



# Sitting height/Height ratio: An indicator for genetic study of the SHOX gene in children with disharmonic short stature.

D. Yeste, N. Vargas, M. Domínguez, A. Campos, M. Clemente, P. Fernández (\*), A. Plaja (\*), A. Carrascosa .  
 Paediatric Endocrinology Unit. (\*) Genetic and Molecular Unit. Hospital Vall d'Hebron. Barcelona. Spain.  
 Paediatric Endocrinology Department, Vall d'Hebron University Hospital, Barcelona.  
 (\*) Genetic and Molecular Unit. Vall d'Hebron University Hospital, Barcelona. Spain.

## INTRODUCTION

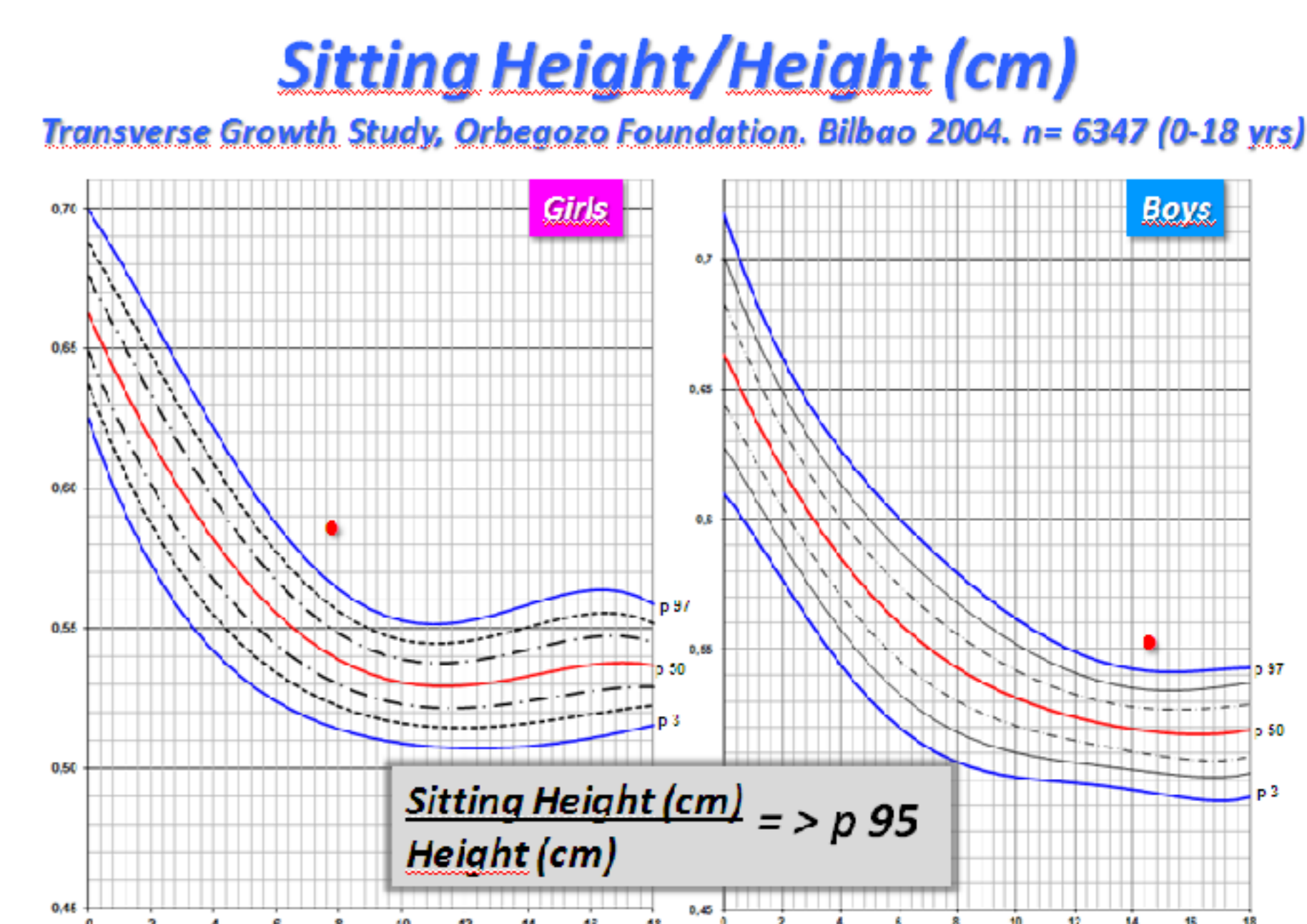
- Gene *SHOX* defects are the most prevalent cause of genetic short stature.
- Gene *SHOX* haploinsufficiency due to deletions or mutations in heterozygosis causes a wide spectrum of phenotypes ranging from very severe disharmonic short stature (S. Léry-Weil, S. Turner) to very mild forms with the appearance of idiopathic short stature (IST) of difficult clinical recognition.
- Auxological study directed at evaluating body disproportions such as the sitting height/height (SH/H) ratio in patients with IST has been postulated as useful for orienting the study of the *SHOX* gene.

## AIMS

- To establish the prevalence of *SHOX* gene defects in children with disharmonic short stature evaluated by the (SH/H) ratio regardless of the presence of dysmorphic features and radiological anomalies.

## PATIENTS & METHODS

- Prospective study of 37 consecutive patients with height < -2 SD and (SH/H) ratio > +2SD.
- All patients initially underwent Genetic Study using:
  - MPLA (P018, MRC Holland) or CGH array (Desing ISCA 8x60, Agilent).
  - Sequencing of all exons of the *SHOX* gene and flanking intronic regions was carried out in patients without *SHOX* gene deletion or its regulating regions.
- Normal Growth Reference Patterns:
  - Height (Spanish Growth Study 2010).
  - Sitting Height/Height (Orbegozo F. Cross-sectional Study. Bilbao 2001) An Esp Ped 2002,56: suppl. 4, 141.



## RESULTS

### SHOX GENE DEFECTS IN CHILDREN WITH DISHARMONIC SHORT STATURE

	n = 11	Age (x ± SD)	Height (x ± SD)	X-ray	Disharmonic Familial Height
<b>SHOX Gene defects</b>	8 (6 M, 2 V) 2 Complete deletion 2 Complex reorganisation of regulating region. 1 Regulating region deletion 1 Partial deletion 1 Duplication 0.56 Mb 1 Mutation in exon 5 (p. Ala257dup)	9.9 ± 3.3	-2.7 ± 0.9	n= 6 -Madelung deformity	6/8
<b>Turner syndrome</b>	3 46 XX(65%), 45 XO 46 X, +mar1 45 XO	9.3 ± 3.3	-3.0 ± 0.9	n=2 -5 <sup>o</sup> MT short -4 <sup>o</sup> MT short -Madelung deformity	0/3

**FREQUENCY GEN SHOX DEFECTS (n= 11): 29.7%**

### PATIENTS WITHOUT SHOX GENE DEFECTS IN CHILDREN WITH DISHARMONIC SHORT STATURE

	n = 26	Age (x ± SD)	Height (x ± SD)	X-ray	Disharmonic Familial Height
<b>Disharmonic</b>	24 (14 M, 10 V) 1 PseudoHPT Phenotype	10.2 ± 1.1	-2.5 ± 1.1	•5 Not done •7 Normal •7 Radial incurvation and/or short metacarpus •4 Madelung def	•Yes - 8 •No - 8 •Not done - 9
<b>Harmonic</b>	2 (1 M, 1 V)	9.3 ± 3.3	-2.7 ± 0.1	1 Radial incurvation and disalignmet	2/2

## CONCLUSIONS

1. The frequency of *SHOX* gene defects in our cohort with disharmonic IST evaluated by the (SH/H) ratio was 29.7%.
2. The (SH/H) ratio is a highly useful parameter for identifying patients with disharmonic IST and orienting *SHOX* gene study.
3. A significant proportion of patients with disharmonic IST remain undiagnosed, which renders this an open field for clinical research.