

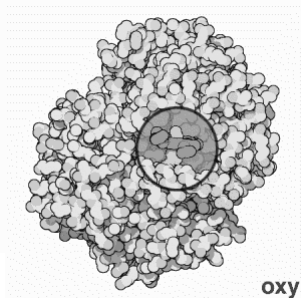
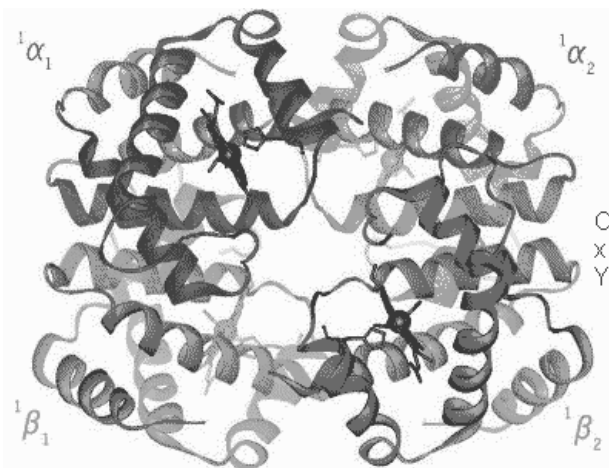
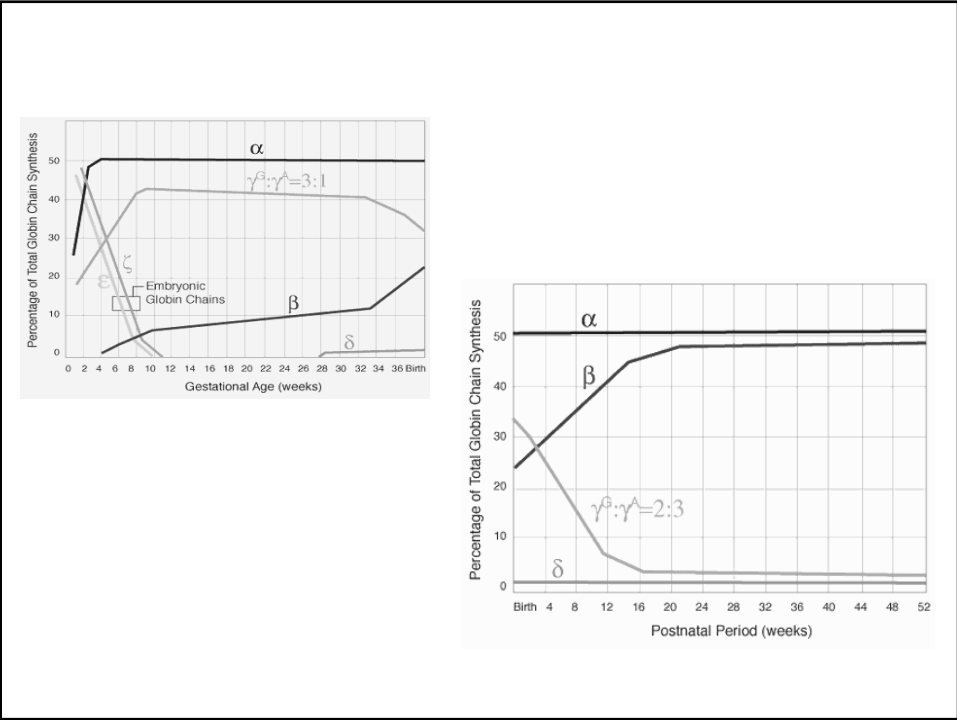
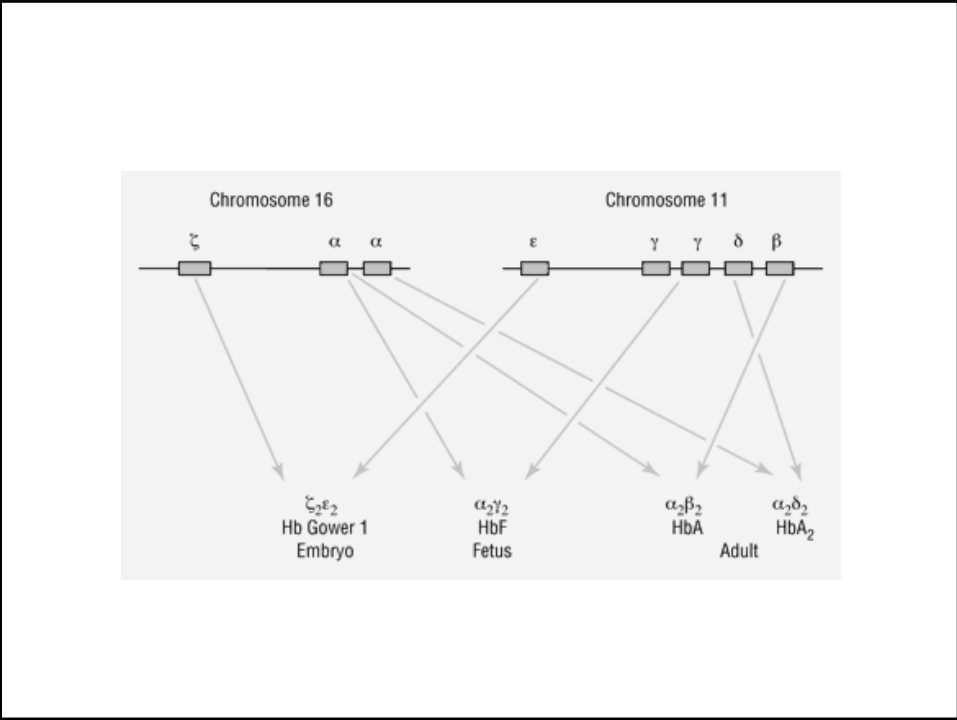
  
 TOSOH  
**INCONTRO DI AGGIORNAMENTO  
 SCIENTIFICO**  
**"EMOGLOBINE: DIAGNOSTICA,  
 STANDARDIZZAZIONE,  
 PROSPETTIVE"**  
**AULA MAGNA OSPEDALE  
 DESENZANO DEL GARDA (BS)**  
**Loc. Montecroce**  
**9 MAGGIO 2009**





**Epidemiologia delle varianti emoglobiniche e  
 della Talassemia**  
*G. Lippi – M. Montagnana (Verona)*





- Hemoglobin variants are a part of the normal embryonic and fetal development, but may also be pathologic mutant forms of hemoglobin in a population, caused by variations in genetics.
- **In the embryo:**
  - Gower 1 ( $\zeta\epsilon\epsilon 2$ )
  - Gower 2 ( $\alpha 2\epsilon 2$ ) (PDB 1A9W)
  - Hemoglobin Portland ( $\zeta 2\gamma 2$ )
- **In the fetus:**
  - Hemoglobin F ( $\alpha 2\gamma 2$ ) (PDB 1FDH)
- **In adults:**
  - Hemoglobin A ( $\alpha 2\beta 2$ ) - The most common (normal amount ~95%)
  - Hemoglobin A2 ( $\alpha 2\delta 2$ ) -  $\delta$  chain synthesis begins late in the third trimester and in adults (normal range 1.5-3.5%)
  - Hemoglobin F ( $\alpha 2\gamma 2$ ) - Restricted to a limited population of red cells called F-cells (normal range 0.3-2.0%).



## DEFINITION, EARLY KNOWLEDGE AND CAUSES OF BIRTH DEFECTS

### Definition

A birth defect is defined as any abnormality affecting body structure or function that is present from birth. It may be clinically obvious at birth or may be diagnosed only later in life. For example, spina bifida is a structural birth defect clinically obvious at birth and hemophilia is a functional birth defect that may present clinically only in infancy or childhood. A few birth defects, like Huntington disease, manifest only in adulthood. Serious birth defects are life-threatening or have the potential to cause lifelong disability (Christianow RE et al., 1981; WHO, 2000b).

### MARCH OF DIMES GLOBAL REPORT ON BIRTH DEFECTS THE HIDDEN TOLL OF DYING AND DISABLED CHILDREN

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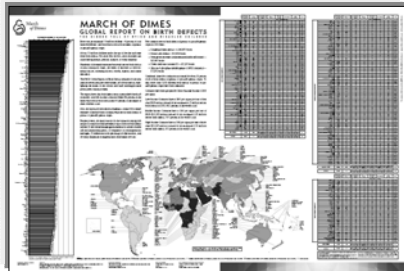
### HEMOGLOBIN DISORDERS

Thalassemia and sickle cell anemia are autosomal recessive disorders that became very common in tropical regions because of heterozygote advantage—carriers are protected against the lethal effects of falciparum malaria. These disorders spread through migration of carriers to other regions. Carriers can be detected by simple hematological tests and global data exist on carrier frequencies (Livingstone, 1985).

The characteristic skeletal changes of Thalassemia have allowed archaeologists to identify the presence of the condition as far as back as the Neolithic period. It must have been very common even at that time.



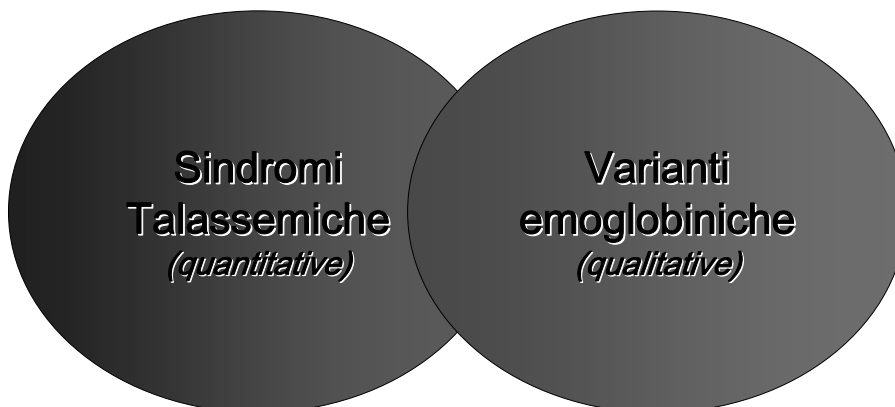
**Choirokoiitia 7000 BC: Over 150 graves, 47% of children died of thalassaemia. The oldest thalassaemic skeleton in Cyprus was of a child who lived about 8300 BC. Kissonerga, Paphos.**



Five common serious birth defects of genetic or partially genetic origin in 2001 were:

- ❖ Congenital heart defects—1,040,835 births
- ❖ Neural tube defects—323,904 births
- ❖ Hemoglobin disorders (thalassemia and sickle cell disease)—307,897 births
- ❖ Down syndrome (trisomy 21)—217,293 births
- ❖ Glucose-6-phosphate dehydrogenase (G6PD) deficiency—177,032 births

	Children Born with Birth Defects Annually <sup>1</sup>	2001 Annual Births (000s)	Birth Defects of the Cardiovascular System <sup>2</sup>	Neural Tube Defects	Prevalence (per 1000 live births)			Total <sup>3</sup>
					Pathological Hemoglobin Disorders	Down Syndrome	G6PD Deficiency <sup>4</sup>	
Italy	21,838	505	7.9	0.5	0.6	1.9	0.0	43.2



## SINDROMI TALASSEMICHE

- Delezioni di segmenti genomici (anche HPFH o High Persistence of Fetal Hemoglobin e Hb Lepore).
- Mutazioni che modificano la trascrizione dei geni globinici
- Mutazioni che modificano la corretta maturazione dell'mRNA
- Mutazioni che alterano i siti di splicing e attivano siti criptici.
- Mutazioni che creano siti di splicing alternativi.
- Mutazioni che alterano la traduzione dell'mRNA

## VARIANTI EMOGLOBINICHE

- Mutazioni che alterano la sequenza aminoacidica della beta-globina (es. anemia falciforme)

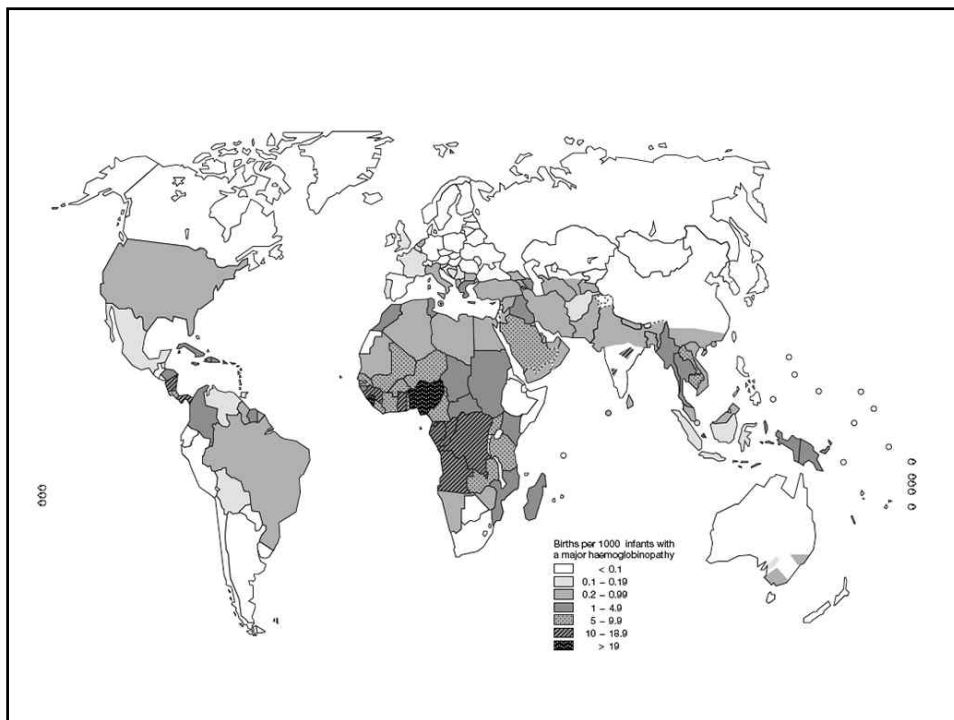
## CLASSIFICAZIONE IN BASE AL MECCANISMO MOLECOLARE RESPONSABILE (MUTAZIONI A LIVELLO DI ESONI)

- 1) MUTAZIONE PUNTIFORME: SINONIMA UAA->CAA LEU  
MISSENSE GUG->GAG GLU -> VAL HB S  
NON-SENSE UAU->UAA TYR-> STOP  
HB MCKEES ROCK
- 2) MUTAZIONE PUNTIFORME DOPPIA:  
2 EVENTI GENETICI  
BETA 6 GLU->LYS  
BETA 95 LYS->GLU  
HB ARLINGTON PARK
- 3) DELEZIONE DI UNA O PIU' TRIPLETTE:  
BETA 6 GLU->O HB LEYDEN
- 4) MUTAZIONI FRAMESHIFT:  
HB TAKOMA +11 AA
- 5) CROSSING OVER INEGUALE. NON OMOLOGO:  
HB LEPORE  
HB ANTI-LEPORE



**Tabella 24** *Classificazione funzionale delle emoglobinopatie*

	Anormalità Funzionale	Luogo Sostituzione	Sintomatologia	Esempio
1	Nessuna	Superficie esterna	Nessuna	Hb G Philadelphia $\alpha 68 \text{ Asn} \rightarrow \text{Lys}$
2	Aggregazione ridotta sol.	Superficie esterna	Anemia emolitica (omozigote)	Hb S $\beta 6 \text{ Glu} \rightarrow \text{Val}$
3	Instabilità ridotta sol.	Eme + contatti subunità	Anemia emolitica	Hb Bristol $\beta 67 \text{ Val} \rightarrow \text{Asp}$
4	MetHb	His prossimale e distale	Cianosi	Hb M Hyde Park $\beta 92 \text{ His} \rightarrow \text{Tyr}$
5	Aumento affinità per l'O <sub>2</sub>	Contatto $\alpha_1\beta_2$ , C-terminale $\beta$	Eritrocitosi	Hb Chesapeake $\alpha 92 \text{ Arg} \rightarrow \text{Leu}$ Hb Hiroshima $\beta 146 \text{ His} \rightarrow \text{Asp}$
6	Diminuita affinità per l'O <sub>2</sub>	Eme oppure $\alpha_1\beta_2$	Cianosi	Hb Kansas $\beta 102 \text{ Asn} \rightarrow \text{Thr}$
7	Ridotta produzione di catene $\alpha$	Mutazione a livello del codone C-terminale	Fenotipo $\alpha$ talassemico	Hb Constant Spring ( $\alpha+31$ )
8	Ridotta produzione di catene $\beta$	Ibrido di fusione	Fenotipo $\beta$ talassemico	Hb Lepore







Carrier Frequencies for Common Hemoglobin Disorders, by World Health Organization Region, 2001  
(percent)

Region	Hb S	Hb C	Hb E	$\beta$ thalassemia	$\alpha^0$ thalassemia	$\alpha^+$ thalassemia
Americas	1-20	0-10	0-20	0-3	0-5	0-40
Eastern Mediterranean	0-60	0-3	0-2	2-18	0-2	1-60
Europe	0-30	0-5	0-20	0-19	1-2	0-12
Southeast Asia	0-40	0	0-70	0-11	1-30	3-40
Sub-Saharan Africa	1-38	0-21	0	0-12	0	10-50
Western Pacific	0	0	0	0-13	0	2-60

Sources: Livingstone 1985; Weatherall and Clegg 2001a, 2001b.  
Note: Many of these data are derived from small population samples.



**Chapter 34**  
**Inherited Disorders of Hemoglobin**  
 David Weatherall, Theodor M. Ruzicka, Lutz Böttcher, Nancy D. Young, and Peter M. Higgs

**Introduction**

As a result of certain protein sequence variants, the inherited hemoglobin disorders are the commonest forms of genetic disease. They are caused by mutations in the genes for the two polypeptide chains of the hemoglobin molecule, the  $\alpha$ -globin and  $\beta$ -globin genes. The  $\alpha$ -globin gene is located on chromosome 16 and the  $\beta$ -globin gene on chromosome 11. Although these disorders are most frequent in tropical regions, they are also present in some temperate zones of the world.

**Structural Variants of Hemoglobin**

Structural variants of hemoglobin are caused by mutations in the genes for the  $\alpha$ -globin and  $\beta$ -globin chains. These variants are usually benign and do not cause disease. They are found in all ethnic groups and are most frequent in tropical regions. The most common structural variant is Hb E, which is found in about 10% of the population in Southeast Asia. Other structural variants include Hb C, Hb D, Hb G, Hb I, Hb K, Hb L, Hb M, Hb N, Hb O, Hb P, Hb Q, Hb R, Hb S, Hb T, Hb U, Hb V, Hb W, Hb X, Hb Y, Hb Z, Hb AA, Hb AB, Hb AC, Hb AD, Hb AE, Hb AF, Hb AG, Hb AH, Hb AI, Hb AJ, Hb AK, Hb AL, Hb AM, Hb AN, Hb AO, Hb AP, Hb AQ, Hb AR, Hb AS, Hb AT, Hb AU, Hb AV, Hb AW, Hb AX, Hb AY, Hb AZ, Hb BA, Hb BB, Hb BC, Hb BD, Hb BE, Hb BF, Hb BG, Hb BH, Hb BI, Hb BJ, Hb BK, Hb BL, Hb BM, Hb BN, Hb BO, Hb BP, Hb BQ, Hb BR, Hb BS, Hb BT, Hb BU, Hb BV, Hb BW, Hb BX, Hb BY, Hb BZ, Hb CA, Hb CB, Hb CC, Hb CD, Hb CE, Hb CF, Hb CG, Hb CH, Hb CI, Hb CJ, Hb CK, Hb CL, Hb CM, Hb CN, Hb CO, Hb CP, Hb CQ, Hb CR, Hb CS, Hb CT, Hb CU, Hb CV, Hb CW, Hb CX, Hb CY, Hb CZ, Hb DA, Hb DB, Hb DC, Hb DD, Hb DE, Hb DF, Hb DG, Hb DH, Hb DI, Hb DJ, Hb DK, Hb DL, Hb DM, Hb DN, Hb DO, Hb DP, Hb DQ, Hb DR, Hb DS, Hb DT, Hb DU, Hb DV, Hb DW, Hb DX, Hb DY, Hb DZ, Hb EA, Hb EB, Hb EC, Hb ED, Hb EE, Hb EF, Hb EG, Hb EH, Hb EI, Hb EJ, Hb EK, Hb EL, Hb EM, Hb EN, Hb EO, Hb EP, Hb EQ, Hb ER, Hb ES, Hb ET, Hb EU, Hb EV, Hb EW, Hb EX, Hb EY, Hb EZ, Hb FA, Hb FB, Hb FC, Hb FD, Hb FE, Hb FF, Hb FG, Hb FH, Hb FI, Hb FJ, Hb FK, Hb FL, Hb FM, Hb FN, Hb FO, Hb FP, Hb FQ, Hb FR, Hb FS, Hb FT, Hb FU, Hb FV, Hb FW, Hb FX, Hb FY, Hb FZ, Hb GA, Hb GB, Hb GC, Hb GD, Hb GE, Hb GF, Hb GG, Hb GH, Hb GI, Hb GJ, Hb GK, Hb GL, Hb GM, Hb GN, Hb GO, Hb GP, Hb GQ, Hb GR, Hb GS, Hb GT, Hb GU, Hb GV, Hb GW, Hb GX, Hb GY, Hb GZ, Hb HA, Hb HB, Hb HC, Hb HD, Hb HE, Hb HF, Hb HG, Hb HH, Hb HI, Hb HJ, Hb HK, Hb HL, Hb HM, Hb HN, Hb HO, Hb HP, Hb HQ, Hb HR, Hb HS, Hb HT, Hb HU, Hb HV, Hb HW, Hb HX, Hb HY, Hb HZ, Hb IA, Hb IB, Hb IC, Hb ID, Hb IE, Hb IF, Hb IG, Hb IH, Hb II, Hb IJ, Hb IK, Hb IL, Hb IM, Hb IN, Hb IO, Hb IP, Hb IQ, Hb IR, Hb IS, Hb IT, Hb IU, Hb IV, Hb IW, Hb IX, Hb IY, Hb IZ, Hb JA, Hb JB, Hb JC, Hb JD, Hb JE, Hb JF, Hb JG, Hb JH, Hb JI, Hb JJ, Hb JK, Hb JL, Hb JM, Hb JN, Hb JO, Hb JP, Hb JQ, Hb JR, Hb JS, Hb JT, Hb JU, Hb JV, Hb JW, Hb JX, Hb JY, Hb JZ, Hb KA, Hb KB, Hb KC, Hb KD, Hb KE, Hb KF, Hb KG, Hb KH, Hb KI, Hb KJ, Hb KK, Hb KL, Hb KM, Hb KN, Hb KO, Hb KP, Hb KQ, Hb KR, Hb KS, Hb KT, Hb KU, Hb KV, Hb KW, Hb KX, Hb KY, Hb KZ, Hb LA, Hb LB, Hb LC, Hb LD, Hb LE, Hb LF, Hb LG, Hb LH, Hb LI, Hb LJ, Hb LK, Hb LL, Hb LM, Hb LN, Hb LO, Hb LP, Hb LQ, Hb LR, Hb LS, Hb LT, Hb LU, Hb LV, Hb LW, Hb LX, Hb LY, Hb LZ, Hb MA, Hb MB, Hb MC, Hb MD, Hb ME, Hb MF, Hb MG, Hb MH, Hb MI, Hb MJ, Hb MK, Hb ML, Hb MM, Hb MN, Hb MO, Hb MP, Hb MQ, Hb MR, Hb MS, Hb MT, Hb MU, Hb MV, Hb MW, Hb MX, Hb MY, Hb MZ, Hb NA, Hb NB, Hb NC, Hb ND, Hb NE, Hb NF, Hb NG, Hb NH, Hb NI, Hb NJ, Hb NK, Hb NL, Hb NM, Hb NN, Hb NO, Hb NP, Hb NQ, Hb NR, Hb NS, Hb NT, Hb NU, Hb NV, Hb NW, Hb NX, Hb NY, Hb NZ, Hb OA, Hb OB, Hb OC, Hb OD, Hb OE, Hb OF, Hb OG, Hb OH, Hb OI, Hb OJ, Hb OK, Hb OL, Hb OM, Hb ON, Hb OO, Hb OP, Hb OQ, Hb OR, Hb OS, Hb OT, Hb OU, Hb OV, Hb OW, Hb OX, Hb OY, Hb OZ, Hb PA, Hb PB, Hb PC, Hb PD, Hb PE, Hb PF, Hb PG, Hb PH, Hb PI, Hb PJ, Hb PK, Hb PL, Hb PM, Hb PN, Hb PO, Hb PP, Hb PQ, Hb PR, Hb PS, Hb PT, Hb PU, Hb PV, Hb PW, Hb PX, Hb PY, Hb PZ, Hb QA, Hb QB, Hb QC, Hb QD, Hb QE, Hb QF, Hb QG, Hb QH, Hb QI, Hb QJ, Hb QK, Hb QL, Hb QM, Hb QN, Hb QO, Hb QP, Hb QQ, Hb QR, Hb QS, Hb QT, Hb QU, Hb QV, Hb QW, Hb QX, Hb QY, Hb QZ, Hb RA, Hb RB, Hb RC, Hb RD, Hb RE, Hb RF, Hb RG, Hb RH, Hb RI, Hb RJ, Hb RK, Hb RL, Hb RM, Hb RN, Hb RO, Hb RP, Hb RQ, Hb RR, Hb RS, Hb RT, Hb RU, Hb RV, Hb RW, Hb RX, Hb RY, Hb RZ, Hb SA, Hb SB, Hb SC, Hb SD, Hb SE, Hb SF, Hb SG, Hb SH, Hb SI, Hb SJ, Hb SK, Hb SL, Hb SM, Hb SN, Hb SO, Hb SP, Hb SQ, Hb SR, Hb SS, Hb ST, Hb SU, Hb SV, Hb SW, Hb SX, Hb SY, Hb SZ, Hb TA, Hb TB, Hb TC, Hb TD, Hb TE, Hb TF, Hb TG, Hb TH, Hb TI, Hb TJ, Hb TK, Hb TL, Hb TM, Hb TN, Hb TO, Hb TP, Hb TQ, Hb TR, Hb TS, Hb TT, Hb TU, Hb TV, Hb TW, Hb TX, Hb TY, Hb TZ, Hb UA, Hb UB, Hb UC, Hb UD, Hb UE, Hb UF, Hb UG, Hb UH, Hb UI, Hb UJ, Hb UK, Hb UL, Hb UM, Hb UN, Hb UO, Hb UP, Hb UQ, Hb UR, Hb US, Hb UT, Hb UU, Hb UV, Hb UW, Hb UX, Hb UY, Hb UZ, Hb VA, Hb VB, Hb VC, Hb VD, Hb VE, Hb VF, Hb VG, Hb VH, Hb VI, Hb VJ, Hb VK, Hb VL, Hb VM, Hb VN, Hb VO, Hb VP, Hb VQ, Hb VR, Hb VS, Hb VT, Hb VU, Hb VV, Hb VW, Hb VX, Hb VY, Hb VZ, Hb WA, Hb WB, Hb WC, Hb WD, Hb WE, Hb WF, Hb WG, Hb WH, Hb WI, Hb WJ, Hb WK, Hb WL, Hb WM, Hb WN, Hb WO, Hb WP, Hb WQ, Hb WR, Hb WS, Hb WT, Hb WU, Hb WV, Hb WW, Hb WX, Hb WY, Hb WZ, Hb XA, Hb XB, Hb XC, Hb XD, Hb XE, Hb XF, Hb XG, Hb XH, Hb XI, Hb XJ, Hb XK, Hb XL, Hb XM, Hb XN, Hb XO, Hb XP, Hb XQ, Hb XR, Hb XS, Hb XT, Hb XU, Hb XV, Hb XW, Hb XX, Hb XY, Hb XZ, Hb YA, Hb YB, Hb YC, Hb YD, Hb YE, Hb YF, Hb YG, Hb YH, Hb YI, Hb YJ, Hb YK, Hb YL, Hb YM, Hb YN, Hb YO, Hb YP, Hb YQ, Hb YR, Hb YS, Hb YT, Hb YU, Hb YV, Hb YW, Hb YX, Hb YY, Hb YZ, Hb ZA, Hb ZB, Hb ZC, Hb ZD, Hb ZE, Hb ZF, Hb ZG, Hb ZH, Hb ZI, Hb ZJ, Hb ZK, Hb ZL, Hb ZM, Hb ZN, Hb ZO, Hb ZP, Hb ZQ, Hb ZR, Hb ZS, Hb ZT, Hb ZU, Hb ZV, Hb ZW, Hb ZX, Hb ZY, Hb ZZ.



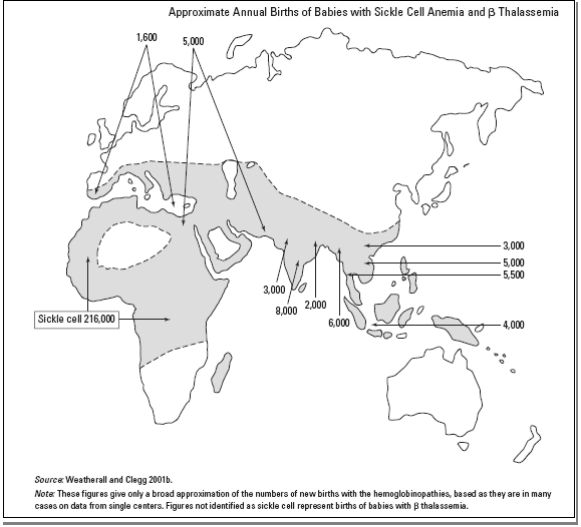
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**Structural Variants of Hemoglobin**

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Other structural variants include Hb C, Hb D, Hb G, Hb I, Hb K, Hb L, Hb M, Hb N, Hb O, Hb P, Hb Q, Hb R, Hb S, Hb T, Hb U, Hb V, Hb W, Hb X, Hb Y, Hb Z, Hb AA, Hb AB, Hb AC, Hb AD, Hb AE, Hb AF, Hb AG, Hb AH, Hb AI, Hb AJ, Hb AK, Hb AL, Hb AM, Hb AN, Hb AO, Hb AP, Hb AQ, Hb AR, Hb AS, Hb AT, Hb AU, Hb AV, Hb AW, Hb AX, Hb AY, Hb AZ, Hb BA, Hb BB, Hb BC, Hb BD, Hb BE, Hb BF, Hb BG, Hb BH, Hb BI, Hb BJ, Hb BK, Hb BL, Hb BM, Hb BN, Hb BO, Hb BP, Hb BQ, Hb BR, Hb BS, Hb BT, Hb BU, Hb BV, Hb BW, Hb BX, Hb BY, Hb BZ, Hb CA, Hb CB, Hb CC, Hb CD, Hb CE, Hb CF, Hb CG, Hb CH, Hb CI, Hb CJ, Hb CK, Hb CL, Hb CM, Hb CN, Hb CO, Hb CP, Hb CQ, Hb CR, Hb CS, Hb CT, Hb CU, Hb CV, Hb CW, Hb CX, Hb CY, Hb CZ, Hb DA, Hb DB, Hb DC, Hb DD, Hb DE, Hb DF, Hb DG, Hb DH, Hb DI, Hb DJ, Hb DK, Hb DL, Hb DM, Hb DN, Hb DO, Hb DP, Hb DQ, Hb DR, Hb DS, Hb DT, Hb DU, Hb DV, Hb DW, Hb DX, Hb DY, Hb DZ, Hb EA, Hb EB, Hb EC, Hb ED, Hb EE, Hb EF, Hb EG, Hb EH, Hb EI, Hb EJ, Hb EK, Hb EL, Hb EM, Hb EN, Hb EO, Hb EP, Hb EQ, Hb ER, Hb ES, Hb ET, Hb EU, Hb EV, Hb EW, Hb EX, Hb EY, Hb EZ, Hb FA, Hb FB, Hb FC, Hb FD, Hb FE, Hb FF, Hb FG, Hb FH, Hb FI, Hb FJ, Hb FK, Hb FL, Hb FM, Hb FN, Hb FO, Hb FP, Hb FQ, Hb FR, Hb FS, Hb FT, Hb FU, Hb FV, Hb FW, Hb FX, Hb FY, Hb FZ, Hb GA, Hb GB, Hb GC, Hb GD, Hb GE, Hb GF, Hb GG, Hb GH, Hb GI, Hb GJ, Hb GK, Hb GL, Hb GM, Hb GN, Hb GO, Hb GP, Hb GQ, Hb GR, Hb GS, Hb GT, Hb GU, Hb GV, Hb GW, Hb GX, Hb GY, Hb GZ, Hb HA, Hb HB, Hb HC, Hb HD, Hb HE, Hb HF, Hb HG, Hb HH, Hb HI, Hb HJ, Hb HK, Hb HL, Hb HM, Hb HN, Hb HO, Hb HP, Hb HQ, Hb HR, Hb HS, Hb HT, Hb HU, Hb HV, Hb HW, Hb HX, Hb HY, Hb HZ, Hb IA, Hb IB, Hb IC, Hb ID, Hb IE, Hb IF, Hb IG, Hb IH, Hb II, Hb IJ, Hb IK, Hb IL, Hb IM, Hb IN, Hb IO, Hb IP, Hb IQ, Hb IR, Hb IS, Hb IT, Hb IU, Hb IV, Hb IW, Hb IX, Hb IY, Hb IZ, Hb JA, Hb JB, Hb JC, Hb JD, Hb JE, Hb JF, Hb JG, Hb JH, Hb JI, Hb JJ, Hb JK, Hb JL, Hb JM, Hb JN, Hb JO, Hb JP, Hb JQ, Hb JR, Hb JS, Hb JT, Hb JU, Hb JV, Hb JW, Hb JX, Hb JY, Hb JZ, Hb KA, Hb KB, Hb KC, Hb KD, Hb KE, Hb KF, Hb KG, Hb KH, Hb KI, Hb KJ, Hb KK, Hb KL, Hb KM, Hb KN, Hb KO, Hb KP, Hb KQ, Hb KR, Hb KS, Hb KT, Hb KU, Hb KV, Hb KW, Hb KX, Hb KY, Hb KZ, Hb LA, Hb LB, Hb LC, Hb LD, Hb LE, Hb LF, Hb LG, Hb LH, Hb LI, Hb LJ, Hb LK, Hb LL, Hb LM, Hb LN, Hb LO, Hb LP, Hb LQ, Hb LR, Hb LS, Hb LT, Hb LU, Hb LV, Hb LW, Hb LX, Hb LY, Hb LZ, Hb MA, Hb MB, Hb MC, Hb MD, Hb ME, Hb MF, Hb MG, Hb MH, Hb MI, Hb MJ, Hb MK, Hb ML, Hb MM, Hb MN, Hb MO, Hb MP, Hb MQ, Hb MR, Hb MS, Hb MT, Hb MU, Hb MV, Hb MW, Hb MX, Hb MY, Hb MZ, Hb NA, Hb NB, Hb NC, Hb ND, Hb NE, Hb NF, Hb NG, Hb NH, Hb NI, Hb NJ, Hb NK, Hb NL, Hb NM, Hb NN, Hb NO, Hb NP, Hb NQ, Hb NR, Hb NS, Hb NT, Hb NU, Hb NV, Hb NW, Hb NX, Hb NY, Hb NZ, Hb OA, Hb OB, Hb OC, Hb OD, Hb OE, Hb OF, Hb OG, Hb OH, Hb OI, Hb OJ, Hb OK, Hb OL, Hb OM, Hb ON, Hb OO, Hb OP, Hb OQ, Hb OR, Hb OS, Hb OT, Hb OU, Hb OV, Hb OW, Hb OX, Hb OY, Hb OZ, Hb PA, Hb PB, Hb PC, Hb PD, Hb PE, Hb PF, Hb PG, Hb PH, Hb PI, Hb PJ, Hb PK, Hb PL, Hb PM, Hb PN, Hb PO, Hb PP, Hb PQ, Hb PR, Hb PS, Hb PT, Hb PU, Hb PV, Hb PW, Hb PX, Hb PY, Hb PZ, Hb QA, Hb QB, Hb QC, Hb QD, Hb QE, Hb QF, Hb QG, Hb QH, Hb QI, Hb QJ, Hb QK, Hb QL, Hb QM, Hb QN, Hb QO, Hb QP, Hb QQ, Hb QR, Hb QS, Hb QT, Hb QU, Hb QV, Hb QW, Hb QX, Hb QY, Hb QZ, Hb RA, Hb RB, Hb RC, Hb RD, Hb RE, Hb RF, Hb RG, Hb RH, Hb RI, Hb RJ, Hb RK, Hb RL, Hb RM, Hb RN, Hb RO, Hb RP, Hb RQ, Hb RR, Hb RS, Hb RT, Hb RU, Hb RV, Hb RW, Hb RX, Hb RY, Hb RZ, Hb SA, Hb SB, Hb SC, Hb SD, Hb SE, Hb SF, Hb SG, Hb SH, Hb SI, Hb SJ, Hb SK, Hb SL, Hb SM, Hb SN, Hb SO, Hb SP, Hb SQ, Hb SR, Hb SS, Hb ST, Hb SU, Hb SV, Hb SW, Hb SX, Hb SY, Hb SZ, Hb TA, Hb TB, Hb TC, Hb TD, Hb TE, Hb TF, Hb TG, Hb TH, Hb TI, Hb TJ, Hb TK, Hb TL, Hb TM, Hb TN, Hb TO, Hb TP, Hb TQ, Hb TR, Hb TS, Hb TT, Hb TU, Hb TV, Hb TW, Hb TX, Hb TY, Hb TZ, Hb UA, Hb UB, Hb UC, Hb UD, Hb UE, Hb UF, Hb UG, Hb UH, Hb UI, Hb UJ, Hb UK, Hb UL, Hb UM, Hb UN, Hb UO, Hb UP, Hb UQ, Hb UR, Hb US, Hb UT, Hb UU, Hb UV, Hb UW, Hb UX, Hb UY, Hb UZ, Hb VA, Hb VB, Hb VC, Hb VD, Hb VE, Hb VF, Hb VG, Hb VH, Hb VI, Hb VJ, Hb VK, Hb VL, Hb VM, Hb VN, Hb VO, Hb VP, Hb VQ, Hb VR, Hb VS, Hb VT, Hb VU, Hb VV, Hb VW, Hb VX, Hb VY, Hb VZ, Hb WA, Hb WB, Hb WC, Hb WD, Hb WE, Hb WF, Hb WG, Hb WH, Hb WI, Hb WJ, Hb WK, Hb WL, Hb WM, Hb WN, Hb WO, Hb WP, Hb WQ, Hb WR, Hb WS, Hb WT, Hb WU, Hb WV, Hb WW, Hb WX, Hb WY, Hb WZ, Hb XA, Hb XB, Hb XC, Hb XD, Hb XE, Hb XF, Hb XG, Hb XH, Hb XI, Hb XJ, Hb XK, Hb XL, Hb XM, Hb XN, Hb XO, Hb XP, Hb XQ, Hb XR, Hb XS, Hb XT, Hb XU, Hb XV, Hb XW, Hb XY, Hb XZ, Hb YA, Hb YB, Hb YC, Hb YD, Hb YE, Hb YF, Hb YG, Hb YH, Hb YI, Hb YJ, Hb YK, Hb YL, Hb YM, Hb YN, Hb YO, Hb YP, Hb YQ, Hb YR, Hb YS, Hb YT, Hb YU, Hb YV, Hb YW, Hb YX, Hb YY, Hb YZ, Hb ZA, Hb ZB, Hb ZC, Hb ZD, Hb ZE, Hb ZF, Hb ZG, Hb ZH, Hb ZI, Hb ZJ, Hb ZK, Hb ZL, Hb ZM, Hb ZN, Hb ZO, Hb ZP, Hb ZQ, Hb ZR, Hb ZS, Hb ZT, Hb ZU, Hb ZV, Hb ZW, Hb ZX, Hb ZY, Hb ZZ.





Burden of Sickle Cell Disease by Age Group, Assuming 1,000 Births per Year and Survival to Various Ages,

Category	Age group (years)										Total or average
	0-4	5-9	10-14	15-19	20-24	25-29	30-34	35-39	40-44	45-49	
Number of survivors	876	834	807	777	727	680	627	564	491	440	682.3
Number of deaths	124	42	27	30	50	47	53	63	73	51	560
Death rate (percent/year)	2.61	0.98	0.66	0.75	1.32	1.33	1.61	2.10	2.73	2.17	1.63
Number of DALYs lost/death	28.90	28.59	27.77	26.84	25.82	24.69	23.43	22.00	20.39	18.58	24.70
Total DALY losses from deaths	3,584	1,201	750	805	1,291	1,161	1,242	1,366	1,468	948	13,856
Number of DALYs lost from background (chronic) anemia	188	171	164	158	150	141	130	119	106	93	1,420
Total DALYs lost from deaths and chronic anemia	3,772	1,372	914	963	1,441	1,302	1,372	1,505	1,594	1,041	15,276
Number of pain crises/year	242.7	381.0	383.8	584.4	866.7	600.5	523.6	473.4	309.6	182.2	4,548
Number of other acute clinical events	77.5	22.2		182.2							281.9
Number of other chronic clinical events	49.8	14.8	12.8	10.9							88.3

Source: Authors' calculations based on Hambleton 2004a, 2004b.



Annual Costs of Hemoglobinopathy per Outpatient, Excluding Transfusion, Toronto (2001 US\$)

Category	Thalassemia		Sickle cell disease	
	Chelated	Nonchelated	Chelated	Nonchelated
Clinic staff salaries	1,011.95	183.68	1,011.95	252.99
Clinic supplies	930.19	25.15	930.19	34.85
Medical and surgical outpatient unit	2,069.57	n.a.	2,069.57	n.a.
Consultations	92.58	88.39	92.58	11.94
Diagnostic tests	742.58	281.44	905.89	210.99
Laboratory costs	413.96	31.04	414.01	42.74
Laboratory costs (medical dayunit visits)	665.81	n.a.	665.81	n.a.
Total	5,926.64	608.70	6,090.00	553.31

Source: Estimated costs provided by Nancy Oliveri of the University of Toronto.  
Note: n.a. = not applicable.

Costs of Treatment of Thalassemia for One Patient Age 7 to 11, Eastern Mediterranean (2001 US\$)

Category	Minimum treatment	Full treatment
<i>Costs other than iron chelation</i>		
Day transfusion: hotel and nursing	375	375
12 transfusions/year	1,088	2,250
Investigations	135	278
Occasional costs (such as operations)	150	645
Staff salaries	300	620
Total if no desferrioxamine therapy	2,048	n.a.
Desferrioxamine therapy (iron chelation)	3,080	6,165
Total with desferrioxamine therapy	5,128	10,333

Source: Ahean and Modell 1997.



Cost-Effectiveness of Treatment for Homozygous  $\beta$  and Transfusion-Dependent Hb E  $\beta$  Thalassemia

Category	Cost/patient (US\$)	DALYs gained/patient		Cost/DALY (US\$)	
		Disability weight = 0.1	Disability weight = 0.25	Disability weight = 0.1	Disability weight = 0.25
<i>Minimal treatment, transfusion only</i>					
Until death at age 10	17,368	6.96–7.60	6.00–6.39	2,285–2,495	2,718–2,896
Until death at age 15	23,840	10.25–10.81	7.52–7.87	2,206–2,325	3,029–3,170
<i>Full treatment with chelation: incremental compared with minimal treatment</i>					
Until age 15	60,467	1.03–3.80	0.55–2.03	15,912–58,706	29,787–109,940
Beyond age 15 to maximum age 80	132,901	17.25	16.35	7,704	8,129
Total lifetime	193,368	18.28–21.05	16.90–18.38	9,186–10,578	10,520–11,442
<i>Full treatment with chelation: total compared with no treatment</i>					
Until age 15	121,284	11.81	10.84	10,273	11,186
Beyond age 15 to maximum age 80	274,662	17.25	16.35	15,922	16,799
Total lifetime	395,946	29.06	27.19	13,625	14,578

Source: Authors' calculations. All costs and DALYs gained are discounted at 3 percent annually, starting at birth.

Public health reviews

Global epidemiology of haemoglobin disorders and derived service indicators

Barbara Machi S. Mouton-Carlier\*

**Abstract** This review provides an overview of the global epidemiology of haemoglobin disorders, including the prevalence of genetic disorders, the burden and prevention of haemoglobin disorders, the clinical features, natural course, and public health and genetic counselling implications for haemoglobin disorders by country, sex, and ethnicity, including the genetic carrier status in relation to the need for genetic counselling and reproductive decisions.

Introduction

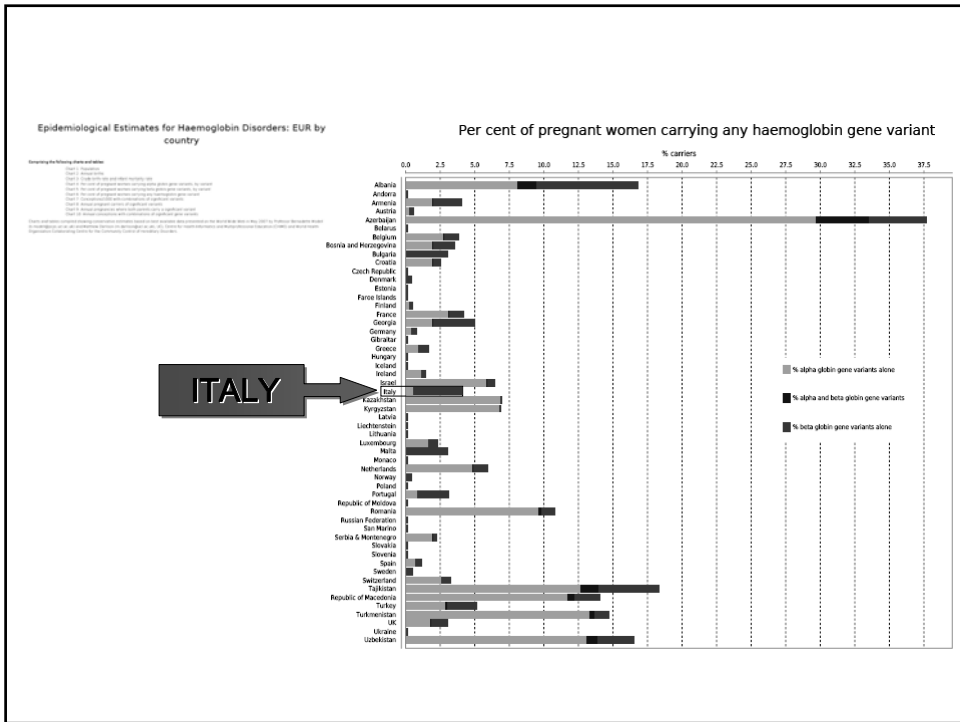
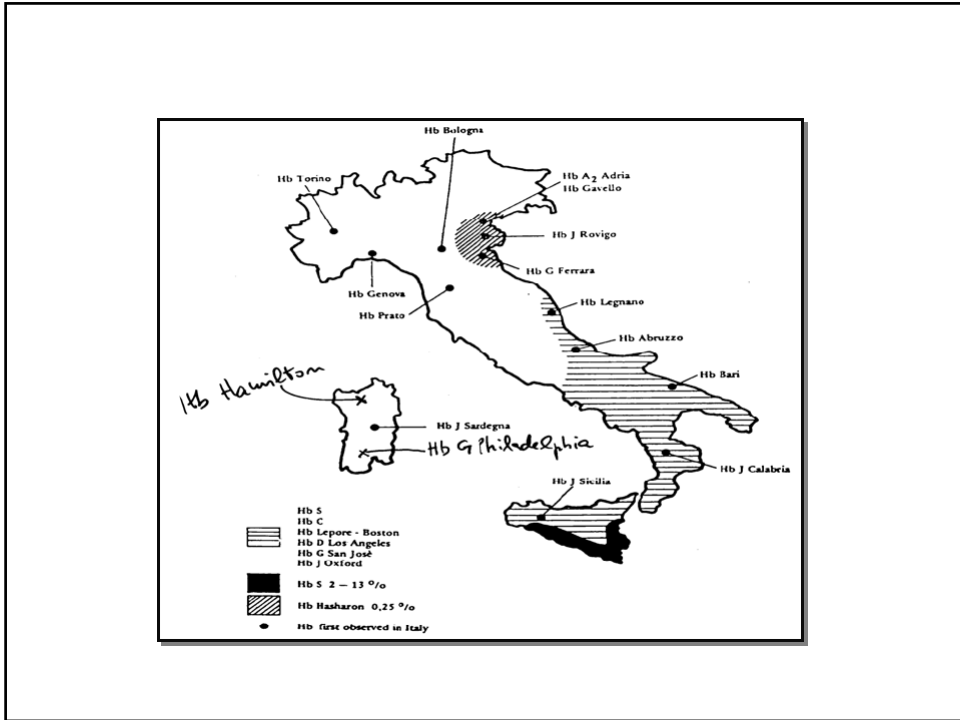
Haemoglobin disorders are inherited genetic conditions that affect the ability of red blood cells to carry oxygen. The most common types are sickle cell disease and thalassemia. These disorders are caused by mutations in the genes that code for the hemoglobin protein. The prevalence of these disorders varies significantly across different populations and ethnic groups. In this review, we will discuss the global epidemiology of haemoglobin disorders, including the prevalence of genetic disorders, the burden and prevention of haemoglobin disorders, the clinical features, natural course, and public health and genetic counselling implications for haemoglobin disorders by country, sex, and ethnicity, including the genetic carrier status in relation to the need for genetic counselling and reproductive decisions.

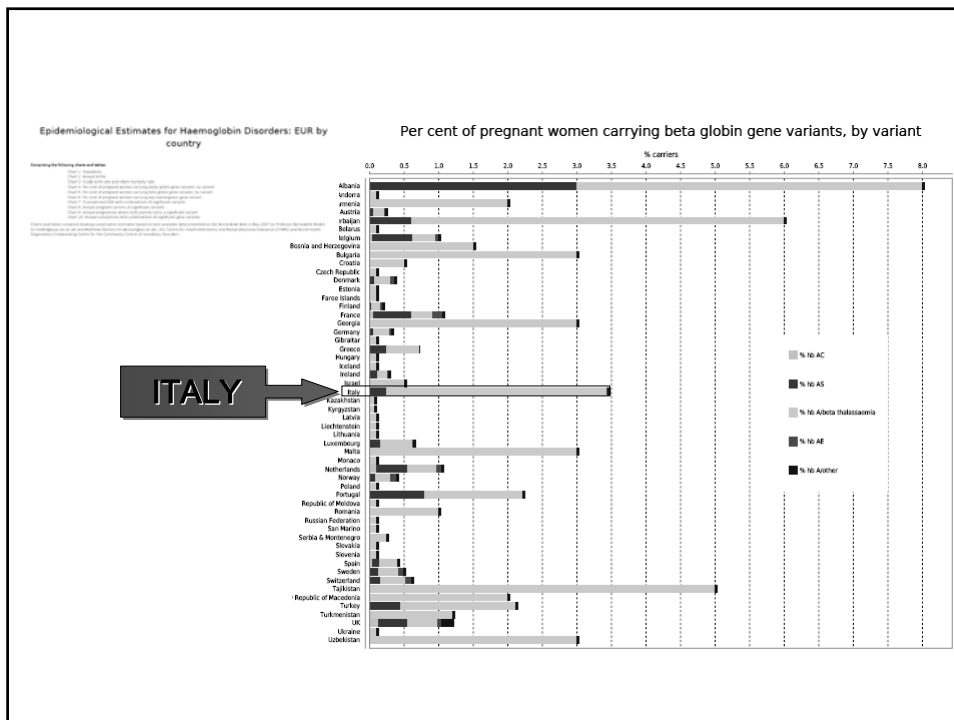
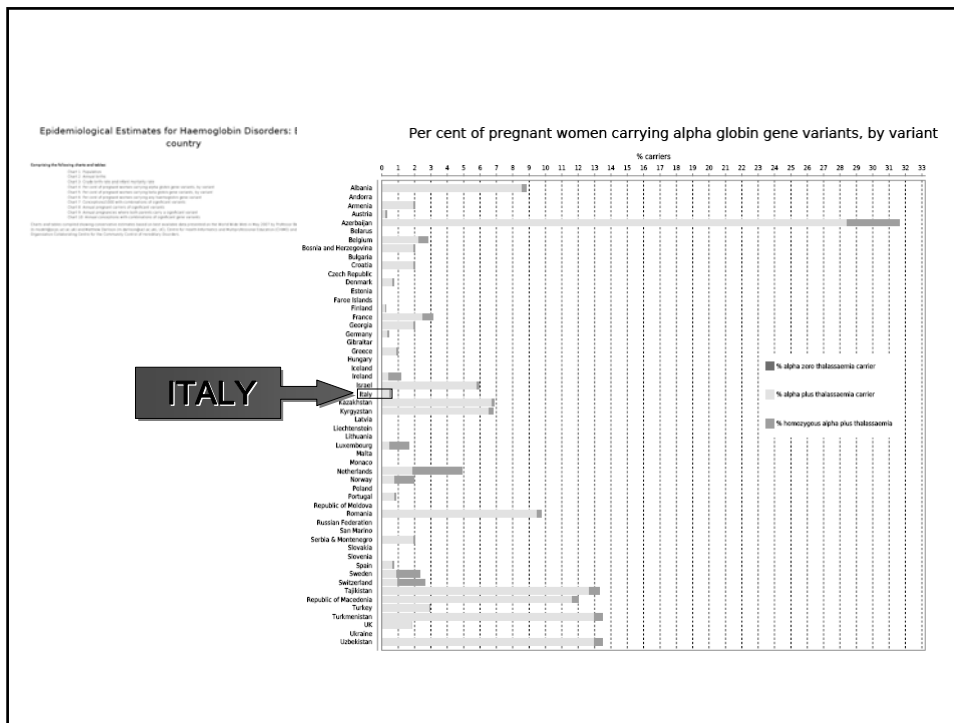
Table 1. Estimated prevalences of carriers of haemoglobin gene variants and affected conceptions

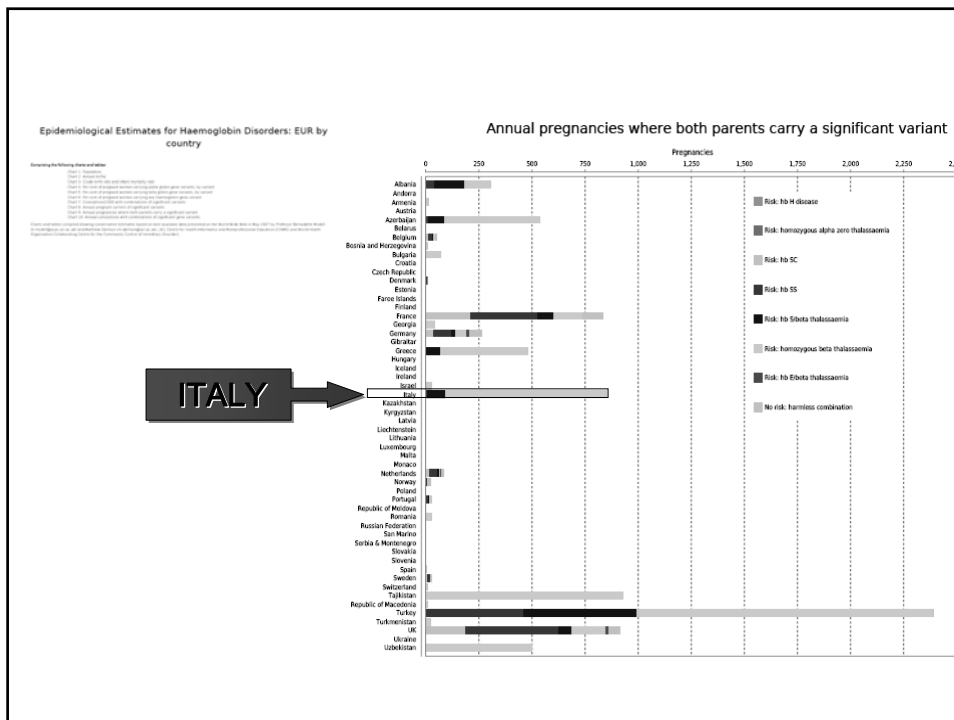
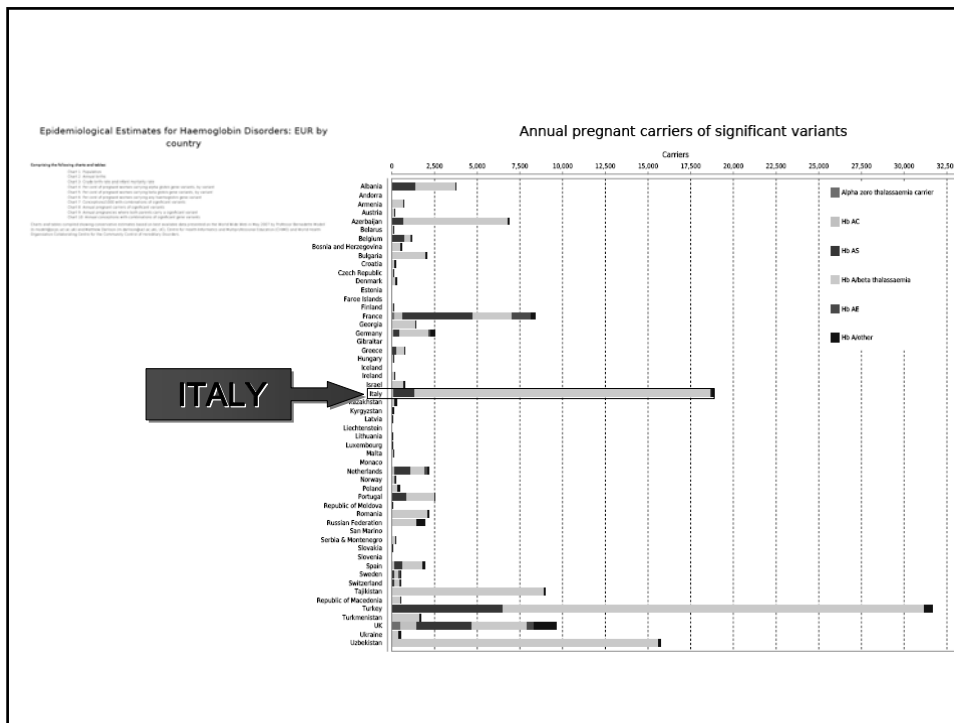
WHO region	Demography 2003			% of the population carrying			Affected conceptions (per 1000)		Affected births (% of under-5 mortality)		
	Population (millions)	Crude birth rate	Annual births (1000s)	Under-5 mortality rate	Significant variant <sup>a</sup>	Any variant <sup>b</sup>	Sickle-cell disorders <sup>c</sup>	Thalassaemias <sup>d</sup>			
African	586	39.0	22 895	168	18.2	41.2	44.4	10.88	0.07	10.74	6.4
American	853	19.5	16 609	27	3.0	4.8	7.5	0.49	0.06	0.54	2.0
Eastern Mediterranean	573	29.3	16 798	108	4.4	19.0	21.7	0.84	0.70	1.54	1.4
European	879	11.9	10 459	25	1.1	2.3	3.3	0.07	0.13	0.20	0.8
South-east Asian	1 564	24.4	38 139	83	6.6	44.6	45.5	0.68	0.66	1.34	1.6
Western Pacific	1 761	13.6	23 914	38	3.2	10.3	13.2	0.00	0.76	0.76	2.0
<b>World</b>	<b>6 217</b>	<b>20.7</b>	<b>128 814</b>	<b>81</b>	<b>5.2</b>	<b>20.7</b>	<b>24.0</b>	<b>2.28</b>	<b>0.46</b>	<b>2.73</b>	<b>3.4</b>

<sup>a</sup> Significant variants include Hb S, Hb C, Hb E, Hb D etc.  $\beta$  thalassaemia,  $\alpha^+$  thalassaemia.  
<sup>b</sup>  $\alpha^+$  thalassaemia includes heterozygous and homozygous  $\alpha^+$  thalassaemia.  
<sup>c</sup> Allows for (1) coexistence of  $\alpha$  and  $\beta$  variants, and (2) harmless combinations of  $\beta$  variants.  
<sup>d</sup> Sickle-cell disorders include SS, SC, S $\beta$  thalassaemia.  
<sup>e</sup> Thalassaemias include homozygous  $\beta$  thalassaemia, haemoglobin E/ $\beta$  thalassaemia, homozygous  $\alpha^+$  thalassaemia,  $\alpha^0/\alpha^+$  thalassaemia (haemoglobin H disease).

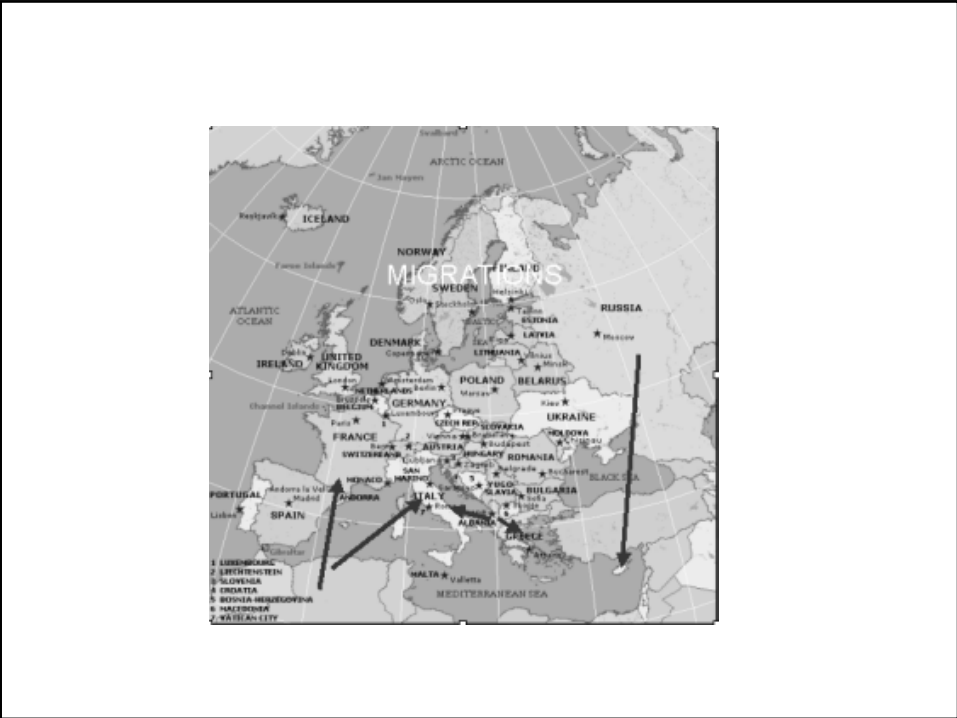






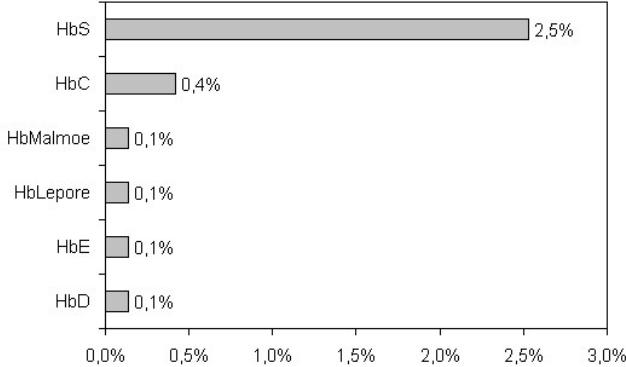






**Frequency and type of newly diagnosed hemoglobin variants in Northern Italy.**

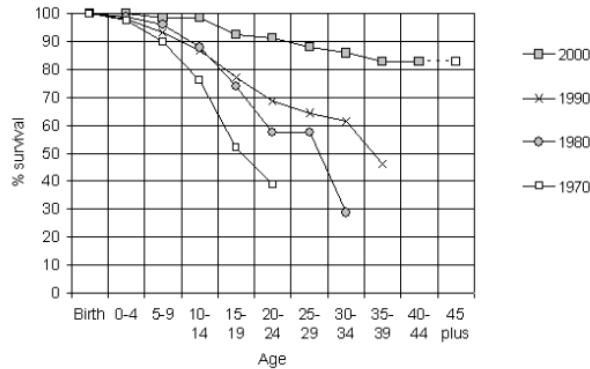
Giuseppe Lippi, Martina Montagnana, Elisa Danese, Gian Luca Salvagno, Francesca Bellorio, Gian Cesare Guidi.  
*Sezione di Chimica e Clinica, Dipartimento di Scienze Morfologico-Biomediche, Università degli Studi di Verona, Italy.*



Model patient register shows use for:

- Supporting equitable delivery of the best possible clinical care of patients with rare disorders
- Monitoring services - both treatment and prevention

Improving life-expectancy in thalassaemia in the UK.



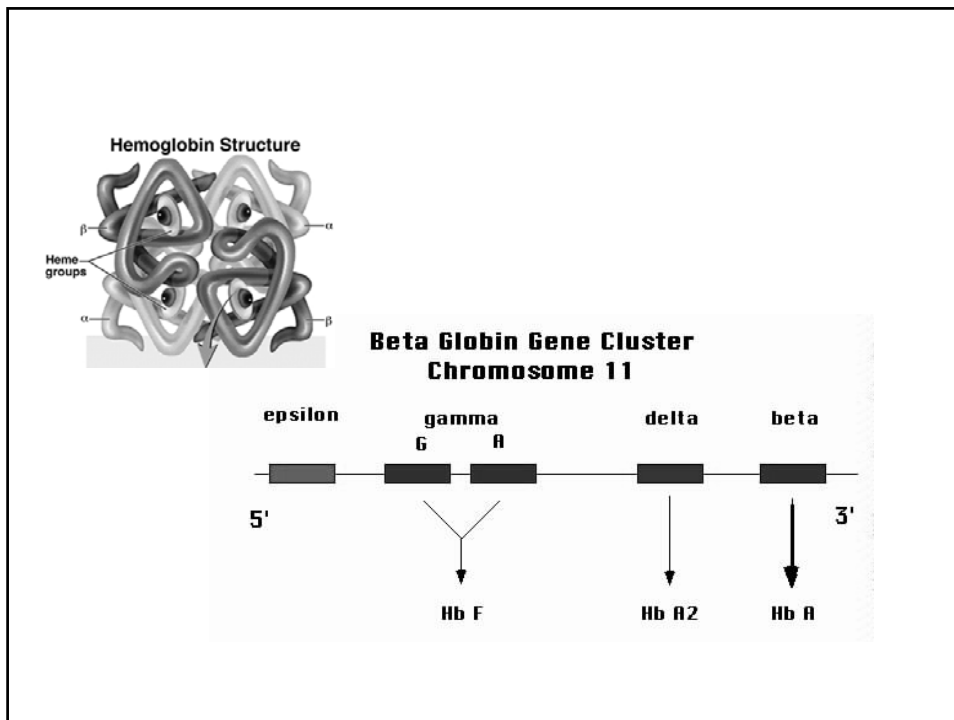
**Public health reviews**  
**Global epidemiology of haemoglobin disorders and derived service indicators**  
 Saraideh Masaki & Matthew Carter

**Abstract** This review is intended for policy-makers and public health professionals responsible for equitable clinical services, including the diagnosis and management of haemoglobin disorders. We present data on the epidemiology and prevalence of gene variants responsible for haemoglobin disorders from diverse populations, treatment outcomes, and published evidence of a clear relationship between haemoglobin disorders by country, including the impact of service indicators to improve the health of patients with haemoglobin disorders. We present data on the epidemiology and prevalence of gene variants responsible for haemoglobin disorders from diverse populations, treatment outcomes, and published evidence of a clear relationship between haemoglobin disorders by country, including the impact of service indicators to improve the health of patients with haemoglobin disorders.

**Introduction** Haemoglobin disorders include sickle cell disease (SCD) and thalassaemia. These are the most common genetic causes of anaemia worldwide. The prevalence of these disorders varies significantly between different ethnic groups and geographical regions. In the United Kingdom, the prevalence of SCD is highest in people of African and Caribbean descent, while thalassaemia is most common in people of Mediterranean, South Asian, and Chinese descent. The impact of these disorders on public health is significant, particularly in terms of the need for specialized clinical services and the potential for complications such as stroke, heart failure, and organ damage. This review examines the global epidemiology of these disorders and discusses the implications for public health practice, including the importance of genetic testing, counselling, and equitable access to clinical services.

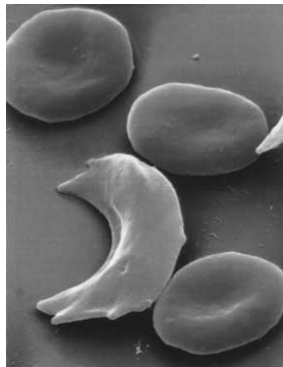
## Conclusion

The data summarized here confirm that screening and genetic counselling for haemoglobin disorders should be an intrinsic part of health care

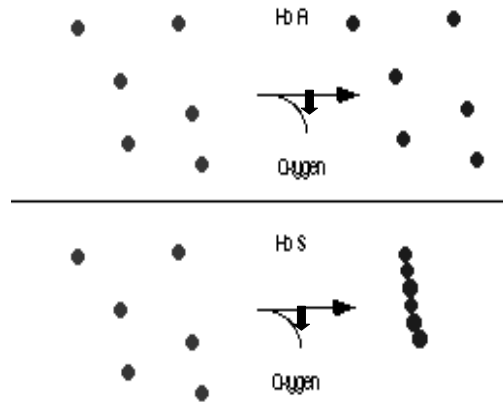


## Hemoglobin S: What's the big deal?

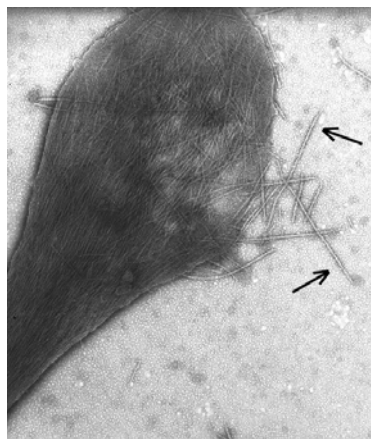
1927 Hahn and Gillespie showed sickling with deoxygenation of red cells in a patient



### Hemoglobin S polymer formation



The hydrophobic valine is exposed in the deoxy conformation and burrows within a hydrophobic pocket of neighboring Beta chains



**Hemoglobin  
polymers**

## Simple pathology-complex disease

- Presence of HbS leads to a variety of clinical manifestations- Cool pathophysiology at work!

Infection

Chest crises

Strokes

Splenic sequestration

Renal failure

Painful crises

Priapism

Pulmonary htn

Avascular necrosis

## Simple pathology-complex disease

- Disease severity variable and difficult to predict
  - Genotype (SS vs SC vs Sthal)
  - Beta globin gene cluster haplotype
    - Senegal most benign, central african republic most severe
  - Fetal hemoglobin levels-Higher is better
  - “Bad” in infant: dactylitis, severe anemia, leukocytosis
- Steinberg MH. Predicting clinical severity in sickle cell anaemia.Br J Haematol. 2005 May;129(4):465-81.

### Incidence of SCD complications based on genotype

Complication	HbSS (per 100 pt-years)	HbSC (per 100 pt- years)	HbS/beta <sup>0</sup> - thalassemia (per 100 pt-years)	HbS/beta <sup>+</sup> - thalassemia (per 100 pt-years)
Acute Chest Syndrome	12.8	5.2	9.4	3.9
Bacteremia	6.5	5.8		
Stroke	0.61	0.17	0.10	0.11
Pain crisis	24.0	8.5		

### Survival by genotype

Sex and Genotype	Median Survival
Males with Hb SS	42 years
Females with Hb SS	48 years
Males with Hb SC	60 years
Females with Hb SC	68 years

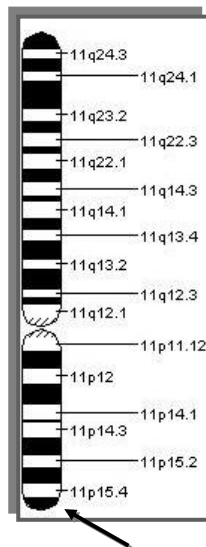
- In 1973 mean survival with SS disease was 14 years!

Care for sickle cell patients has begun to focus on screening and prevention in addition to symptom management

*Determinazione degli Aplotipi del cluster genico della  $\beta$ -globina in pazienti affetti da Anemia Falciforme*

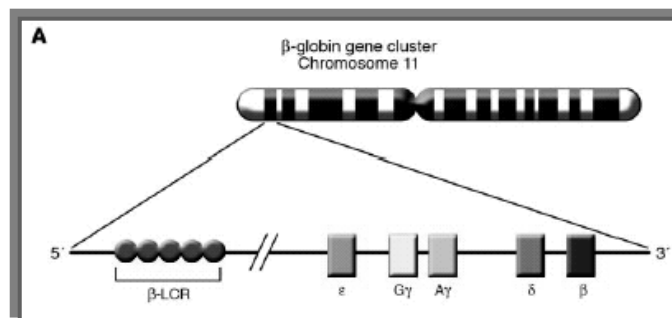


*Martina Montagnana, Giuseppe Lippi*



Premessa (1)

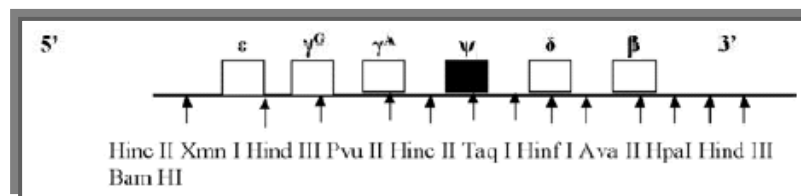
- Il cluster genico della  $\beta$ -globina è localizzato in posizione 11p15.5
- DNA 70Kb



## Premessa (2)

Lungo il cluster sono stati individuati dei siti specifici (più di 20) che possono essere riconosciuti e quindi tagliati da determinati enzimi (endonucleasi) di restrizione

(Antonarakis et al. *Hum Genet* 1985;69:1)



## Aplotipo



Si definisce aplotipo una combinazione di varianti alleliche lungo un cromosoma o segmento cromosomico contenente loci in linkage disequilibrium

Sickle Hemoglobin (*Hb S*) Allele and Sickle Cell Disease: A HuGE Review

American Journal of  
EPIDEMIOLOGY

A. Ashley-Koch,<sup>1</sup> Q. Yang,<sup>2</sup> and R. S. Olney<sup>2</sup>



## Determinazione degli aplotipi



### 1.1. $\beta$ -Globin-like Gene Cluster Haplotypes in Biology, Medicine, and Anthropology

Haplotypes of the  $\beta$ -globin-like clusters have been used for the following purposes:

1. **To provide anthropological correlations:** They have been used to give evidence of and/or define the common origin and the likelihood of an ancestral home for the tribals of India (13) and their potential origin in the Harappa culture, in the margins of the Indus River; to give a biological basis to the linguistic basis of the Bantu expansion hypothesis in Africa (8); and determine the Indian tribal origin and east African origin of the sickle gene, respectively, in Indian and African inhabitants of Mauritius Island (14).
2. **To provide a source of clinical diversity among SC patients:** Evidence exists that the linkage of the  $\beta^S$  gene to the Senegal and Arab Indian India haplotypes is associated with higher expression of HbS in SS and more benign hematological profile (15-17). Conversely, the Bantu haplotype has the most severe course (18).

## Correlazione genotipo-fenotipo nell'anemia falciforme



Stessa mutazione ma diversa clinica

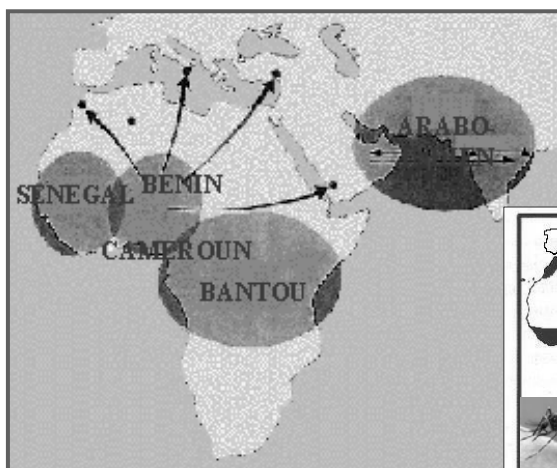


Aplotipo?

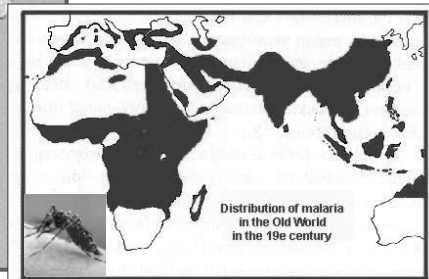


- Nei pazienti affetti da anemia falciforme sono stati identificati cinque diversi aplotipi maggiori che risultano in L.D. con la mutazione caratteristica  $\beta^S$ .

While many haplotypes exist for the  $\beta$ -globin cluster region, only specific haplotypes are found on chromosomes that carry the *Hb S* variant (18). These haplotypes are named for the geographic regions of Africa and the Middle East where they predominate (18).



Origine degli aplotipi



(Pagnier et al. Proc Natl Acad Sci USA 1984;81:1771-73)

## Altri aplotipi riconosciuti nel cluster $\beta$ -globinico

### Linkage of $\beta$ -thalassaemia mutations and $\beta$ -globin gene polymorphisms with DNA polymorphisms in human $\beta$ -globin gene cluster

Stuart H. Orkin<sup>1</sup>, Haig H. Kazazian Jr<sup>1</sup>, Stylianos E. Antonarakis<sup>1</sup>,  
Sabra C. Goff<sup>1</sup>, Corinne D. Boehm<sup>1</sup>, Julianne P. Sexton<sup>1</sup>,  
Pamela G. Waber<sup>1</sup> & Patricia J. V. Giardina<sup>1</sup>



Nature Vol. 296 15 April 1982

Haplotype	Thalassaemia defect identified
I	IVS-1 $\beta^+$
II	Nonsense codon 39
III	5' IVS-2 splice
IV	Frameshift codon 8
V	5' IVS-1 splice 5' IVS-2 splice
VI	IVS-1 consensus substitution
VII	IVS-2 position 745
VIII	-87 substitution
IX	Nonsense codon 39

### Sickle Hemoglobin (*Hb S*) Allele and Sickle Cell Disease: A HuGE Review

The  $\beta$ -globin cluster haplotypes are associated with differing clinical severities in sickle cell disease. This is probably due to variation in hemoglobin and fetal hemoglobin concentrations. Both hemoglobin and fetal hemoglobin levels vary with respect to haplotype (19) and are correlated with clinical expression of sickle cell disease (14, 20).

A. Ashley-Koch,<sup>1</sup> Q. Yang,<sup>2</sup> and R. S. Olney<sup>2</sup>

American Journal of  
**EPIDEMIOLOGY**

**The Senegal haplotype remains associated with a higher level of HbF compared to other African haplotypes**

[Nagel et al. N Engl J Med 1985;312:880-884]

Among the three most common haplotypes in sickle cell disease, the Senegal haplotype is associated with the most benign form of sickle cell disease, followed by the Benin haplotype. The Central African Republic haplotype is associated with the most severe form of the disease (19). In Africa, as well as in the United States, sickle cell patients with the Central African Republic haplotype have a twofold increased risk of complications and early mortality when compared with sickle cell patients with other haplotypes (19).



**Flussi migratori  
dall'Africa**

**Studi di popolazione  
hanno mostrato  
che il 13% dei  
neonati affetti da  
emoglobinopatie si  
trovano nei paesi  
Occidentali  
(Europa ed Italia),  
a causa dei flussi  
migratori**



It is therefore vital that international health agencies and governments of countries where the haemoglobin disorders occur at a high frequency become aware of the future extent of this problem and develop programmes for their control and management.

## **Inherited haemoglobin disorders: an increasing global health problem**

D.J. Weatherall<sup>1</sup> & J.B. Clegg<sup>2</sup>

*Bulletin of the World Health Organization, 2001, 79: 704–712.*



## Scopo dello studio



- Sviluppare una metodica di laboratorio per lo studio degli aplotipi, allo scopo di determinare la prevalenza e la distribuzione degli aplotipi nei soggetti con anemia falciforme residenti in Italia

## Popolazione studiata



- 68 pz.
- Sesso: 30 maschi e 38 femmine
- Età: 1-59 anni
- 18 SS, 38 AS, 4 SC, 8 S/beta talassemia



62 Africa  
1 Italia  
1 Cuba  
1 Rep. Dom.  
1 Albania  
2 Brasile

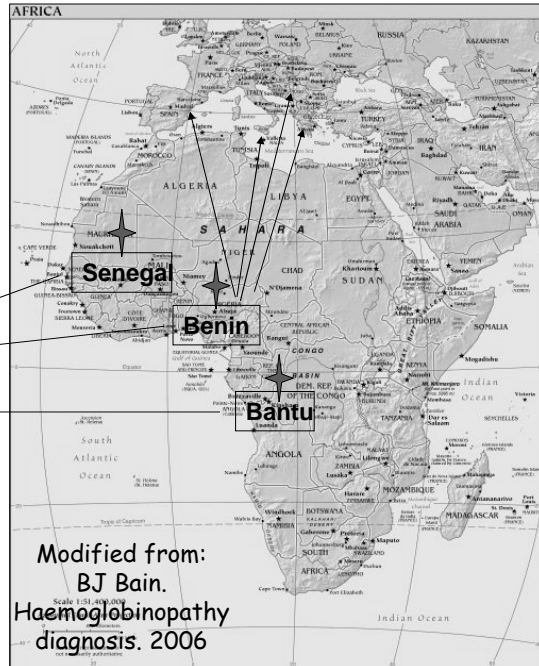
Rogers ZR, et al.  
Nonblack patients with sickle cell disease have African beta S gene cluster haplotypes.  
JAMA. 1989;261:2991-4.

18 pazienti non africani affetti da AF.

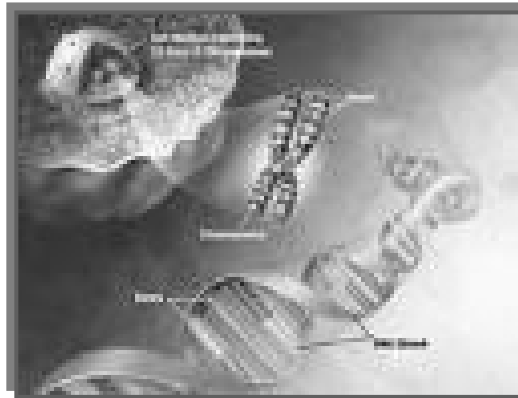
"These data strongly support the concept that the beta s gene on chromosome 11 of these individuals is of African origin.

The clinical severity of the disease in these nonblack patients is appropriate to their haplotype and is comparable with that of black patients."

## Migration from Africa to...



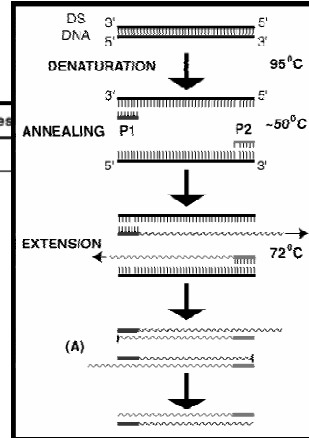
## 1. Estrazione del DNA



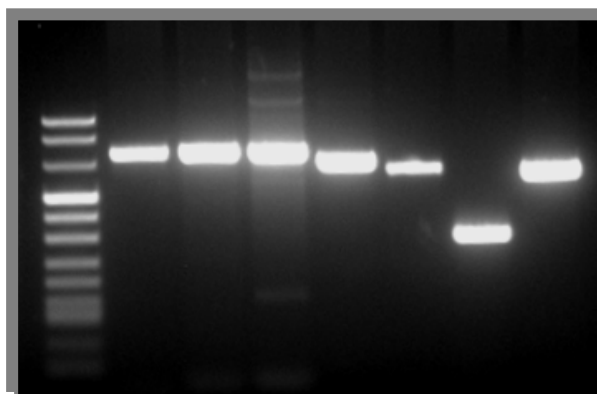
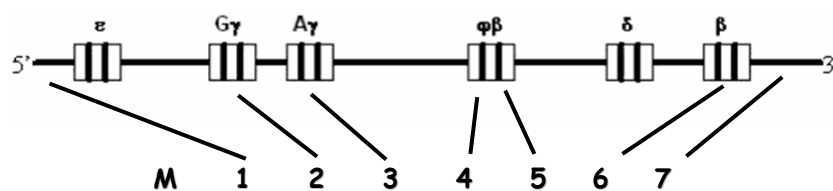
## 2. Amplificazione di 7 frammenti del cluster beta-globinico

TABLE I. Oligonucleotide Primers Used for PCR-RFLP Analysis of Seven RFLP Sites

Genes	Primer sequence 5-3	Product size (bp)
$\epsilon$ 5'	TCTCTGTTTGATGACAAATTC	760
$\alpha_1\gamma_1/\alpha_2\gamma_2$	AGTCATTGGTCAAGGCTGACC	1/2 = 781 2/3 = 766
	1. AAGTGTGGAGTGTACATGA	
	2. TGCTGCTAATGCTTCATTACAA 3. TAAATGAGGAGCATGCACAC	
5' $\psi$ $\beta$	GAACAGAAGTTGAGATAGAGA	701
3' $\psi$ $\beta$	ACTCAGTGGTCTGTGGGCT	592
	TCGCAATGACTCTGTAGC	
5' $\beta$	GGACCCTAACTGATATAACTA	328
	GTTGGTCTACCCCTGGACCCAGAGG	
3' $\beta$	TTCGTCGTGTTCCCATTCCTAACT	638
	AGTTAGAGGCTTGATTTGGAGG	
	GTTAAGGGGTTGATGTGAAC	



Primer sequences were those given by Old [Old JM. Hemoglobinopathies. In: Elles R, editor. *Methods in molecular medicine: molecular diagnosis of genetic disease*. Totowa, NJ: Humana Press; 1996:169-183] with minor modifications.





## Digestione con enzimi di restrizione

I prodotti della PCR di ciascun pz. (7 per ogni pz.) sono stati incubati con l'appropriato enzima di restrizione (3h 37°C) ed i frammenti risultanti sono stati fatti migrare su gel di agarosio al 3%.

Siti di restrizione studiati:

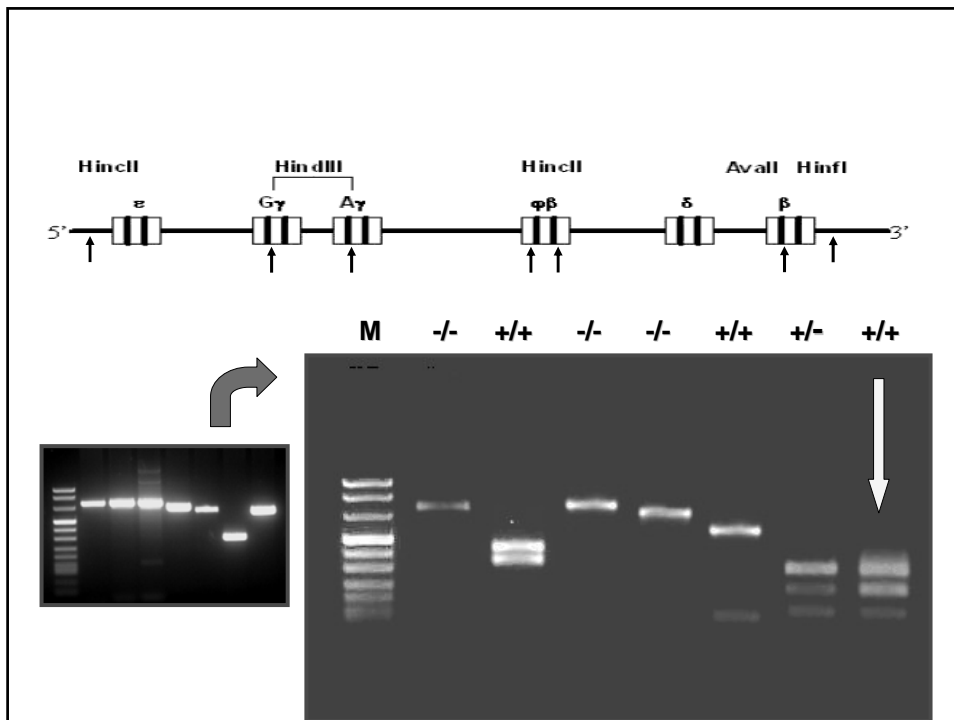
HincII (5' to  $\epsilon$ ), HindIII (within IVS2 of the  $G\gamma$  and  $A\gamma$ ), HincII (in 5' and 3' of  $\psi\beta$ ), AvaII (5' to  $\beta$ ) and HinfI (3' to  $\beta$ ).



## Analisi dei risultati

Per determinare gli aplotipi eterozigoti, abbiamo assunto che i pazienti eterozigoti possedessero un aplotipo comune ed un aplotipo raro, come riportato da Steinberg, piuttosto che due diversi aplotipi rari.

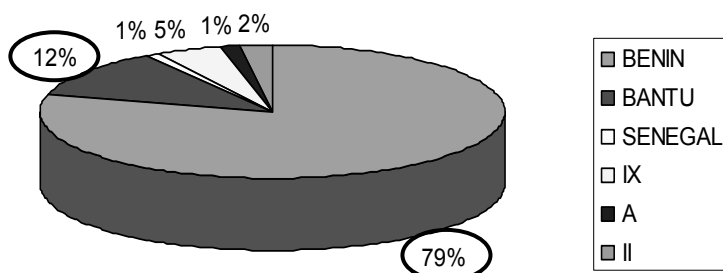
(Steinberg et al. Am J Hematol 1995;48:175-181)



**Costruzione dell'aplotipo**

	Hinc II	Hind III	Hind III	Hinc II	Hinc II	Ava II	Hinf I
<b>Benin</b>	-	-	-	-	+	+	+
<b>Bantu</b>	-	+	-	-	+	+	+
<b>Arab-Indian</b>	+	+	-	+	+	+	+
<b>Senegal</b>	+	+	-	+	+	+	+
<b>Cameroon</b>	-	+	+	-	+	+	-

## Frequenza determinata sugli 86 cromosomi S



### Original Research Article

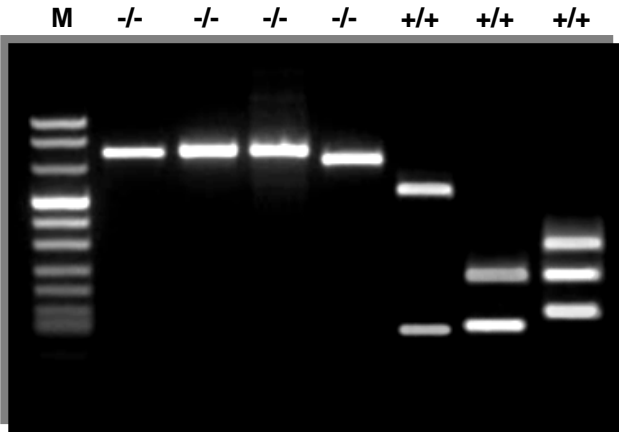
AMERICAN JOURNAL OF HUMAN BIOLOGY 00:000-000 (2008)

### $\beta$ -Globin Gene Cluster Haplotypes and $\alpha$ -Thalassemia in Sickle Cell Disease Patients from Trinidad

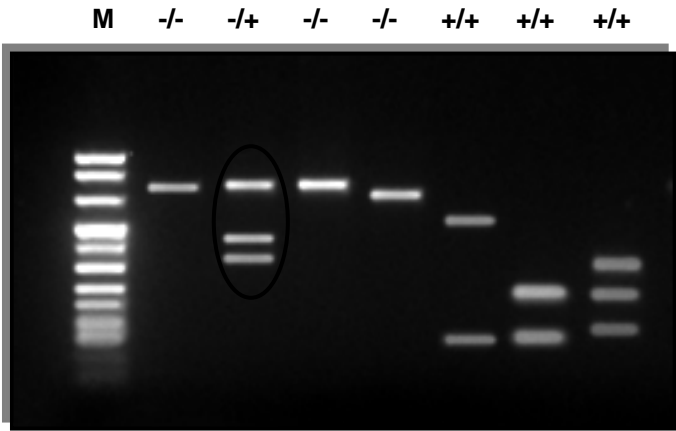
ALTHEIA JONES-LECOINTE,<sup>1</sup> ERSKINE SMITH,<sup>1</sup> MARC ROMANA,<sup>2</sup> MARIE-GEORGES GILBERT,<sup>1</sup> WAVENEY P. CHARLES,<sup>1</sup> CHRISTIAN SAINT-MARTIN,<sup>3</sup> AND LISIANE KÉCLARD<sup>2\*</sup>

$\beta^S$ Haplotype	Cuba <sup>a</sup>	Guadeloupe <sup>b</sup>	Jamaica <sup>c</sup>	Trinidad <sup>d</sup>
N <sup>e</sup> 86	198	832	244	283
Benin	<b>79%</b> 101 (51%)	622 (74.8%)	184 (75%)	175 (61.8%)
Bantu	<b>12%</b> 81 (41%)	92 (11.1%)	24 (10%)	49 (17.3%)
Senegal	<b>1%</b> 16 (8%)	51 (6.1%)	7 (3%)	24 (8.5%)
Cameroon	NA	19 (2.3%)	5 (2%)	10 (3.2%)
Arab-Indian	NA	6 (0.7%)	2 (1%)	9 (3.2%)
Other	NA	42 (5%)	22 (9%)	16 (5.7%)

Es. 1: aplotipo Benin-Benin



Es. 2: aplotipo Benin-Bantu



## Conclusioni

I nostri dati rivelano delle frequenze sovrapponibili a quelle osservate in altri paesi



Rimaniamo in attesa dei dati clinici per verificare la nostra ipotesi di partenza...