



Published in final edited form as:

*Clin Endocrinol (Oxf)*. 2017 December ; 87(6): 874–876. doi:10.1111/cen.13400.

## Isolated Growth Hormone Deficiency due to the R183H Mutation in *GH1*: Clinical Analysis of a Four-Generation Family

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### Dear Editor

The R183H mutation in the growth hormone gene (*GHI*) is a well-described genetic variant that causes autosomal dominant isolated growth hormone deficiency (IGHD) type II. Previous studies have demonstrated that individuals with this mutation have releasable growth hormone (GH) stores, but such release is severely impaired<sup>1</sup>. Hess et al. reported variable height deficits (−4.5 to −1.0 SDS), variable IGF-I concentrations (−2.9 to −0.8 SDS), and low but detectable, or even normal stimulated peak GH in several patients with the R183H mutation<sup>2</sup>. In contrast, IGHD type IB (caused by homozygous *GHI* or *GHRHR* mutations) results in very low but measurable stimulated GH, while IGHD type IA (caused by homozygous deletions and nonsense mutations in *GHI*), results in complete absence of GH leading to a more severe phenotype, reflecting a spectrum of growth hormone deficiency (GHD).

Adult GHD (AGHD) causes a distinct phenotype with significant morbidity including increased fat mass, decreased muscle mass and exercise capacity, decreased bone mineral density (BMD), and decreased quality of life, in addition to an abnormal cardiovascular risk profile. However, the majority of studies supporting these findings were conducted in patients with hypopituitarism, which is the most common cause of AGHD. There is limited information regarding the clinical phenotype of patients with IGHD - especially related to the comorbidities seen in AGHD. As the R183H mutation leads to decreased GH secretion, but not a complete absence of GH, it is unknown what the long-term effects of this mutation are in adult carriers. To gain insights into this question, we performed a comprehensive clinical evaluation of a four-generation family with six members affected by the R183H mutation in *GHI*.

The proband (Subject IV.2) was referred to the endocrinology clinic at age 5 years for evaluation of proportional short stature (height SDS −3.0). Her initial workup revealed low

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**Conflict of Interest Statement:** The authors have nothing to disclose.

IGF-I and low IGF binding protein-3 concerning for GHD. However, her GH stimulation test with arginine and clonidine revealed a normal peak GH concentration of 8.7 ng/mL (normal 5.2 using chemiluminescence assay). On further history we identified several family members with short stature that followed a dominant mode of inheritance (Figure 1). Via whole exome sequencing, we found that the affected individuals carried the heterozygous R183H mutation in *GHI*. The adults underwent a comprehensive clinical evaluation including GH stimulation test using glucagon, bone densitometry, and complete cardiovascular risk assessment (Supplementary Data). The affected children underwent GH stimulation testing using arginine and clonidine.

The results are summarized in Table 1. We found that our patients had variable degrees of short stature and low-normal IGF-I concentrations. Interestingly, the patients had marked variability on the GH stimulation test; the affected children and one adult had delayed stimulated peak GH secretion, and two of the affected adults had absent responses. Hess et al. hypothesized that the variability in GH secretion could be explained by a mild dominant-negative effect of R183H-GH on the wild-type GH, and possibly due to compensation of the GH-insufficiency by additional genes<sup>2</sup>. Additionally, Zhu et al. proposed that the R183H mutation results in prolonged retention of GH molecules in the secretory granules affecting the rate of secretory granule release<sup>3</sup>. This mechanism could explain the delayed peak GH response observed in some of the patients from this family.

A thorough cardiovascular risk assessment including echocardiography and noninvasive arterial imaging for carotid thickness and arterial stiffness (pulse wave velocity, augmentation index and brachial flow-mediated dilation) did not suggest increased cardiovascular risk among the adult patients in this family with IGHD secondary to the R183H mutation (Supplemental Table 1). In addition, all three adult patients had normal biochemical parameters including lipid profile, apolipoprotein B, and inflammatory markers. One individual with obesity was found to have evidence of insulin resistance but it is difficult to ascribe this to her GHD given that the other two adult subjects had normal glucose metabolism. Similarly, previous studies have shown that patients with IGHD type IB due to GHRHR mutations have normal echocardiographic parameters and normal carotid thickness with no evidence of accelerated atherosclerosis<sup>4</sup> and not surprisingly, normal longevity<sup>5</sup>. However, these patients exhibit increased total and LDL cholesterol, in addition to increased inflammatory markers which is not present in our patients. Nonetheless, using a DXA scan, we found that the two younger adult patients from our family had central obesity and increased fat mass, comparable to previous reports in IGHD type IB<sup>6</sup>. However, the oldest individual had a normal BMI and body composition suggesting that the findings on the younger patients could be at least partially due to lifestyle choices. Lastly, we found that all the patients had normal bone mineral density (BMD) compared to age- and sex-matched controls. Likewise, patients with IGHD type IB have normal volumetric BMD but they have a high prevalence of hip joint problems and genu valgum<sup>7</sup> which was not present in our patients.

In summary, this is the first study describing in detail the adult phenotype of a family with several members affected by familial IGHD type II. We acknowledge that it is difficult to draw definitive generalizable conclusions from a single family study. However, the detailed

investigation of this family with a known pathogenic *GHI* mutation does provide some important insights into the potential effects of this mutation on adult manifestations of IGHD. While there are many factors that can affect co-morbidities such as hyperlipidemia, decreased BMD, and increased cardiovascular risk, it is notable that our patients do not consistently manifest any of these co-morbidities, even in the elderly great-grandmother. This suggests that in this family, while the R183H mutation has significant effects on childhood linear growth, there is sufficient residual GH effect to meet the metabolic needs of adults carrying this mutation. Combining our present results with those from earlier studies, we conclude that familial IGHD results in different phenotypes leading to diverse outcomes, where IGHD type IA causes the most severe phenotype with undetectable GH concentrations resulting in severe short stature and reduced longevity<sup>8</sup>; followed by IGHD type IB, which results in short stature with some metabolic abnormalities; and lastly, the least severe phenotype, IGHD type II. The R183H mutation in *GHI* results in partial GHD that appears to have a milder presentation when compared to other forms of IGHD. Therefore, GH therapy may not be required for the adult patients that participated in this study. In addition, patients with IGHD type II can have normal stimulated GH concentrations that can result in inappropriate diagnosis and treatment, thus pointing out the limitations of current growth hormone stimulation tests and the potential benefits of genetic testing. Furthermore, our findings support the need for additional research into the manifestations of adult GHD in other patients with the R183H *GHI* mutation and other genetic etiologies of GHD.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

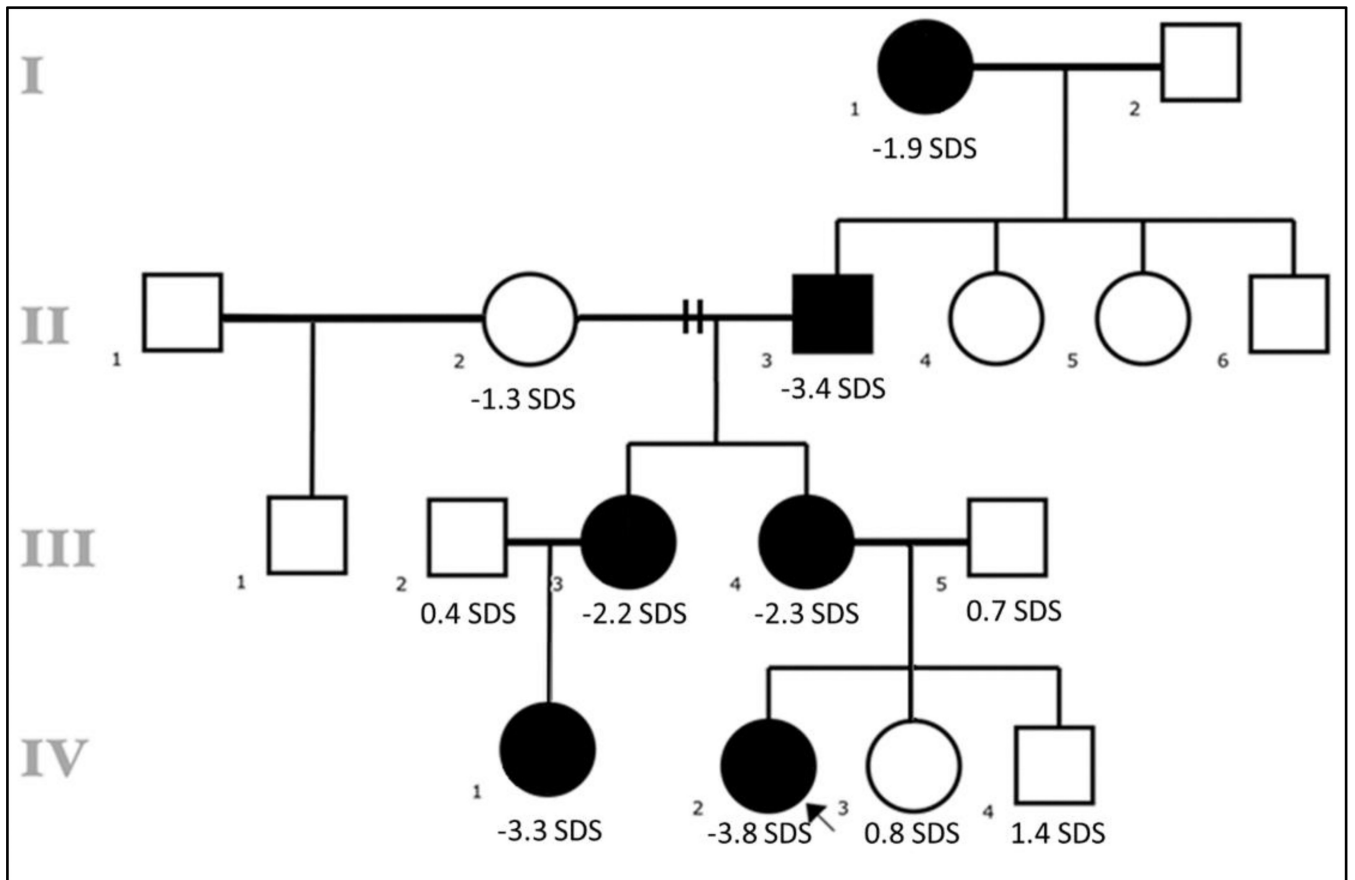
## Acknowledgments

This work was supported by grant K23HD07335 (to A.D.) from the Eunice Kennedy Shriver National Institute of Child Health & Human Development of the National Institutes of Health. Additional support was provided by the National Center for Advancing Translational Sciences, National Institutes of Health, through Grant 1UL1 TR001425. The content is solely the responsibility of the authors and does not necessarily represent the official views of the NIH.

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**Figure 1.** Family pedigree with affected individuals indicated by black squares and circles. When known, the height SD score is reported.

Table 1

Anthropometric characteristics, laboratory measurements, BMD and body composition.

	IV.1	IV.2	III.3	III.4	I.1
Gender	Female	Female	Female	Female	Female
Age (years)	6.6	5.9	30	33	73
Weight (kg)	19	12	72	96	43
Height (cm)	103	96	148	149	151
Height Z-score	-3.3	-3.8	-2.3	-2.2	-1.9
BMI (kg/m <sup>2</sup> )	18	14	33	43	18
<b>Laboratory measurements</b>					
IGF-I (µg/L)	53.3	27.4	95	117	47
IGF-I ref. range for age	45 – 254	33–276	75 – 253	72 – 241	36 – 164
IGFBP-3 (µg/L)	2801	1260	4702	4222	2518
IGFBP-3 ref. range for age	1983–5461	2169–4790	2654 – 5982	2562–5783	1999 – 5543
Peak GH (µg/L)	5.2	8.7	0.5	0.4	9.3
ALS (mg/L)			20	15	6.5
Total cholesterol (mmol/L)			4.2	4.4	5.3
LDL-C (mmol/L)			2.2	2.8	3.6
HDL-C (mmol/L)			1.3	1.2	1.1
Triglycerides (mmol/L)			1.5	1.0	1.4
Apolipoprotein B (g/L)			0.88	0.94	1.1
Glucose (mmol/L)			5.1	6.5	4.3
Insulin (pmol/L)			75	144	13
Hemoglobin A1c (%)			4.8	5.5	4.8
IL-6 (pg/mL)			0.0	0.0	6.3
CRP (µg/mL)			1.2	1.8	0.4
VW factor Ag (kIU/L)			30	46	146
<b>BMD and body composition</b>					
LS BMD T- score			0	0.1	-0.9
Z-score			0	0.1	1.4
Hip BMD T-score			0.7	0.9	-2.3

	IV.1	IV.2	III.3	III.4	I.1
Z score			0.8	1	-0.3
FA BMD T-score			-0.4	-0.4	-0.8
Z-score			-0.2	-0.2	1.6
TB BMD T score			-0.6	0.5	-1.8
Z score			-0.7	0.3	-0.2
% Body fat (SDS for age)			44 (1.4)	50 (2.0)	44 (0.8)

Abbreviations: ALS, acid labile subunit. BMD, bone mineral density. LS, lumbar spine. TB, total body. FA, forearm. Normal reference ranges: ALS: 7–16 mg/L, total cholesterol: <5.2 mmol/L, LDL-C: <4.1 mmol/L, HDL-C: >1.0 mmol/L, triglycerides: <2.2 mmol/L, apolipoprotein B: 0.6 – 1.17 g/L, glucose <5.5 mmol/L, insulin: <102 pmol/L, Hemoglobin A1C: <5.7%, IL-6: <7 pg/mL, CRP: <5 µg/mL, WV factor Ag: 50–100 kIU/L.