



Museo Storico della Città
Martedì 2 Giugno 2015
Lushnja, Ore 10

CONFERENZA SCIENTIFICA SULLA TALASSEMIA

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Hemoglobin Structure

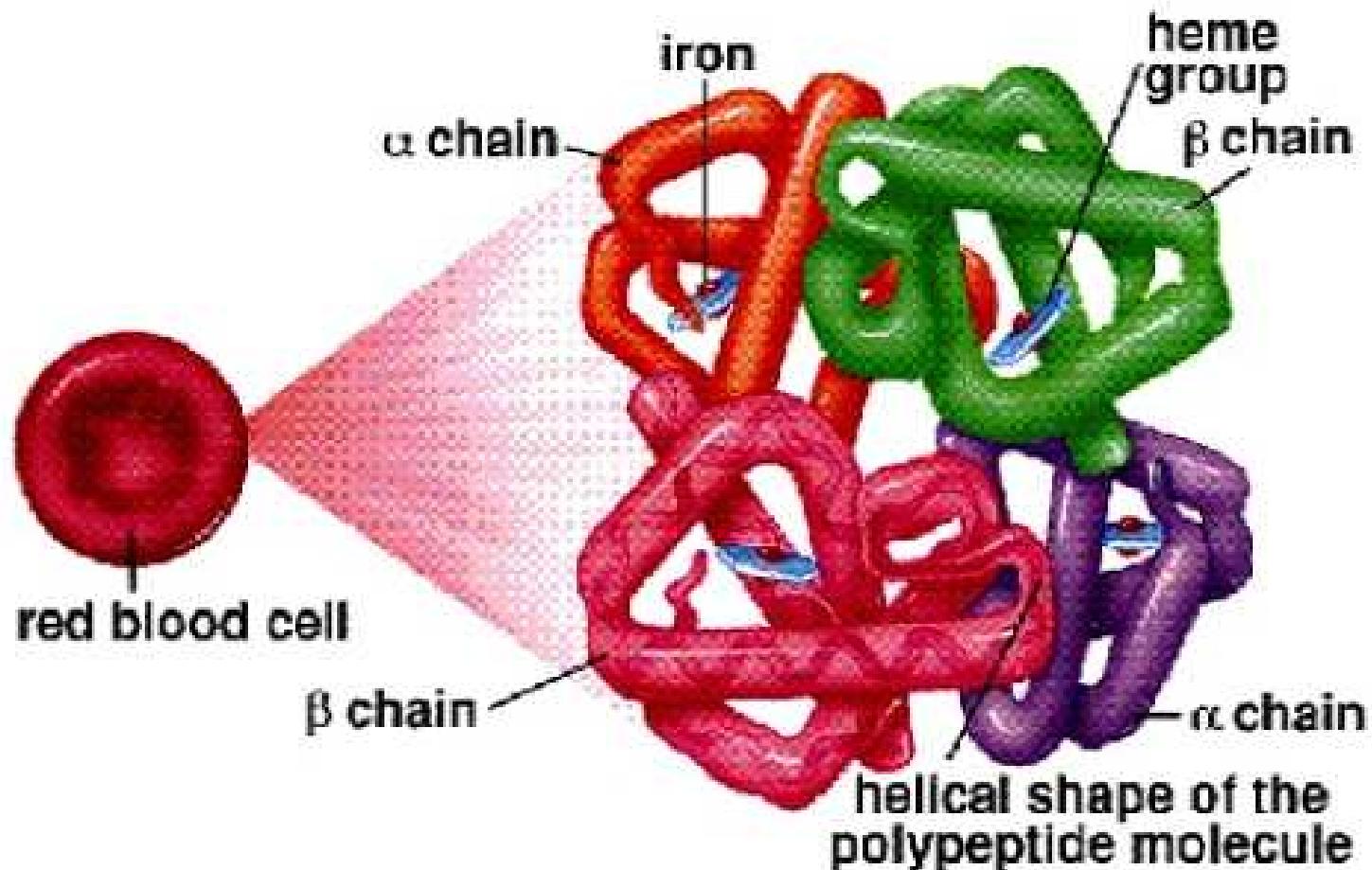
- Four subunits

- two α
 - two β

- Iron

- Heme

- Binds 4 O₂

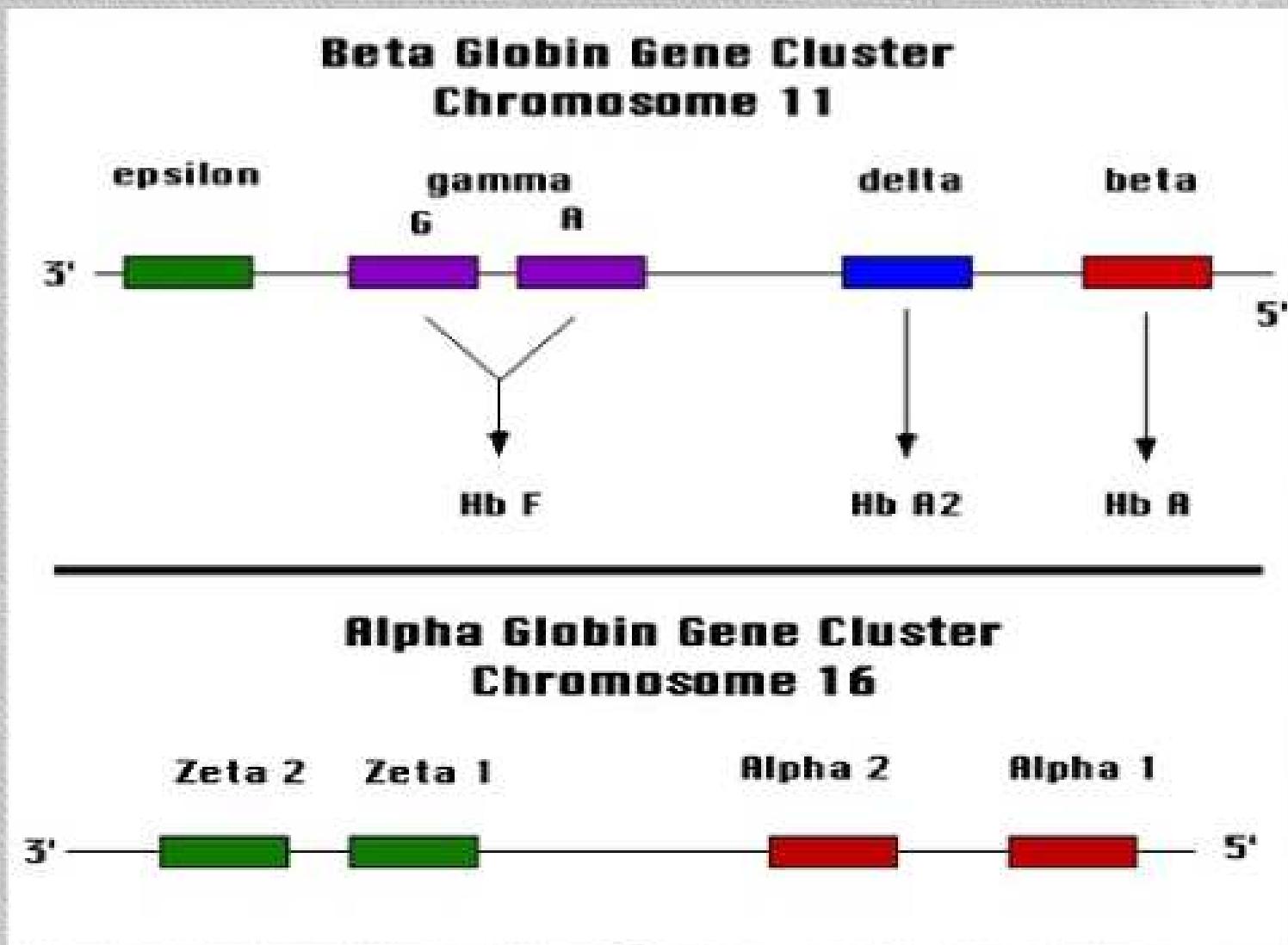


Emoglobine fisiologiche nell'uomo

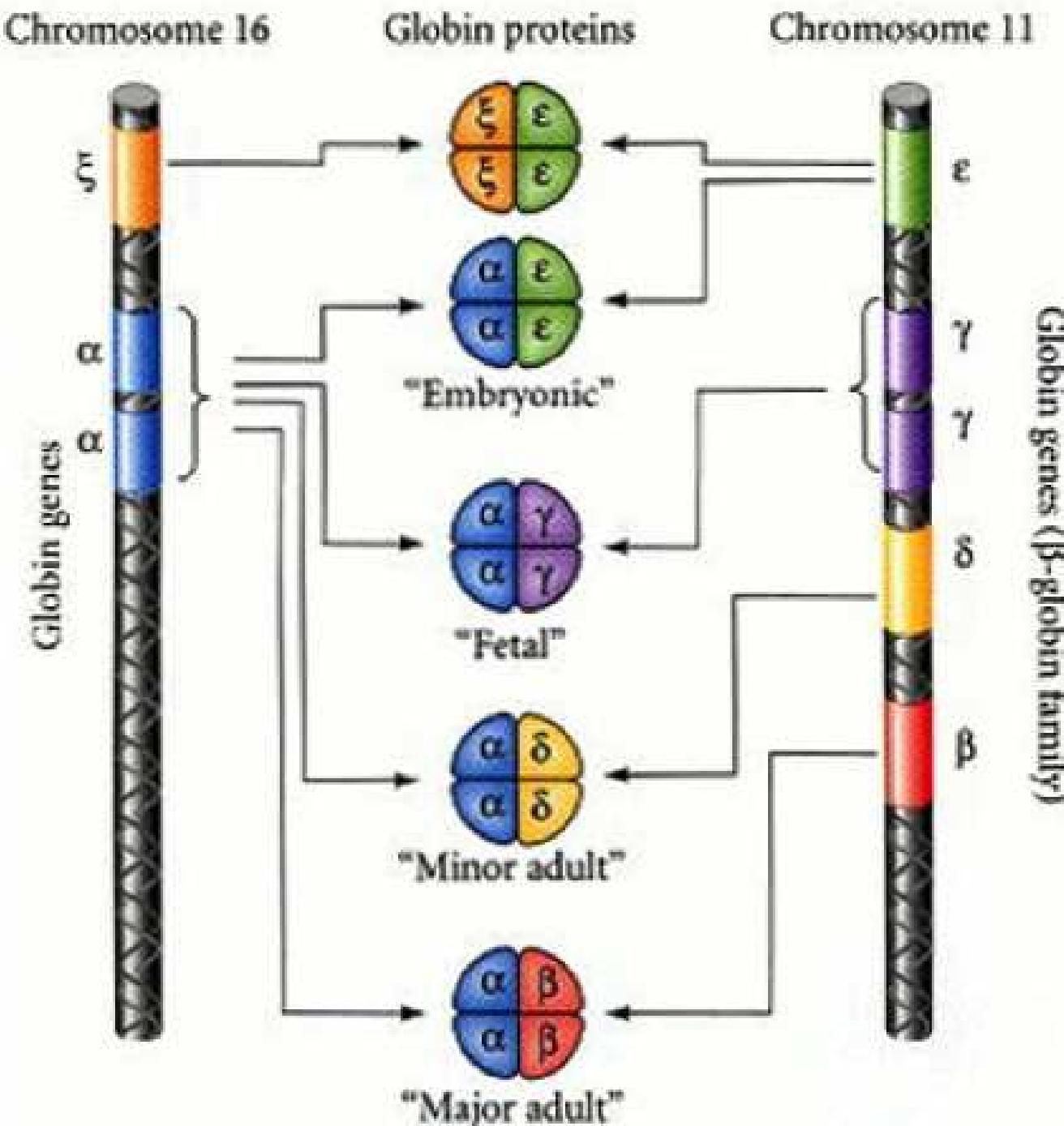
Haemoglobin	Globin chains	Period of life when present
Gower 1	$\zeta_2\epsilon_2$	Embryo
Gower 2	$\alpha_2\epsilon_2$	Embryo
Portland 1	$\zeta_2\gamma_2$	Embryo
Haemoglobin F	$\alpha_2\gamma_2$	Embryo, fetus and neonate; minor component during adult life
Haemoglobin A	$\alpha_2\beta_2$	Minor component in fetus, increasing late in gestation and in the neonatal period to become the major haemoglobin during infancy, childhood and adult life
Haemoglobin A ₂	$\alpha_2\delta_2$	Very low levels in infancy; minor component in childhood and adult life

Geni delle catene globiniche

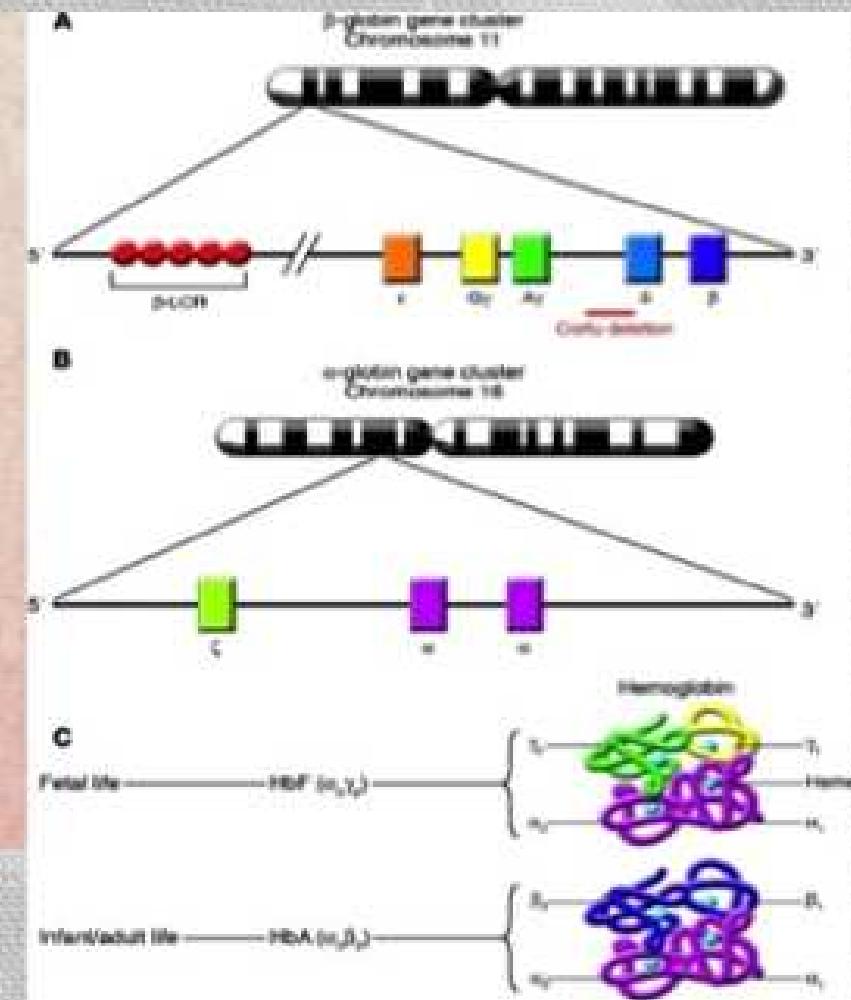
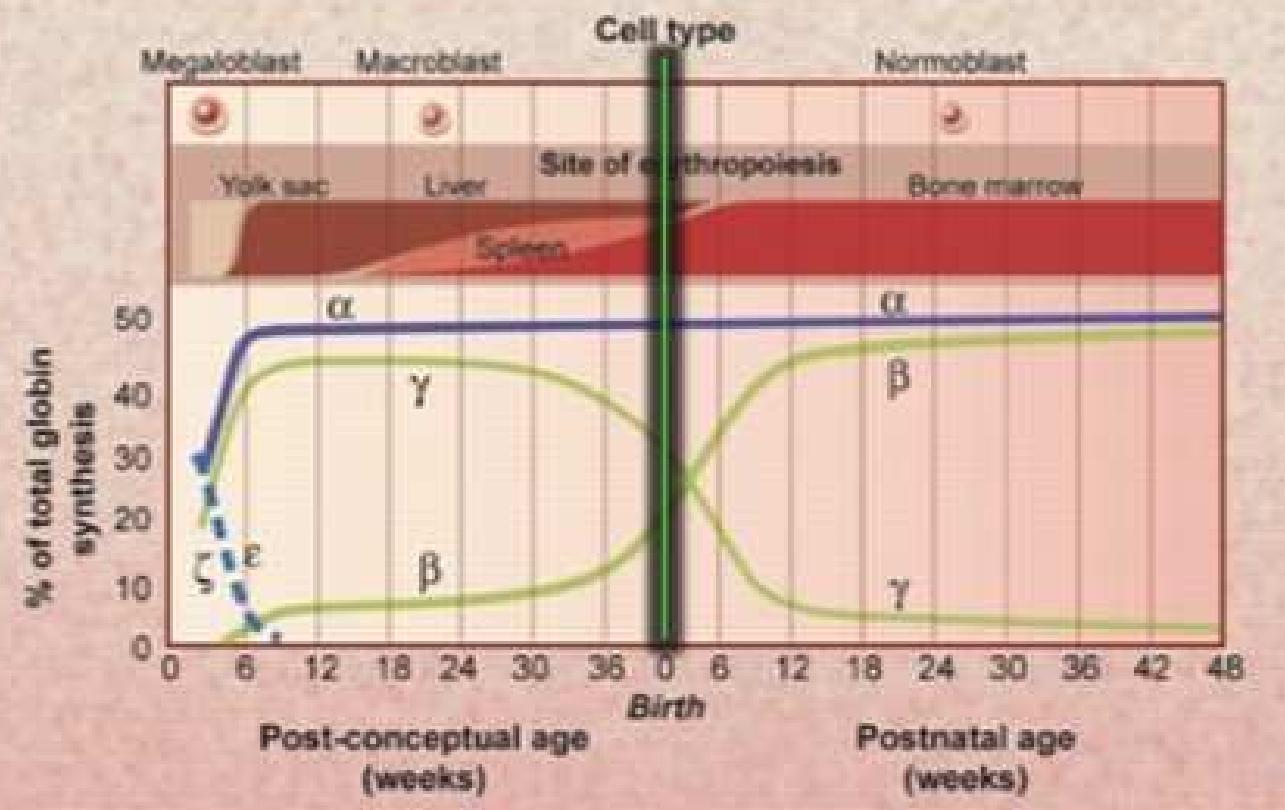
Chr. 11



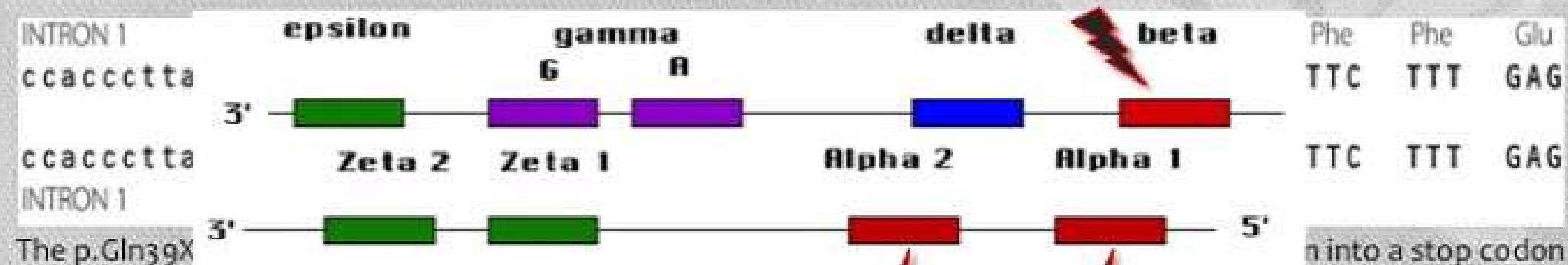
Dai Geni alle Globine



Un emoglobina per ogni età...



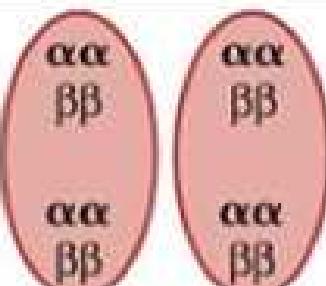
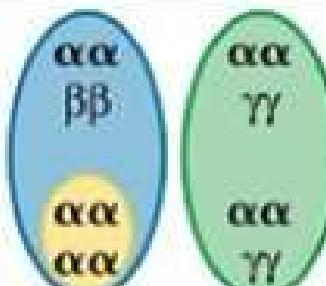
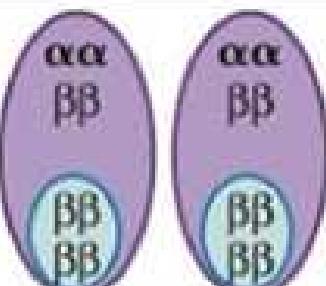
Mutazioni nei geni globinici → fenotipo



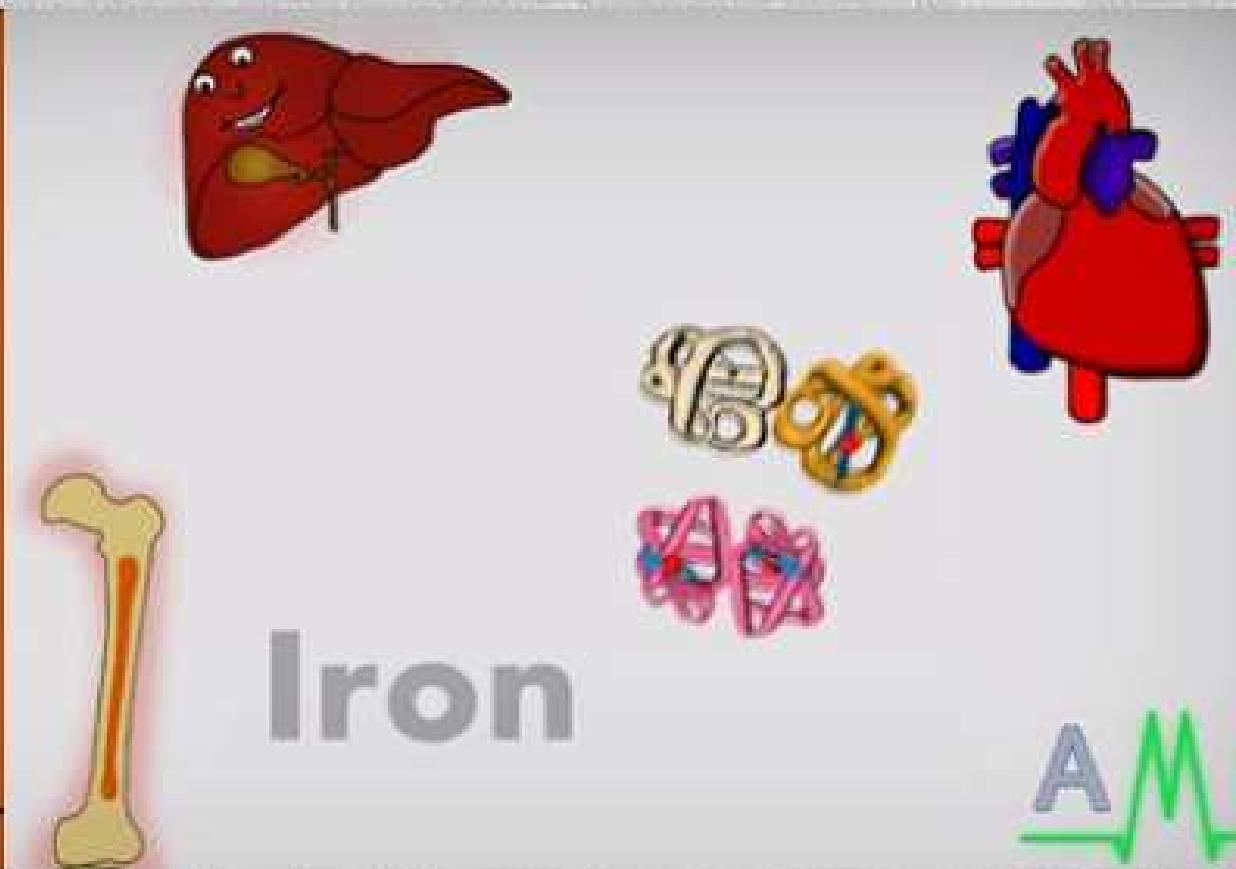
Diverse mutazioni → Diverse globine

- **Hemoglobin S** ($\alpha_2\beta^S$, severe). This is the predominant hemoglobin in people with sickle cell disease.
- **Hemoglobin Constant Spring** named after isolation in a Chinese family from the Constant Spring district of Jamaica (severe). In this variant, a mutation in the alpha globin gene produces an alpha globin chain that is abnormally long. Both the mRNA and the alpha chain protein are unstable.
- **Hemoglobin H** (β_4 , mild). This is a tetramer composed of four beta globin chains: it occurs only with extreme limitation of alpha chain availability. Hemoglobin H forms in people with three-gene alpha thalassemia as well as in people with the combination of two-gene deletion alpha thalassemia and hemoglobin Constant Spring.
- **Hemoglobin Barts** (γ_4 , lethal). With four-gene deletion alpha thalassemia no alpha chain is produced. The gamma chains produced during fetal development combine to form gamma chain tetramers. Individuals with four-gene deletion thalassemia and consequent hemoglobin Barts die in utero (hydrops fetalis).

Alterata sintesi delle globine - patogenesi

	Normal	Severe β thalassemia	Severe α thalassemia (hemoglobin H disease)
Globin production	Balanced α and β production	Decreased β production	Decreased α production
Bone marrow		 <p>Excess α damages precursors in marrow causing ineffective erythropoiesis</p>	
Peripheral blood	Normal hemoglobin composition (97% hemoglobin A)	Selective survival of cells producing hemoglobin F	Hemoglobin H detectable in blood. Hemoglobin H precipitates and damages RBCs, causing hemolysis

Alterata sintesi delle globine - patogenesi



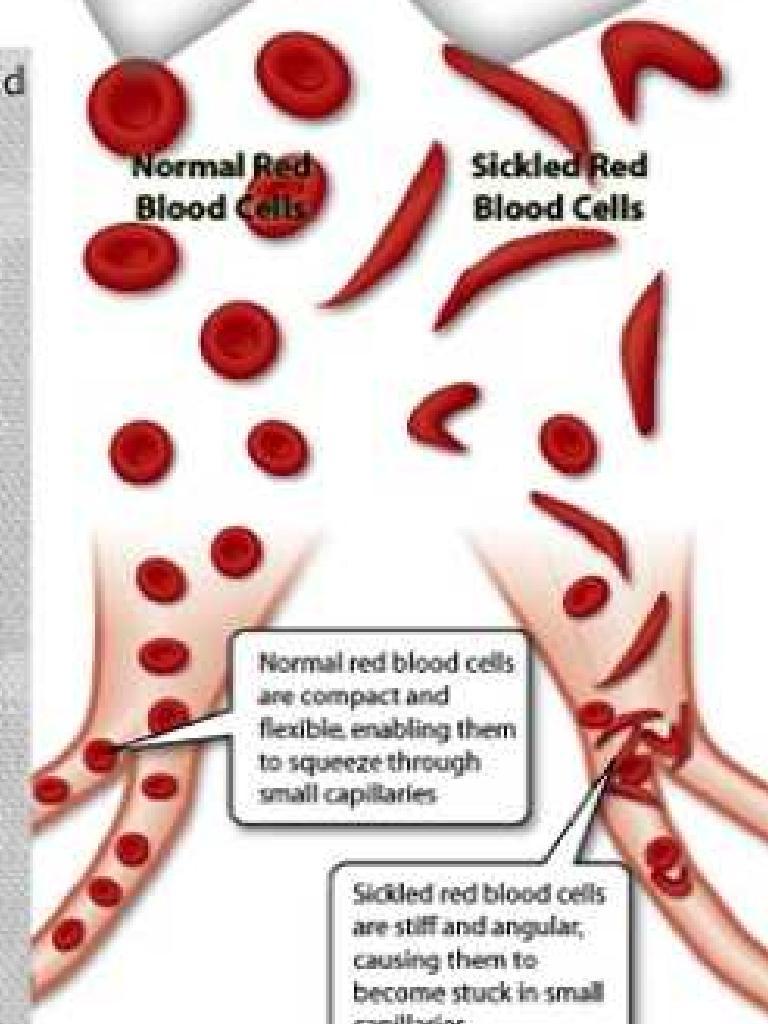
About Sickle Cell



	Val	His	Leu	Thr	Pro
NORMAL	GTG	CAT	CTG	ACT	CCT
SICKLE	GTG	CAT	CTG	ACT	CCT
	Val	His	Leu	Thr	Pro

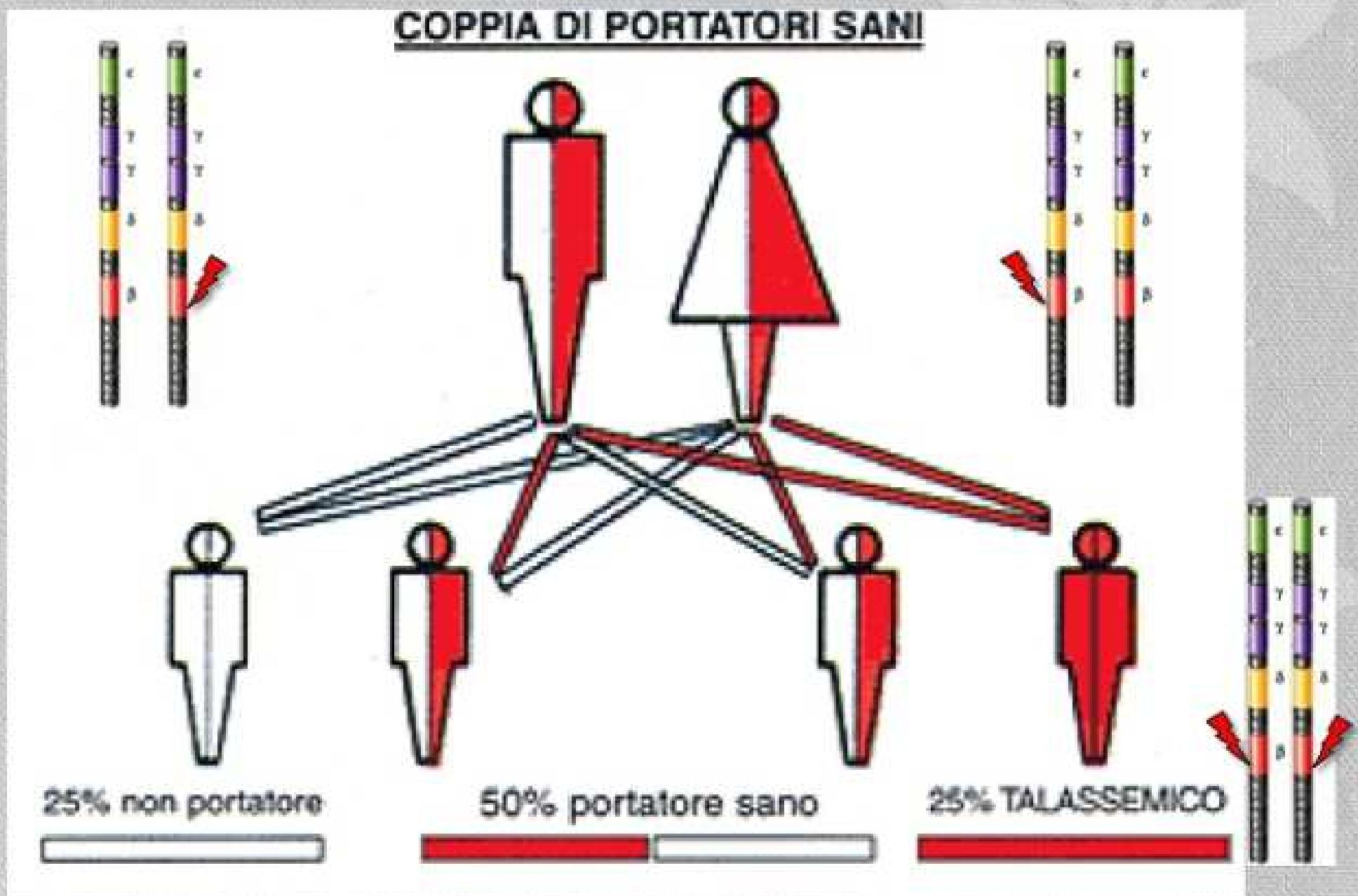
In sickle cell disease an A>T nucleotide substitution causes glutamic acid to be replaced by valine in the β -globin protein.

Anemia falciforme - HbS



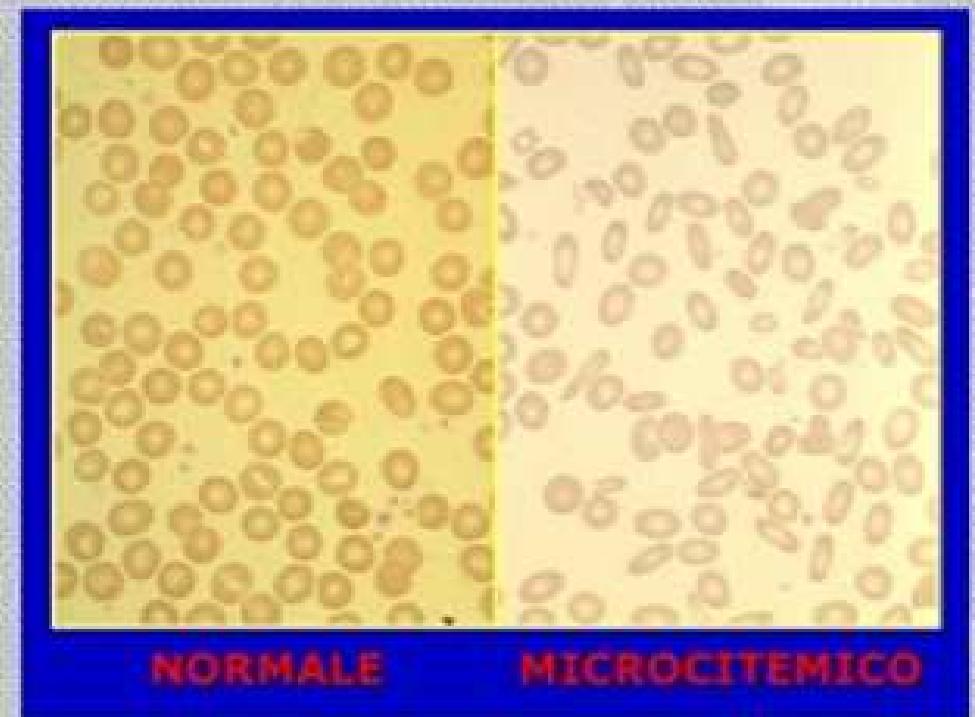
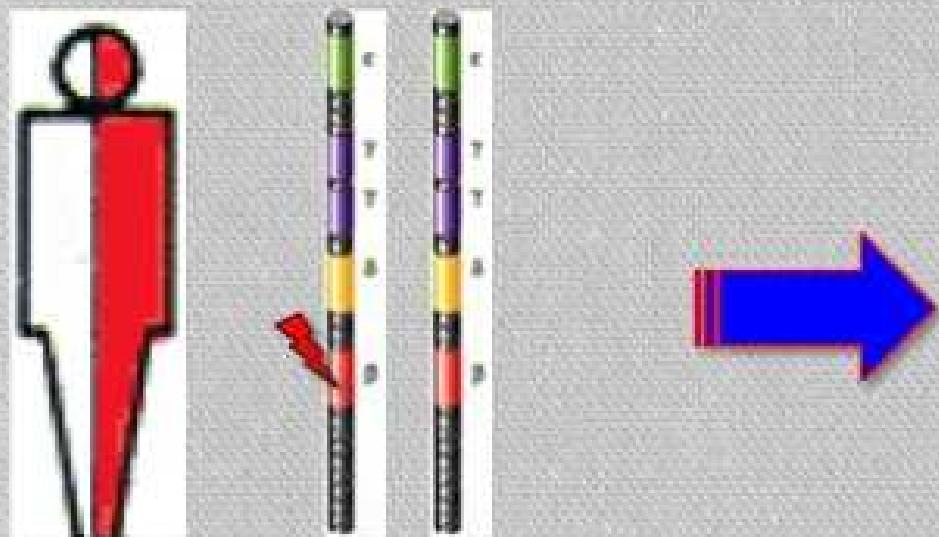
Sickle Cell Anemia

Modello di ereditarietà della talassemia



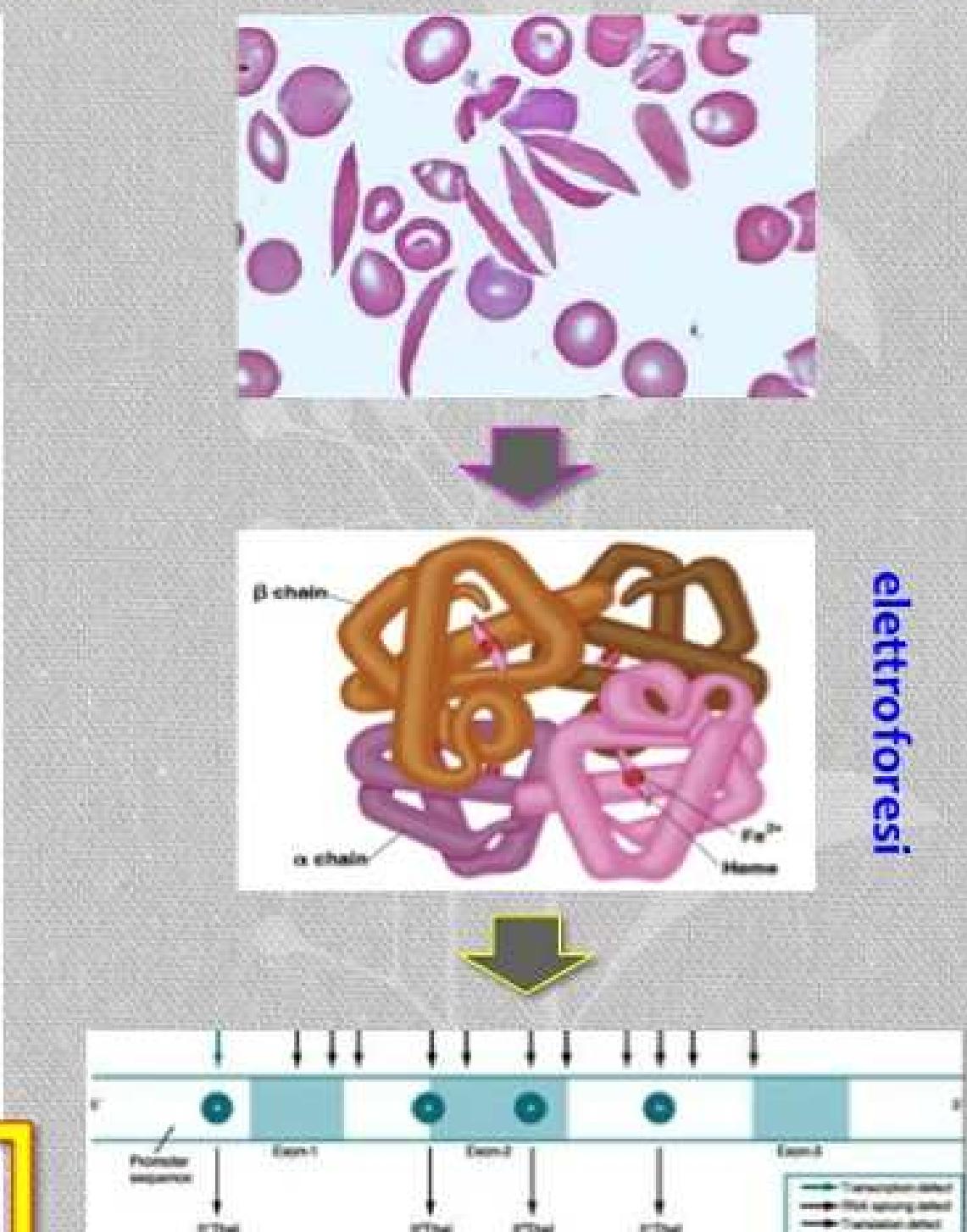
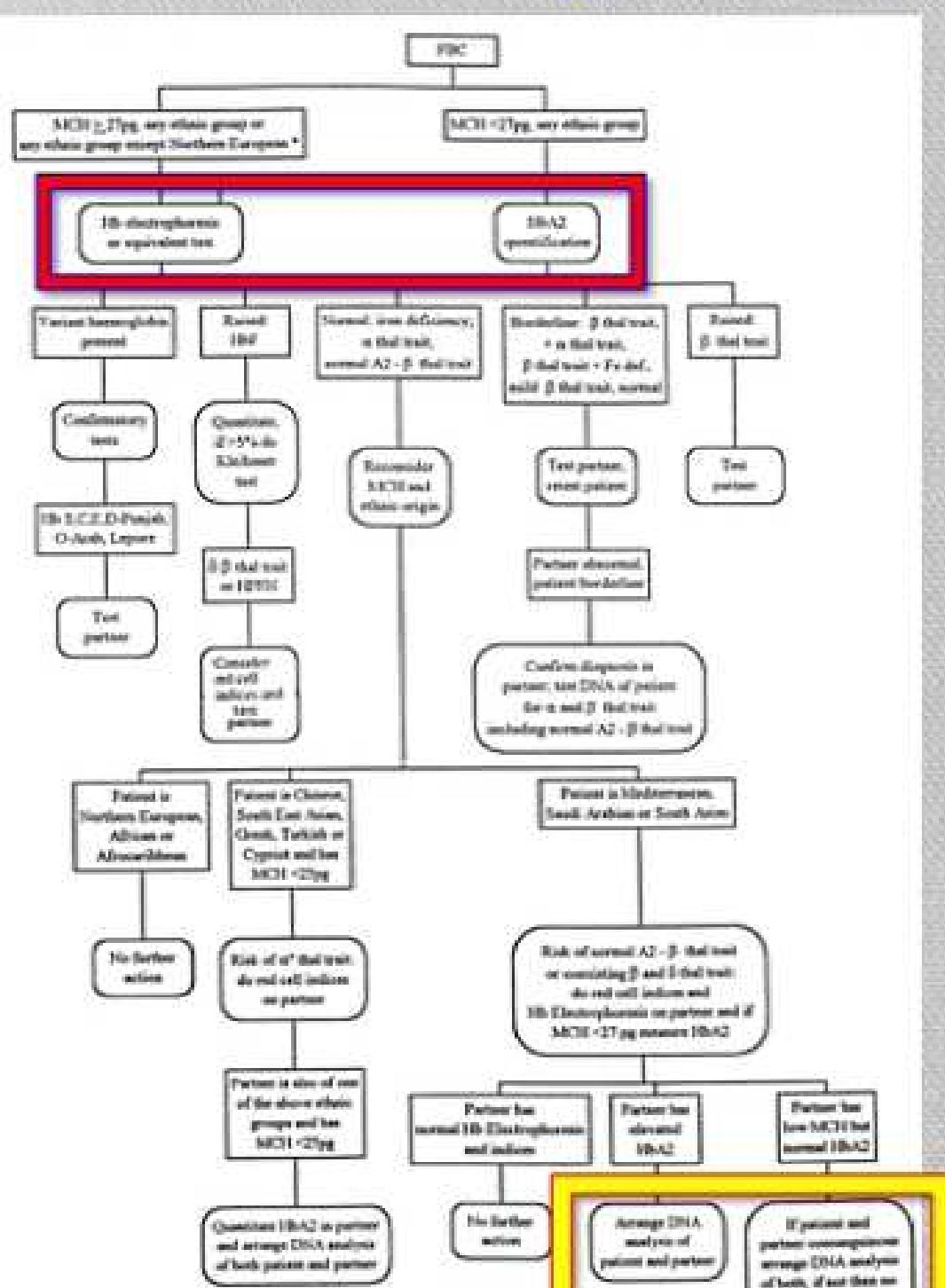
Endofenotipo talassemia: microcitemia

Fenotipo cellulare



Possibile identificare i portatori sani mediante test NON GENETICI !

Flow-chart diagnostica no genetics without chemistry!



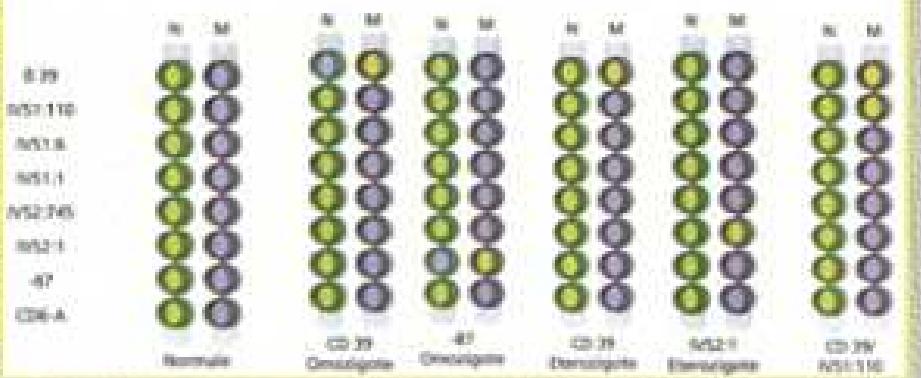
Analisi genetica: non una ma diversi tipi

Table 5.1 The principal methods of DNA diagnosis of the haemoglobinopathies.

DISORDER AND MUTATION TYPE	DIAGNOSTIC METHOD
α^0 -thalassæmia	Gap-PCR, MLPA
α^+ -thalassæmia: deletion non deletion	Gap-PCR, MLPA ASO, RE, DGGE, Sanger sequencing
β -thalassæmia: deletion non deletion	Gap-PCR, MLPA ASO, RDB, ARMS, RE-PCR, Sanger sequencing
$\delta\beta$ -thalassæmia	Gap-PCR, MLPA
HPPH deletion non deletion	Gap-PCR, MLPA ASO, ARMS, RE-PCR, Sanger sequencing
Hb Lepore	Gap-PCR, MLPA
HbS	ASO, RDB, ARMS, RE-PCR, pyrosequencing
HbC	ASO, RDB, ARMS, pyrosequencing
HbE	ASO, RDB, ARMS, RE-PCR, pyrosequencing
Hb D-Punjab	ASO, RDB, ARMS, RE-PCR, Sanger sequencing
Hb O-Arab	ASO, ARMS, RE-PCR, Sanger sequencing
Hb variants	Sanger sequencing

Reverse dot-blot

La rivelazione dell'avvenuta ibridazione attraverso una reazione enzimatico-colorimetrica (colore giallo), indica quali sequenze (normali o mutate) sono presenti nel campione.



(...)

Sequenziamento diretto

$\alpha 2$ -globin gene →

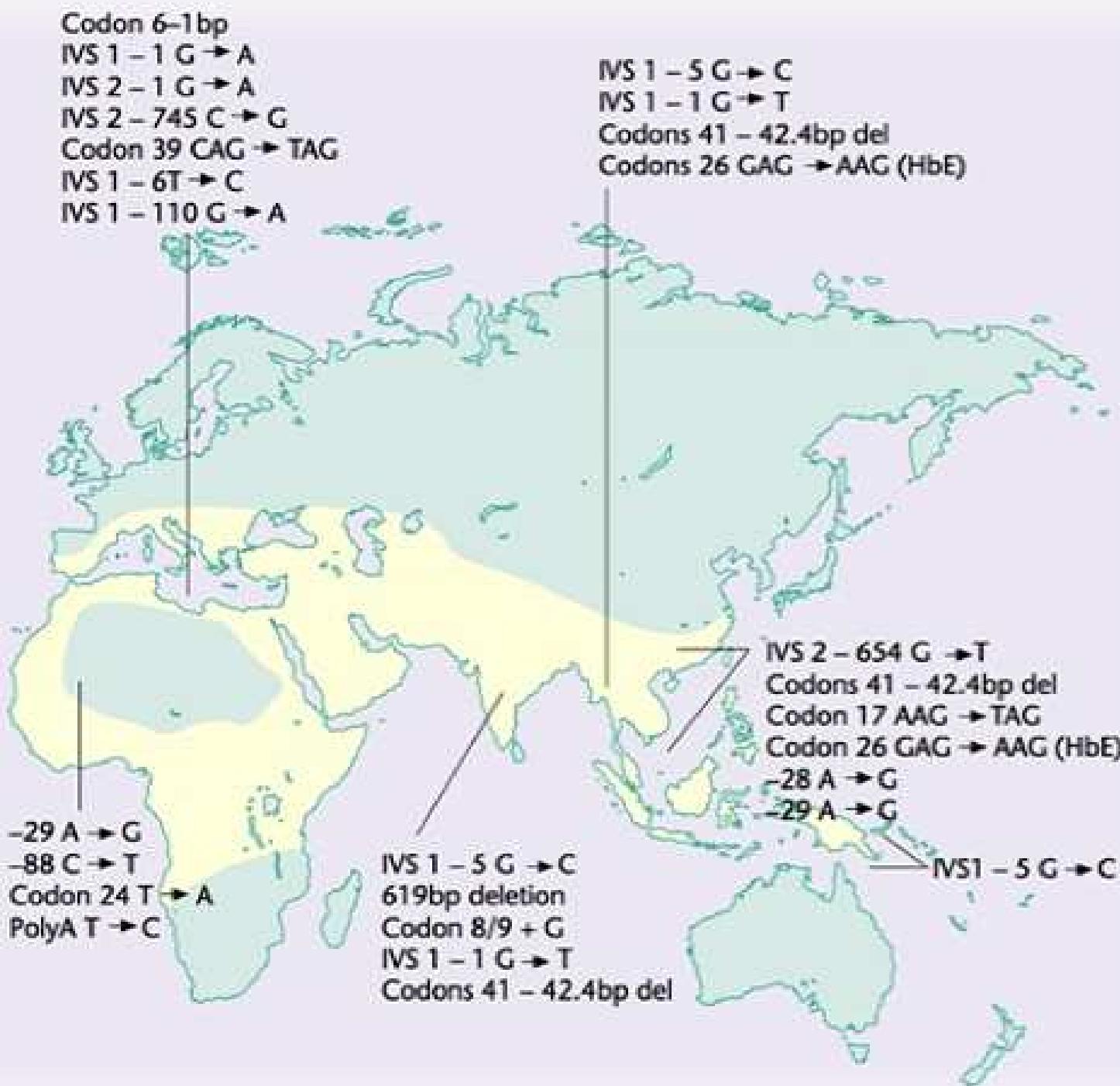
ATTCCTGTCCTTCCCCCCCCAAAC TACT TTCCCCCTTC CAACTG AACG
35 36 37 38 39 40 41 42 43 44 45 46 47 48 49

Normal: TCC TTC CCC ACC AAAG ACC TAC TTC CGG CAC TTC GAC CTG AGC

CRM37-C: TCC TTC CCC CCA AGA CCT ACT TCC CGC ACT TGG ACC TGA GGC
Stop

(C)

Distribuzione nel mondo delle mutazioni della talassemia



Letteratura Talassemia – Falcemia (Albania)

Boletini E, Svobodova M, Divoky V, Curuk M, Dimovski A, Liang R, Adekile AD, Huisman THJ.
Sickle cell anemia, sickle cell β -thalassemia, and thalassemia major in Albania:
characterization of mutations. Hum Genet. 1994; 93: 182-187.

Angioletti M, Lacerra G, Boletini E, Di Noce F, Musollino G, Carestia C.
 β - and α -globin genotypes in Albanian patients affected by β -globin disorders.
Haematologica 2002; 87(9): 1002-1003.

Mokini V, Duka D, Rosatelli C, Tuveri T, Demurtas M, Babameto-Laku A, Cao A.
Molecular characterization of β -thalassemia mutations in Albania. The UNEPSA and European Congress of Paediatrics, 2000. Abstract Book, Haematology and Oncology. 2000; HO-265: 143.

Babameto-Laku A, Mitre A, Berisha S, Mokini V, Roko D
Molecular genetic characterization of β -thalassemia and sickle cell syndrome in the Albanian population. Balkan J Med Genet 2011; 14: 45–50.

Letteratura Talassemia – Falcemia (Albania)

#	β - and/or α -globin genotypes	Patients no. and age (yr)	Transmission	
A	β -BS+110 homozygote	0/0/0/0	5 (10m, 3) 3 (14, 3)	regular
	β -cod 38/ β -BS+1	0/0/0/0	1 (17)	sporadic
	β /-	0/0/0/0	1	regular
	β homozygote	0/0/0/0	2 (14, 6)	sporadic
	β cod 38/ β	0/0/0/0	1	regular
	β -BS+110/ β	0/0/0/0	3 (16, 15, 7) 1 (10)	sporadic regular
	β -BS+110/ γ	0/0/0/0	7 (24, 27, 12, 6) 1 (8)	sporadic regular
B	β -BS+110/ β cod 38	0/0/0/0	4	regular
	β -BS+110/ β cod 44	0/0/0/0	2	regular
	β -BS+110 homozygote	0/0/0 ^{BS+110} /0/0	1	regular
	β -cod 38 homozygote	0/0/0/0	1	regular
	β -BS+110/ β cod 82-83	0/0/0/0	1	regular
	β -cod 38/ β cod 5	0/0/0/0	1	regular
	β -cod 38/ β -BS+1	0/0/0/0	1	regular
	β -BS+74S/ β -BS+1	0/0/0/0	1	regular
	β -BS+4/ β cod 37	0/0/0/0	1	regular
C	β -BS+4/ β -BS+110	0/0/0/0	5 (8, 13, 15, 3)	sporadic
	β -BS+4 homozygote	0/0/0/0	2 (7, 4)	sporadic
	β -poly A/ β -BS+110	0/0/0/0	2	sporadic
	β -BS+4/ β cod 38	0/0/0/0	1 (15)	sporadic
	β -BS+4/ β cod 44	0/0/0/0	1 (4)	sporadic
	β -cod 38 homozygote	0 ^{BS+110} /0/0/0	1 (10)	sporadic
	β /-	0/0/0/0	2 (23, 24)	sporadic
	β -BS+1/ β	0/0/0/0	1	sporadic
D	β -cod 38/ β	0/0/0 ^{BS+110} /0/0	1 (9)	none
	β -cod 44/ β	0/0/0 ^{BS+110} /0/0	1 (45)	none
	β / β	0/0/0/0	1 (2)	none
	β -BS+110/ β	0/0/0/0	1	none
	β -BS+4/ β	0/0/0/0	1	none

... alta eterogeneità allelica nei pazienti Albanesi con emoglobinopatie...

Angioletti M b- and α -globin genotypes in Albanian patients affected by b-globin disorders.
Haematologica 2002; 87(2): 1622-1629.

Studi pilota in Albania - Lushnjë

Acta
Haematologica

Accepted manuscript / Accepted manuscript / DOI: 10.1160/000123456-2009

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A Pilot Beta-Thalassaemia Screening Program in the Albanian Population for a Health Planning Program

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Federica Sangiuliano^{a,c} Vane Molkin^b Giuseppe Novelli^{a,c}

217 students
Saliva DNA extraction (DNA only)

Results:

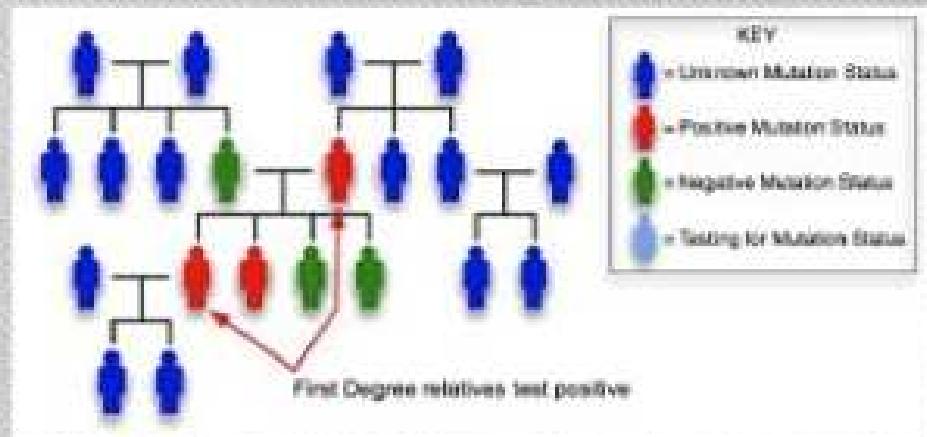
- | | |
|-------------------|------|
| - HbS c.20A→T | 3.2% |
| - IVS-I-110 (G→A) | 1% |

DNA only (selected mutations)

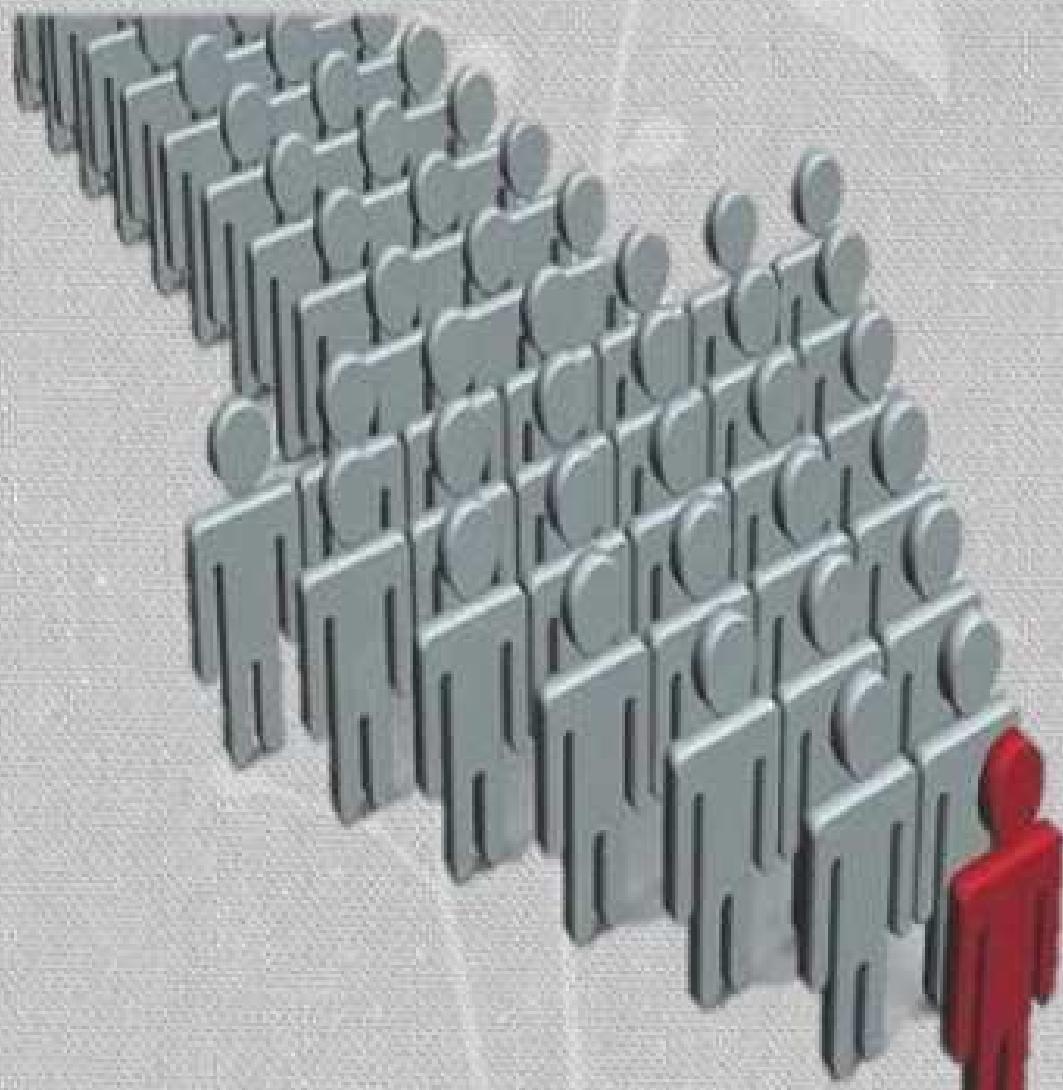


Strategie di SCREENING genetico

a cascata



di popolazione



CONSULENZA GENETICA



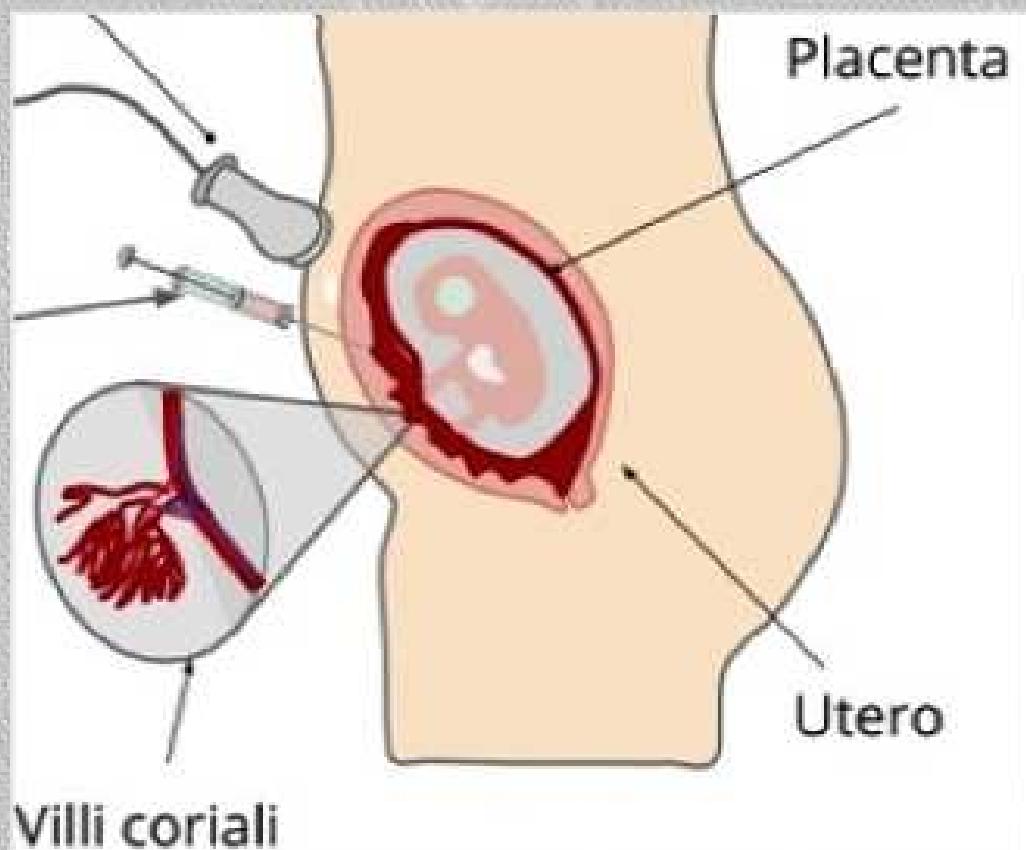
Diagnosi prenatale

*... ricerca delle specifiche mutazioni
CONFIRMATE NEI GENITORI su prelievo di villi
coriali a 11 settimane di gestazione...*

TeleGenetics



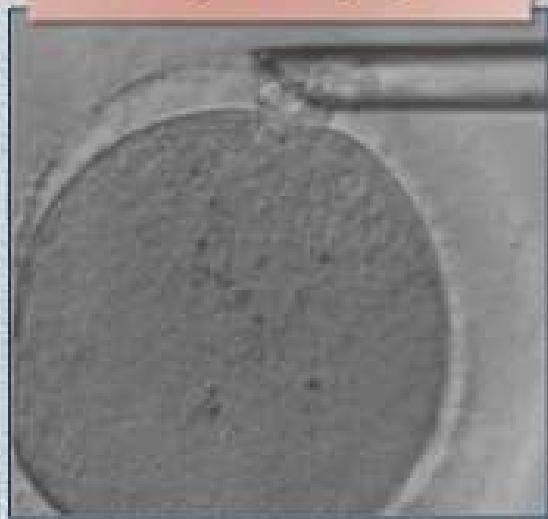
LABORATORIO ALTA
SPECIALIZZAZIONE



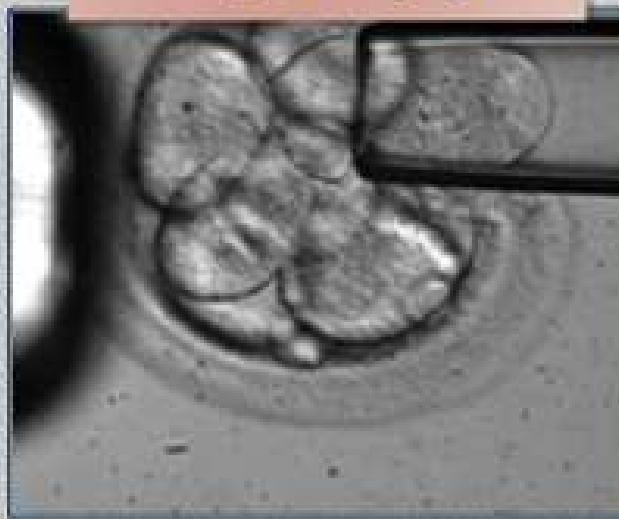
Diagnosi Genetica Preimpianto

progressi, ricerca e tecnologia

Day 0 biopsy



Day 3 biopsy



Day 5 biopsy



- Paternal and post-zygotic errors not detected
- Need of 2 PB biopsy
- High false positive diagnostic rate
- Impact on embryo development
- Most expensive and time-consuming approach

- High worldwide experience
- Small reduction in embryo viability
- High impact of mosaicism
- Single cell analysis

- More robust genetic analysis
- No impact of biopsy
- Low impact of mosaicism
- Reduced number of embryos/cycles
- Less expensive

PGD: take home message

Ginecologo



Non laboratori ma Servizi di Medicina della Riproduzione

Effetti dello SCREENING genetico

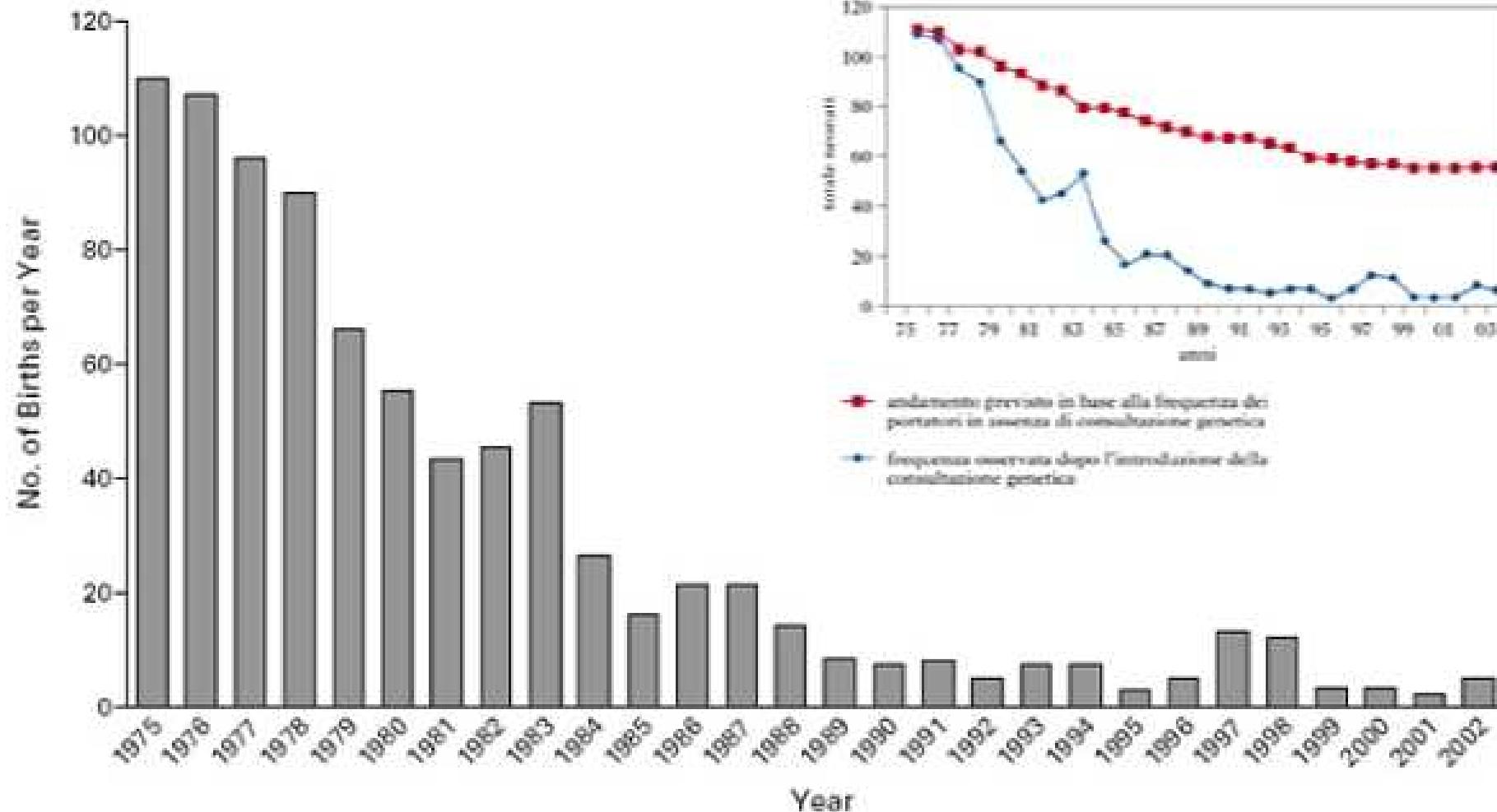


Figure 1. Declining Rate of Birth of Infants Homozygous for β -Thalassemia in Sardinia since 1975, When the Screening Program Began.

Grazie!