 0005737] [evidence NR];
go_function: structural constituent of muscle [goid 0008307] [evidence IAS] [pmid 9350988];
go_process: sarcomere alignment [goid 0006938] [evidence IAS] [pmid 9817758];
go_process: cell shape and cell size control [goid 0007148] [evidence E] [pmid 9817758];
go_process: protein complex assembly [goid 0006461] [evidence TAS] [pmid 9817758]"
/codon_start=1
/product="telethonin"
/protein_id="WP_003664.1" ← Link alla proteina
/db_xref="GI:4507435"
/db_xref="GeneID:8557"
/db_xref="LocusID:8557"
/db_xref="MIM:604488"

/translation="MATSSELSCEVSEENCEDREAFWAEWEDITLSTDPHECCSLHEED
TQPHETTHQQGQCQVVLVQRSPWLMRRGILGRGLQEQYQLPYQEVLPPIFTPAKMGAT
KEEREDTPIQLQELLALETALGGQCVDREQEVAEITKQLPYPVFSKPGALRRSLSESM
SQEAQDC"
```

**componente cellulare**  
**funzione**

**processo biologico**

**GENE ONTOLOGY**  
<http://www.geneontology.org/>

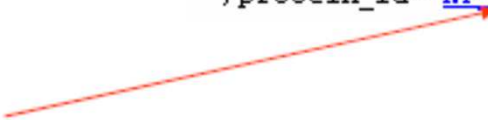
**Sequenza Proteina**

## LINK ALLA PROTEINA

Clickando sul link `protein_id` si arriva al record della proteina corrispondente

```
LOCUS       NP_003664                167 aa             linear   PRI 20-DEC-2003
DEFINITION  telethonin; 19 kDa sarcomeric protein [Homo sapiens].
ACCESSION   NP_003664
VERSION     NP_003664.1  CI:4507435
DESOURCE    EMBSEQ: accession NM\_003673.2
KEYWORDS    .
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Cranialia; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (residues 1 to 167)
AUTHORS     Zou,P., Gubel,M., Geerlof,A., Wilmanns,M., Koch,M.H. and
            Svergun,D.I.
TITLE       Solution scattering suggests cross-linking function of telethonin
            in the complex with titin
JOURNAL     J. Biol. Chem. 278 (4), 2636-2644 (2003)
PUBMED     12445666
REMARK     GeneRIF: telethonin may play a role in linking titin filaments at
            the Z-disk periphery
UNRECORDED 2  (residues 1 to 167)
```

`/protein_id="NP_003664.1"`




# PROTEINA

Modalità grafica. Vedere il funzionamento delle opzioni

Display Graphics Show: 1 Send to File Get Subsequence

1: [NP\\_003664](#) telothronin, 19 kD... [gi.450743]

CDS with gene and mRNA  Other features  Hide sequence Refresh [Hide Toolbar](#)



Legend:  
— protein — other feature

Sequence:

```
1 HRTSELSCEV SEENCERRR FARELIKILT STRPEEGCSL HEEDQRHET YHRRGCDAL telothronin
61 VDRSPALMYR IICILGRGLDE YQLPYDRMLP LPIFTPTNIG RTKEEREDTP IQLDLLEALE telothronin
121 TFLGGSCVDR QEVREITNGL PPWPUSKPG ALRESLSRSM SZEFRRG telothronin
      phosphorylation
```

[Protein](#) [Gene](#) [Structure](#) [FAC](#) [Taxonomy](#) [Links](#)  
 For rubisco AND 1a AND arabidopsis thaliana [organi]    
[Limits](#) [Preview/Index](#) [History](#) [Clipboard](#) [Details](#)

Database: cdd, v. 1.6.5

Click on boxes for multiple alignments

1: [P10795](#)  
 Ribulose b...  
 gi2773322

2: [NP\\_974098](#)  
 ribulose bi...  
 thaliana]  
 gi4257201

3: [NP\\_176880](#)  
 Ribulose biphosphate carboxylase small chain 1A / RuBisCO small subunit 1A (RBCS-1A) (ATS1A) [Arabidopsis thaliana]  
 gi15219826[ref][NP\_176880.1][15219826]

[BLink](#) [Domains](#) [Links](#)

Domini funzionali della proteina

Legend:  
 — protein

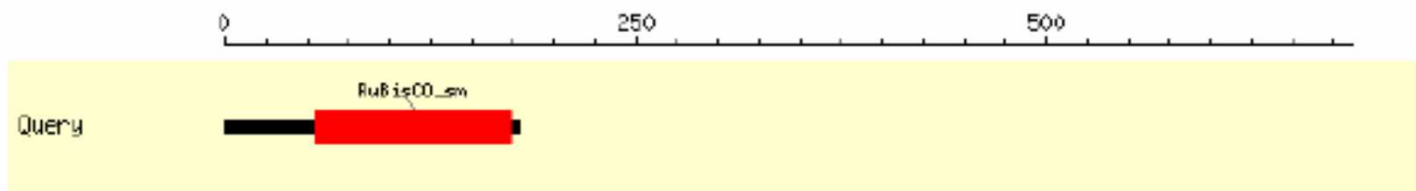
Sequence:

1 MRSSTLSERT IIVSRPQRTD IIAFFNGLKSS RAFPATIKRN NDLTSTISNG GRVNCIDMHP  
 61 PIGKKKFKETL SYLPDLTDE LAKEVDYLIR NKHIPDVEFE LEHCFVYREH DNSPCYYDGR  
 121 YNTINKLPLF GCTDSRAULK EMSECKKEYP NAFIRIIGFD NTRQVDCISF IAYKPPSFTG

ribulose bisphos  
 ribulose bisphospha  
 ribulose bisphospha  
 ribulose bisphospha

# PROTEINE CON GLI STESSI DOMINI

Sono immediatamente accessibili anche proteine aventi gli stessi domini



## Similar domain architectures



http://www.ncbi.nlm.nih.gov/genome/guide/human/

The screenshot displays the NCBI Entrez search engine interface. At the top left is the NCBI logo. The main header features the Entrez logo and the text "Entrez, The Life Sciences Search Engine". Below this is a navigation bar with tabs for "HOME", "SEARCH", "SITE MAP", "PubMed", "All Databases", "Human Genome", "GenBank", "Map Viewer", and "BLAST". The "Human Genome" tab is selected. A search bar contains the text "pd38", with "GO" and "CLEAR" buttons and a "Help" link to its right. The search results are organized into two columns of database entries, each with a count, an icon, a name, a description, and a help icon.

Count	Icon	Database Name	Description	Help
3114	M	PubMed	biomedical literature citations and abstracts	?
726	Open Book	PubMed Central	free, full text journal articles	?
none	W	Site Search	NCBI web and FTP sites	?
37	B	Books	online books	?
21	Person	OMIM	online Mendelian Inheritance in Man	?
none	Hand	OMIA	Online Mendelian Inheritance in Animals	?
1942	Globe	Nucleotide	sequence database (includes GenBank)	?
199	Protein	Protein	sequence database	?
6	Genome	Genome	whole genome sequences	?
13	Structure	Structure	three-dimensional macromolecular structures	?
none	Microscope	Taxonomy	organisms in GenBank	?
549	SNP	SNP	single nucleotide polymorphism	?
34	Gene	Gene	gene-centered information	?
13	HomoloGene	HomoloGene	eukaryotic homology groups	?
none	PubChem	PubChem Compound	unique small molecule chemical structures	?
11	UniGene	UniGene	gene-oriented clusters of transcript sequences	?
1	CDD	CDD	conserved protein domain database	?
28	3D Domains	3D Domains	domains from Entrez Structure	?
17	UniSTS	UniSTS	markers and mapping data	?
15	PopSet	PopSet	population study data sets	?
46432	GEO Profiles	GEO Profiles	expression and molecular abundance profiles	?
23	GEO DataSets	GEO DataSets	experimental sets of GEO data	?
53	Cancer Chromosomes	Cancer Chromosomes	cytogenetic databases	?
none	PubChem BioAssay	PubChem BioAssay	bioactivity screens of chemical substances	?

<http://www.ncbi.nlm.nih.gov/omim/>



- OMIM: online mendelian inheritance in man
- Creato da Victor McKusick
- Continuamente aggiornato
- Punto chiave per acquisire informazione sui caratteri mendeliani umani, patologici e non
- A ogni carattere viene attribuito un numero a 6 cifre  
MIM



**NCBI**

- MIM +143100
- Description
- Clinical Features
- Biochemical Features
- Other Features
- Inheritance
- Mapping
- Molecular Genetics
- Pathogenesis
- Diagnosis
- Population Genetics
- Clinical Management
- Animal Model
- History
- Allelic Variants
- View List
- See Also
- References
- Contributors
- Creation Date
- Edit History
- Clinical Synopsis
- Gene map
- Entrez Gene
- Nomenclature
- MIM +143100
- Description
- Clinical Features
- Biochemical Features
- Other Features
- Inheritance
- Mapping
- Molecular Genetics
- Pathogenesis
- Diagnosis
- Population Genetics
- Clinical Management



All Databases: PubMed Nucleotide Protein Structure PMC Taxonomy OMIM

Search OMIM for [ ] Go Clear

Limits Preview/Index History Clipboard Details

Display Detailed Show 20 Send to

All: 1

**+143100**  
**HUNTINGTON DISEASE; HD**

Alternative titles; symbols

- HUNTINGTON CHOREA
- HUNTINGTIN, INCLUDED; HD, INCLUDED
- HTT, INCLUDED
- IT15, INCLUDED

Gene map locus [4p16.3](#)

**TEXT**

**DESCRIPTION**

Huntington disease (HTT) is inherited as an autosomal dominant disease that gives rise to progressive, selective (non-size-4) neuronal cell death associated with choreic movements and dementia. The disease is associated with increases in the length of a CAG triplet repeat present in a gene called 'huntingtin' located on chromosome 4p16.3.

**CLINICAL FEATURES**

The classic signs of Huntington disease are progressive chorea, rigidity, and dementia, frequently associated with seizures. A characteristic atrophy of the caudate nucleus is seen radiographically. Typically, there is a prodromal phase of mild psychotic and behavioral symptoms which precedes frank chorea by up to 10 years. The results of a study by [Shwach and Norbury \(1994\)](#) clashed with the conventional wisdom that psychiatric symptoms are a frequent presentation of Huntington disease before the development of neurologic symptoms. They performed a control study of 93 neurologically healthy individuals at risk for Huntington disease. The 20 asymptomatic heterozygotes showed no increased incidence of psychiatric disease of any sort when compared to the 33 normal homozygotes in the same group. However, the whole group of heterozygous and homozygous normal at-risk individuals showed a significantly greater number of psychiatric episodes than did their 43 spouses, suggesting stress from the uncertainty associated with belonging to a family segregating this disorder. [Shwach and Norbury \(1994\)](#) concluded that neither depression nor psychiatric disorders are likely to be significant preneurologic indicators of heterozygous expression of the disease gene.

GeneTests, Links

MIM Gene map - Microsoft Internet Explorer

File Edit View Favorites Tools Help

Back Forward Stop Home Search Favorites

Address <http://www.ncbi.nlm.nih.gov/Omim/getmap.cgi?143100>

Google omim

Web Accelerator

NCBI Online Mendelian Inheritance in Man

OMIM

Johns Hopkins University

PubMed Nucleotide Protein Genome PopSet Taxonomy OMIM

The OMIM Gene map presents the cytogenetic map location of disease genes and other expressed genes described in OMIM. See the [OMIM Morbid Map](#) for a list of disease genes organized by disease. For more refined maps of genes and DNA segments click on the **Location** to invoke NCBI Entrez [Map Viewer](#).

Search for:    (from the current location)

- Enter gene symbol, chromosomal location, or disorder keyword to search for, e.g. "CYP1", "5", "1pter", "Xq", or "alzheimer".
- You must capitalize X and Y to search for those chromosomes.

**4p16.3, HD to 4p16.1, HMX1**

<<Move Up Move Down>>

Location	Symbol	Title	MIM #	Disorder	Comments	Method	Mouse
<a href="#">4p16.3</a>	HD, IT15	Huntingtin	<a href="#">143100</a>	Huntington disease (3)	distal to D4S10	Fd	<a href="#">5(Hdh)</a>
<a href="#">4p16.3</a>	IDUA, IDA	Iduronidase, alpha-L-	<a href="#">252800</a>	Mucopolysaccharidosis Ih, <a href="#">607014 (3)</a> ; Mucopolysaccharidosis Is, <a href="#">607016 (3)</a> ; Mucopolysaccharidosis Ih/s, <a href="#">607015 (3)</a>		REa, A, S	<a href="#">5(Ldua)</a>
<a href="#">4p16.3</a>	LETM1	Leucine zipper/EF-hand-containing transmembrane protein 1	<a href="#">604407</a>			A	
<a href="#">4p16.3</a>	LRPAP1, A2MRAP	Low density lipoprotein-related protein-associated protein 1 (alpha-2-macroglobulin receptor-associated protein 1)	<a href="#">104225</a>			?involved in Wolf-Hirschhorn syndrome	
<a href="#">4p16.3</a>	MYL5	Myosin, light polypeptide-5, regulatory	<a href="#">160782</a>			A, REa	
<a href="#">4p16.3</a>	PDE6B, PDEB, CSNB3	Phosphodiesterase-6B, cGMP-specific, rod, beta	<a href="#">180072</a>	Night blindness, congenital stationary, type 3, <a href="#">163500 (3)</a> ; Retinitis pigmentosa, autosomal recessive (3)		RE	
<a href="#">4p16.3</a>						REa, A, Fd	<a href="#">5(Pdeb, rd)</a>

The OMIM Morbid Map presents the cytogenetic map location of disease genes described in OMIM. For a map organized by chromosome, see the [OMIM Gene Map](#). For more refined maps of genes and DNA segments, use NCBI Entrez [Map Viewer](#) and the [Genome Database](#).

Search for:    (from the current location)

- Enter gene symbol, chromosomal location, or disorder keyword to search for, e.g. "recessive", "CYP1" "5", "1pter", "5", "1pter", or "Xq".
- You must capitalize X and Y to search for those chromosomes.

[<<Move Up](#) [Move Down>>](#)

Disorder	Symbol(s)	OMIM	Location
Huntington disease (3)	HD, IT15	<a href="#">143100</a>	4p16.3
Huntington disease-like 1, <a href="#">603218</a> (3)	PRNP, PRIP	<a href="#">176640</a>	20pter-p12
Huntington disease-like 2, <a href="#">606438</a> (3)	JPH3, JP3, HDL2	<a href="#">605268</a>	16q24.3
Huntington disease-like 3 (2)	HDL3, HLN2	<a href="#">604802</a>	4p15.3
Huntington disease-like-4, <a href="#">607136</a> (3)	TBP, SCA17	<a href="#">600075</a>	6q27
Huniez syndrome (2)	TYS, HRZ	<a href="#">181600</a>	4q23
Hyalinosis, infantile systemic, <a href="#">236490</a> (3)	ANTXR2, CMG2, JHF, ISH	<a href="#">608041</a>	4q21
Hydatidiform mole, <a href="#">231090</a> (3)	NALP7, NOD12, PYPAF3, HYDM	<a href="#">609661</a>	19q13.4
Hydrocephalus due to aqueductal stenosis, <a href="#">307000</a> (3)	L1CAM, CAML1, HSAS1	<a href="#">308840</a>	Xq28
Hydrocephalus with Hirschsprung disease and cleft palate, <a href="#">142623</a> (3)	L1CAM, CAML1, HSAS1	<a href="#">308840</a>	Xq28
Hydrocephalus with congenital idiopathic intestinal pseudoobstruction, <a href="#">307000</a> (3)	L1CAM, CAML1, HSAS1	<a href="#">308840</a>	Xq28




The OMIM Morbid Map presents the cytogenetic map location of disease genes described in OMIM. For a map organized by chromosome, see the [OMIM Gene Map](#). For more refined maps of genes and DNA segments, use NCBI Entrez [Map Viewer](#) and the [Genome Database](#).

Search for:    (from the current location)

- Enter gene symbol, chromosomal location, or disorder keyword to search for, e.g. "recessive", "CYP1", "5", "1pter", or "Xq".
- You must capitalize X and Y to search for those chromosomes.

[<<Move Up](#) [Move Down>>](#)

Disorder	Symbol(s)	OMIM	Location
{Obesity, susceptibility to}, <a href="#">601665</a> (3)	ADRB2	<a href="#">109690</a>	5q32-q34
{Obesity, susceptibility to}, <a href="#">601665</a> (3)	ADRB3	<a href="#">109691</a>	8p12-p11.2
{Obesity, susceptibility to}, <a href="#">601665</a> (3)	CART	<a href="#">602606</a>	5q13.2
{Obesity, susceptibility to}, <a href="#">601665</a> (3)	ENPP1, PDNRP1, NPPS, M6S1, PCA1	<a href="#">173335</a>	6q22-q23
{Obesity, susceptibility to}, <a href="#">601665</a> (3)	GHRL	<a href="#">605353</a>	3p26-p25
{Obesity, susceptibility to}, <a href="#">601665</a> (3)	UCP1	<a href="#">113730</a>	4q31
{Obesity, susceptibility to}, <a href="#">601665</a> (3)	UCP2	<a href="#">601693</a>	11q13
{Obesity/hyperinsulinism, susceptibility to} (2)	OQTL	<a href="#">602025</a>	20q13.11-q13.2
{Obsessive-compulsive disorder 1}, <a href="#">164230</a> (3)	SLC6A4, HTT, OCD1	<a href="#">182138</a>	17q11.1-q12
{Obsessive-compulsive disorder, protection against}, <a href="#">164230</a> (3)	BDNF	<a href="#">113505</a>	11p13
{Obsessive-compulsive disorder, susceptibility to}, <a href="#">164230</a> (3)	HTR2A	<a href="#">182135</a>	13q14-q21



# NCBI Map Viewer

PubMed
Nucleotide
Protein
Genome
Gene
Structure
PopSet
Taxonomy
Help

Search for  on chromosome(s)

Show related entries

**Map Viewer**  
Map Viewer Home

Map Viewer Help  
Human Maps Help

Release Notes

**NCBI Resources**  
Genome Project  
TaxPlot  
Consensus Coding Sequence (CCDS)  
Human Genome Resources  
NCBI Handbook  
RefSeq  
Whole Genome Association (WGA)

[Organism Data in](#)

**BLAST search the human genome**

**Homo sapiens (human) genome view**  
[Build 36.1 statistics](#) [Switch to previous build](#)

**Lineage:** [Eukaryota](#), [Metazoa](#), [Chordata](#), [Cranata](#), [Vertebrata](#), [Euteleostomi](#), [Mammalia](#), [Eutheria](#), [Euarchontoglires](#), [Primates](#), [Haplorrhini](#), [Catarrhini](#), [Hominoidea](#), [Homo](#), [Homo sapiens](#)

**March 2006:** NCBI released an update for the human genome (NCBI Build 36.1) that includes some changes to the reference genome assembly as well as updated annotation. This release includes a major change to the Map Viewer in that the previous build (NCBI Build 35.1) can still be accessed for Map Viewer display and for BLAST. For additional information about changes, statistics, and the status of the CCDS project please refer to:

- [Release Notes](#)
- [Statistics](#)
- [CCDS Project](#)

The NCBI Map Viewer provides graphical displays of features on the human genome sequence assembly as well as cytogenetic, genetic, physical, and radiation hybrid maps. Extensive documentation is provided to describe the resource features and methods used, tutorials, and statistics.

Map features that can be seen along the sequence include genes, transcripts, NCBI contigs (the 'Contig' map), the BAC tiling path (the 'Component' map), STSs, FISH mapped clones, ESTs and transcripts from several different organisms, [Gnomon](#) predicted gene models, and more.

You can find genes or markers of interest by submitting a query against the whole genome, or a chromosome at a time. Use the Advanced Search form for more complex

Overview page (Build 5.1)

Map Viewer Home

Map Viewer Help

Human Maps Help

TP

Data As Table View

Maps & Options

Compress Map

Region Shown:

Go

out  
 zoom  
 in

You are here:

Ideogram



Region Displayed: 0-181M bp

Ideogram

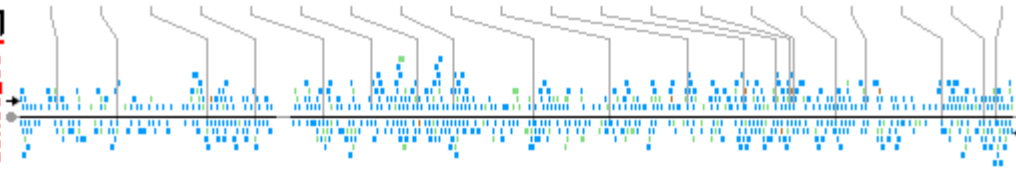
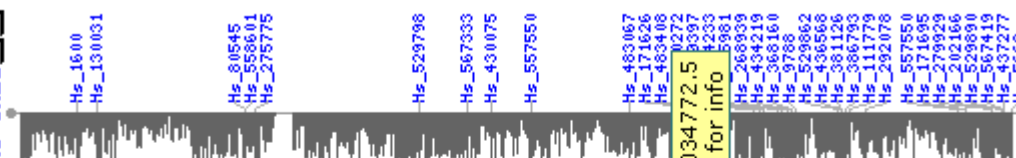
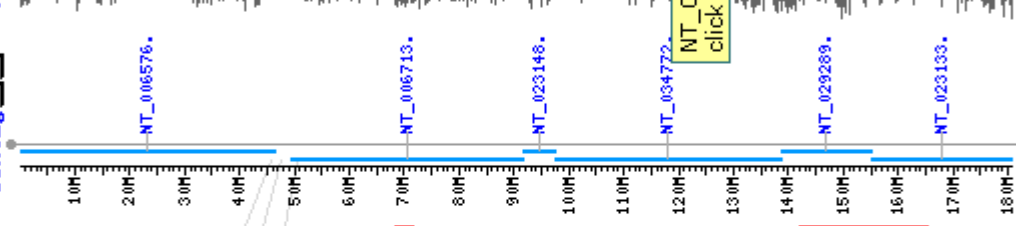
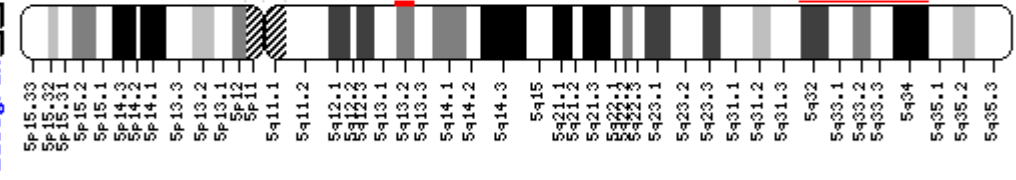
Contig

Hs Unig

Genes-seq

Symbol

Links



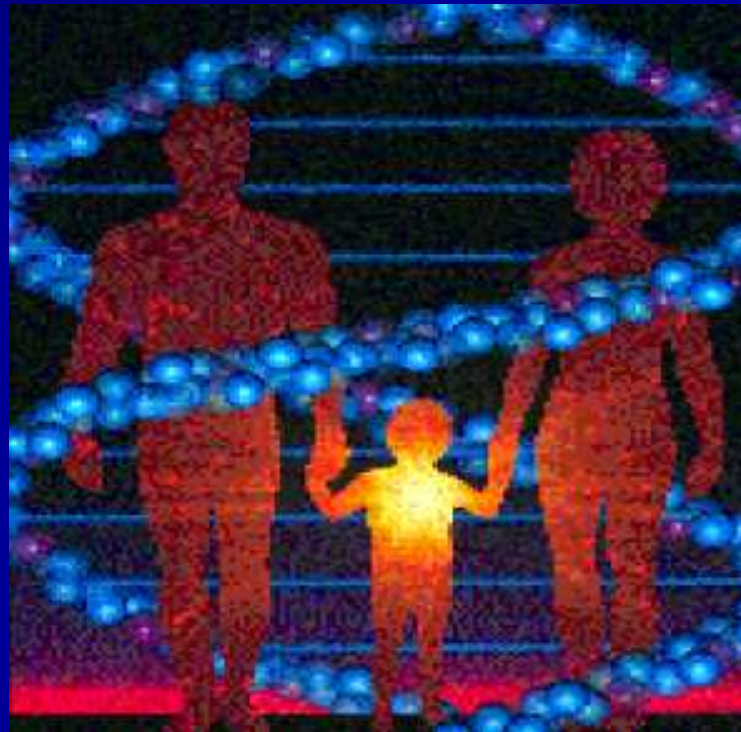
Symbol	Links	RefSeq
<a href="#">SRD5A1</a>	<a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">sts</a> <a href="#">CCDS</a>	best RefSeq
<a href="#">LOC646126</a>	<a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a>	protein
<a href="#">C1QTNF3</a>	<a href="#">HGNC</a> <a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">sts</a> <a href="#">CCDS</a>	best RefSeq
<a href="#">SEPP1</a>	<a href="#">HGNC</a> <a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">sts</a>	best RefSeq
<a href="#">IL6ST</a>	<a href="#">OMIM</a> <a href="#">HGNC</a> <a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">sts</a> <a href="#">CCDS</a>	best RefSeq
<a href="#">MRPL49P1</a>	<a href="#">sv</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a>	best RefSeq
<a href="#">PRP2</a>	<a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">CCDS</a>	best RefSeq
<a href="#">PAPP4</a>	<a href="#">HGNC</a> <a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">sts</a> <a href="#">CCDS</a>	best RefSeq
<a href="#">LOC642488</a>	<a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a>	mRNA
<a href="#">FBXL17</a>	<a href="#">OMIM</a> <a href="#">HGNC</a> <a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">sts</a>	best RefSeq
<a href="#">FLJ32921</a>	<a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">CCDS</a>	best RefSeq
<a href="#">RAD50</a>	<a href="#">OMIM</a> <a href="#">HGNC</a> <a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">sts</a>	best RefSeq
<a href="#">KIF20A</a>	<a href="#">OMIM</a> <a href="#">HGNC</a> <a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">sts</a> <a href="#">CCDS</a>	best RefSeq
<a href="#">PCDHA6</a>	<a href="#">OMIM</a> <a href="#">HGNC</a> <a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">sts</a>	best RefSeq
<a href="#">PCDHGC5</a>	<a href="#">OMIM</a> <a href="#">HGNC</a> <a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">sts</a> <a href="#">CCDS</a>	best RefSeq
<a href="#">IL17B</a>	<a href="#">OMIM</a> <a href="#">HGNC</a> <a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">CCDS</a>	best RefSeq
<a href="#">LARP1</a>	<a href="#">HGNC</a> <a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">sts</a> <a href="#">CCDS</a>	best RefSeq
<a href="#">PANK3</a>	<a href="#">OMIM</a> <a href="#">HGNC</a> <a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">sts</a> <a href="#">CCDS</a>	best RefSeq
<a href="#">HSPC111</a>	<a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">sts</a>	best RefSeq
<a href="#">COL23A1</a>	<a href="#">HGNC</a> <a href="#">sv</a> <a href="#">pr</a> <a href="#">dl</a> <a href="#">ev</a> <a href="#">mm</a> <a href="#">hm</a> <a href="#">sts</a> <a href="#">CCDS</a>	best RefSeq

default

# url per bioinformatica IN GENETICA MEDICA

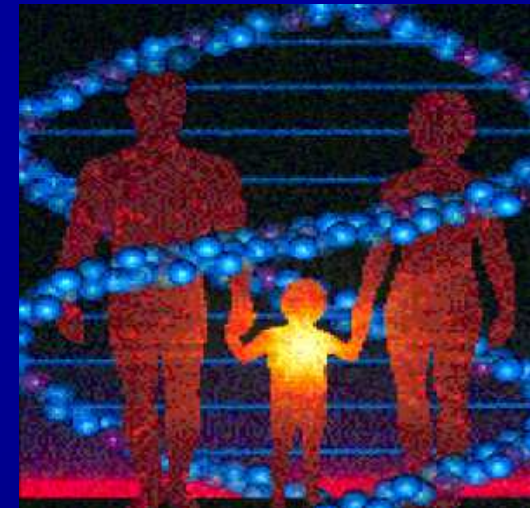
- OMIM
- <http://www.ncbi.nlm.nih.gov/omim>
- NCBI
- <http://www.ncbi.nlm.nih.gov/>
- GENECARDS
- <http://www.genecards.org/>
- GENEREVIEWS
- <http://www.ncbi.nlm.nih.gov/sites/GeneTests/review?db=GeneTests>
- ORPHANET
- <http://www.orphanet-italia.it/national/IT-IT/index/homepage/>
- EUCERD
- <http://www.eucerd.eu/>

# ETICA IN GENETICA MEDICA





# Introduciamo il concetto di BioEtica



# BioEtica : le origini

## Quali sono le origini della bioetica?

1. Il *Codice di Norimberga* (1947) con la condanna di ogni **sperimentazione** sull'uomo senza il suo consenso
2. Allarme suscitato da talune **sperimentazioni** negli USA
3. La *Dichiarazione di Helsinki* sulla **sperimentazione** clinica (1964-2000)



# BioEtica: il termine

## Chi ha coniato il termine *Bioetica*?

- 1970-71: l'oncologo americano V. R. POTTER

– prima volta del termine “BIOETICA”


- 1970 - *Bioethics. The science of survival*
- 1971 - *Bioethics. Bridge to the future*

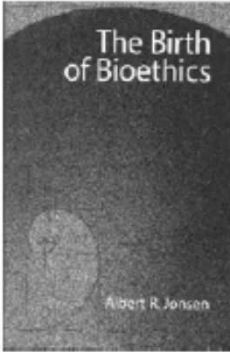
– Di fronte al pericolo per la sopravvivenza dell'intero ecosistema, la BIOETICA come “una nuova disciplina che combinasse la conoscenza biologica con la conoscenza del sistema dei valori umani”

•Un “ponte” tra due culture: scientifica e umanistica




# BioEtica e sperimentazione


 A parte il nome ... i problemi che portarono allo sviluppo della bioetica. 1




- **La sperimentazione sull'uomo**

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 Gli episodi di sperimentazione "selvaggia"



- Il processo di Norimberga contro i crimini dei medici nazisti
- Ulteriori rilevazione di abusi su soggetti umani:
  - 800 bambini handicappati psichici vengono inoculati con siero infetto per studiare la patogenesi dell'epatite virale e sviluppare metodi per l'immunizzazione al *Willowbrook State Hospital di NY*, 1956-1970
  - 600 pazienti ("black men, mostly poor and uneducated") vengono reclutati nel 1932 in Alabama, per valutare la storia naturale della sifilide e lasciati senza terapia anche quando la penicillina divenne disponibile, fino all'inizio degli anni '70 (*Tuskegee Syphilis Study*)
  - ...

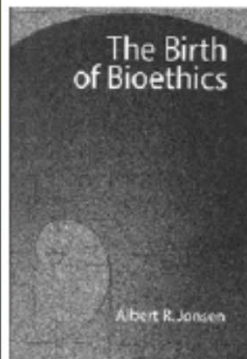
  
Charles Richard, one of the study subjects, at the Tuskegee Institute. (1968)

12

# BioEtica e genetica



A parte il nome ... i problemi che portarono allo sviluppo della bioetica. 2



- ♦ **La sperimentazione sull'uomo**
- ♦ **Le scoperte della genetica**



## La genetica e la nascita della *Bioetica*



- ♦ *Stanford University, inizi degli anni '70: programmazione di esperimento di integrazione nell'Escherichia Coli del genoma del virus SV40 (tumori nella scimmia, non infettivo per l'uomo, in associazione a tumori cerebrali)*

Interrogativi:

- 1) Se l'E.Coli fosse sfuggito al controllo?
- 2) Se avesse infettato i ricercatori?
- 3) Se si fossero scatenate epidemie di cancro?

**L'ESPERIMENTO NON FU PIÙ ESEGUITO!**

- ♦ *La genetica entra nei Parlamenti: audizioni di scienziati e ricercatori (lo stesso Watson nel 1971, che illustra gli enzimi di restrizione)*

# Genetica e bioetica nel terzo millennio

# I pericoli della biogenetica: la storia

- Piano **eugenetico nazista**: eliminazione degli ebrei abitanti in Europa, dei gruppi “inferiori” e di quelli “difettosi”
- “Legge di sterilizzazione genetica”: sterilizzazione di 225.000 persone per ordine della Corte di Sanità Ereditaria
- Leggi di Norimberga, 1935: piano diventa esplicitamente antisemita, proibiti i matrimoni tra persone di provenienza etnica differente
- 1939: programma di eutanasia per gli individui “inutili”

# Principi guida per i test genetici

- Tutela della confidenzialità dei dati, rispetto dell'autonomia personale, conoscenza più approfondita della genetica, previsione di beneficio, equità d'accesso (President's Commission, *Linee guida*, 1981)
- Screening genetici obbligatori: mai per ottenere un *pool* genetico sano o una riduzione dei costi sanitari



# Documenti internazionali

- Consiglio d'Europa, *Convenzione di bioetica* :  
“Non si potrà procedere a *test predittivi di malattie genetiche* o che permettano di *identificare il soggetto come portatore* di un gene responsabile di una malattia o di scoprire una predisposizione o suscettibilità genetica ad una malattia, se non *a fini medici o di ricerca medica*, e sotto riserva di un *counseling genetico appropriato*”

# Terapie geniche

- Somatiche: si interviene sul patrimonio genetico di cellule dell'organismo (soma=corpo)
- Germinali: si interviene nelle prime fasi di formazione dello zigote o dell'embrione, o sui gameti, e si elimina il difetto genetico, trasmettendo così la modificazione anche agli eventuali discendenti

# A favore delle terapie geniche

- Doveri di cura
- utilità terapeutica
- efficienza preventiva
- rispetto della scelta autonoma delle coppie
- rispetto della libertà della ricerca

# Contro l'uso delle terapie geniche

- Imprevedibilità dei rischi
- effetti sulle generazioni future
- costi alti delle pratiche: questioni di giustizia?
- integrità del patrimonio genetico
- “pendio scivoloso”: da cura a manipolazione?
- Eugenetica?
- Confine terapia - potenziamento?
- Discrimine salute - malattia?

# La clonazione



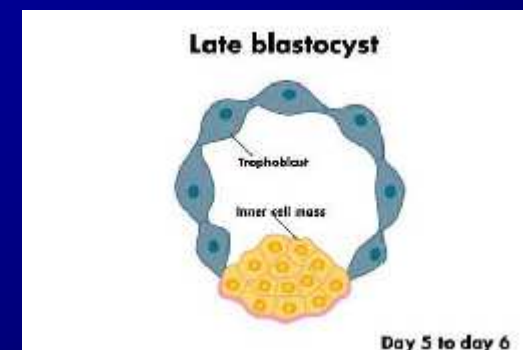
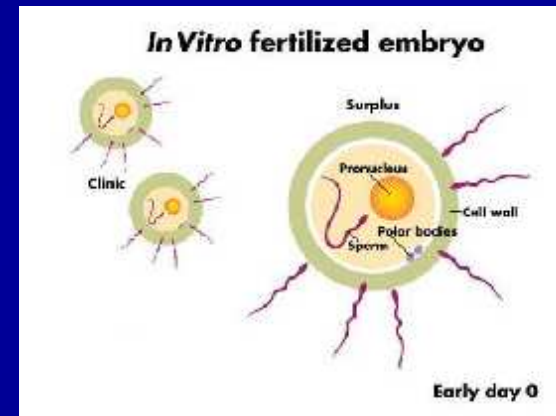
Anno II Numero 29 del 21 Febbraio 2003

**Articolo**

**Gb. E' morta Dolly, un mito, la prima pecora clonata**

# La clonazione

- Clone: dalla biologia vegetale, la talea, che può duplicare un'intera pianta (greco *clon* = germoglio)
- “Clonazione”: tecniche di laboratorio che comportano la produzione di un insieme di individui (molecole di DNA, cellule o interi organismi) derivanti per duplicazioni successive da un unico progenitore, di cui risultano copie identiche
- Febbraio 1997, Roslin Institute, Edimburgo: Ian Wilmut pubblica i risultati dell'esperimento di clonazione di un agnellino, la pecora Dolly (per trasferimento nucleare)



# Tecniche di clonazione

- Clonazione per scissione embrionale: formazione di più embrioni per separazione delle cellule, all'inizio della divisione (*embryo splitting*)
- Clonazione per trapianto/trasferimento nucleare: inserimento del nucleo di una cellula somatica in un ovocita enucleato

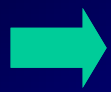
# Genetica medica: luci e ombre

- Consulenza genetica e diagnosi prenatale
- Le malattie ereditarie
- Test genetici
- La rivoluzione del progetto genoma

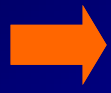


# Conclusioni – l'importanza della consulenza genetica

L'obiettivo della consulenza genetica è quello di aiutare il consultante, la coppia o la famiglia a....



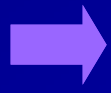
comprendere le informazioni mediche, inclusa la diagnosi, la prognosi e le terapie disponibili



rendersi conto del contributo ereditario alla malattia e del rischio di ricorrenza



prendere le decisioni che sembrano appropriate in rapporto al rischio di ricorrenza, ai progetti familiari, agli standard etici e religiosi e ad agire in accordo con queste decisioni



ottenere il miglior possibile adattamento alla malattia (in un soggetto affetto) o al rischio di ricorrenza

Ogni consulenza è un caso a sé, con problematiche diverse legate sia alla patologia sia alla particolare situazione di chi richiede la consulenza

### Chi ha coniato il termine *Bioetica*?

- 1970-71: l'oncologo americano V. R. POTTER

– prima volta del termine “BIOETICA”

- 1970 - *Bioethics. The science of survival*
- 1971 - *Bioethics. Bridge to the future*



– Di fronte al pericolo per la sopravvivenza dell'intero ecosistema, la BIOETICA come “una nuova disciplina che combinasse la conoscenza biologica con la conoscenza del sistema dei valori umani”

•Un “ponte” tra due culture: scientifica e umanistica

# TEST GENETICI : ASPETTI ETICI “GENI e SREGOLATEZZA”?

## Il mito delle perfezione

### II “PACCHETTO BAMBINO SANO” IN DIAGNOSI PRENATALE

- Effetto della riduzione della fitness Darwiniana nella popolazione Caucasica (riduzione del numero medio di figli)
  - Conseguente investimento emotivo e sociale nell'unica gravidanza
- Tendenza ad uniformare socialmente (social mimicry) il proprio bambino alle aspettative della società
  - Ruolo dei Media sulla informazione di strumenti preventivi
  - Aumento dell'offerta di test genetici > aumento della richiesta

### PROBLEMI

- Se i test genetici **non sono esaustivi** (detection rate # 99%)  
-possono comunque generare un falso senso di sicurezza nella coppia  
-non sono considerati “affidabili”





## **TEST GENETICI : ASPETTI ETICI GENI e REGOLATEZZA?**

# **Il mito della conoscenza Test genetici razionali e razionalizzati**

- esami esaustivi >> detection rate vicina al 100%**
- tecniche diagnostiche validate e armonizzate con le linee guida europee**
- abbattimento dell'errore diagnostico tramite tecniche non PCR-BASED**
- costi sostenibili**
- offerta test genetici :  
a chi?  
quando?**



# TEST GENETICI

## A CHI POSSONO ESSERE RIVOLTI E QUANDO?

### TEST GENETICI SUL PRODOTTO DEL CONCEPIMENTO

#### *PROS*

**TEST SINGOLO  
IDENTIFICAZIONE DI  
MUTAZIONI “DE NOVO”**

#### *CONS*

- INVASIVO**
- PROBLEMI INTERPRETATIVI  
(POLIMORFISMI, NUOVE  
VARIAZIONI)**
- PROBLEMI TECNICI (LEGATI  
ALL'UTILIZZO DEL  
MATERIALE FETALE)**

### TEST GENETICI SULLA COPPIA

#### *CONS*

**TEST DOPPIO**

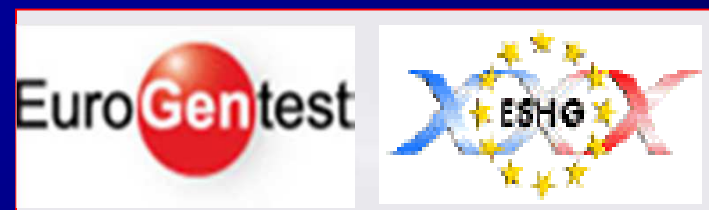
#### *PROS*

- ESEGUIBILE SIA IN EPOCA  
PRECONCEZIONALE CHE  
PRENATALE**
- NON COINVOLGE IL FETO**
  - NON IDENTIFICA  
MUTAZIONI “DE NOVO”**
- GESTIONE EMOTIVA DEL  
RISULTATO**
- GESTIONE DELLE  
CONSEGUENZE**



# LA GENETICA DEL TERZO MILLENNIO HIGHTHROUGHPUT OMIC ANALYSIS

- Harmonizing genetic testing across Europe
- Harmonizing guidelines for genetic testing across Europe



<http://www.eurogentest.org/>

# COSTI A CONFRONTO

Table 1 Second-generation DNA sequencing technologies

	Feature generation	Sequencing by synthesis	Cost per megabase	Cost per instrument	Paired ends?	1° error modality	Read-length	References
454	Emulsion PCR	Polymerase (pyrosequencing)	~\$60	\$500,000	Yes	Indel	250 bp	14,20
Solexa	Bridge PCR	Polymerase (reversible terminators)	~\$2	\$430,000	Yes	Subst.	36 bp	17,22
SOLID	Emulsion PCR	Ligase (octamers with two-base encoding)	~\$2	\$591,000	Yes	Subst.	35 bp	13,26
Polonator	Emulsion PCR	Ligase (nonamers)	~\$1	\$155,000	Yes	Subst.	13 bp	13,20
HellScope	Single molecule	Polymerase (asynchronous extensions)	~\$1	\$1,350,000	Yes	Del	30 bp	18,30

The pace with which the field is moving makes it likely that estimates for costs and read-lengths will be quickly outdated. Vendors including Roche Applied Science, Illumina, and Applied Biosystems have major upgrade releases currently in progress. Estimated costs-per-megabase are approximate and inclusive only of reagents. Read-lengths are for single tags. Subst., substitutions; indel, insertions or deletions; del, deletions.



# SCREENING GENETICI?

**PATOLOGIE MENDELIANE PROPONIBILI PER SCREENING PRECONCEZIONALI O PRENATALI DI COPPIA IN BASE ALL'INICIDENZA**

**AMIOTROFIA SPINALE (SMA) 1:10.000**

**FIBROSI CISTICA 1:2.000-1:6.000**

**TALASSEMIA 1/250 \*\***

**SORDITA' EREDITARIE (TUTTE CIRCA 35 GENI) 1/1000 nati**

**DISTROFIE MUSCOLARI (TUTTE, CIRCA 70 GENI) 1:10.000**



# E LE PATOLOGIE POLIGENICHE?

MAJOR GENES

GENE SUSCETTIBILITA'

POTENZIALMENTE COINVOLGONO CIRCA IL 50% DELLA POPOLAZIONE

**MA A TUTT'OGGI**

DATI SCIENTIFICI NON SUFFICIENTI PER LA PREDICIBILITA' DEL FENOTIPO

OCCORRE UNA QUANTIFICAZIONE DELLA PENETTRANZA ACCURATA  
CHE SIA IN GRADO DI STABILIRE  
IL REALE RISCHIO DI SVILUPPARE LA PATOLOGIA



## Se i geni sono regolati...

- Come “gestire” i test genetici “senza sregolatezza”??
- La Genetica medica deve avere delle linee guida?
- I test genetici devono essere offerti a tutti (principio di autonomia)?
- Quali sono i costi sostenibili per i test genetici?
- Può il sistema sanitario nazionale sostenere questi costi?
- Se sì, per quali “livelli di rischio”?
- Quale rapporto fra pubblico e privato?
- RIFLESSIONI.....

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- Quali sono i costi sostenibili per i test genetici?
- Ruolo dei LEA (livelli essenziali di assistenza) nell’offerta SSN dei test genetici
- Può il SSN sostenere questi costi?
- Se sì, per quali “livelli di rischio”?
- Quale rapporto fra pubblico e privato?
- RIFLESSIONI.....

# GENI E SREGOLATEZZA

- STIAMO ANDANDO VERSO....
- **Capacità di sequenziare l'intero genoma di 1 individuo (300 milioni di basi) in 48 ore con un costo di 1000 euro**
- CHE RIFLESSIONI FARE?
- QUALI LE RIPERCUSSIONI SULLA SANITA' PUBBLICA?
- QUALI RIPERCUSSIONE SULLA BIOETICA?