Analysis of the clinical and molecular characteristics of a child with achondroplasia: A case report

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Abstract. Achondroplasia (ACH) is a hereditary dwarfism caused by the disturbed proliferation and differentiation of growth plate chondrocytes, followed by impaired endochondral bone growth. ACH is caused by mutations in the gene encoding the transmembrane receptor, fibroblast growth factor receptor 3 (FGFR3). In total, >90% of patients with ACH have a G1138A mutation in the transmembrane domain of the FGFR3 gene. Patients with ACH usually have no growth hormone (GH) deficiency. The current study presents the case of a four-year-old male with clinical manifestations suggestive of ACH, including a large head, prominent forehead, short upper arms and legs, and short hands with fingers assuming a trident position. The patient showed normal responses to GH provocation tests with L-dopa (peak GH concentration, 42.38 ng/ml) and insulin (peak GH concentration, 23.29 ng/ml during hypoglycemia), but a blunted response to a GH provocation test with arginine (peak GH concentration, 7.31 ng/ml). Furthermore, the GH concentration during exercise was low (4.8 ng/ml). Magnetic resonance imaging revealed a decreased pituitary volume. Thyroid function tests and the levels of sex hormones (follicle stimulating hormone, luteinizing hormone, estradiol, prolactin and progesterone), cortisol and adrenocorticotropic hormone were normal. A heterozygous G1138A mutation within the FGFR3 gene was detected, confirming the diagnosis of ACH. Thus, recombinant human GH therapy (0.1 IU/kg/day) was initiated. At the six-month follow-up, the height, arm span-to-height ratio and lower limb length-to-height ratio of the patient had increased, while the head circumference had decreased. The present results corroborate the finding that the G1138A mutation within FGFR3 is the most common ACH-causing mutation in different populations. GH may be beneficial in the treatment of short stature in ACH patients with subnormal GH secretion.

Introduction

Achondroplasia (ACH) is a hereditary dwarfism caused by a disturbance in the proliferation and differentiation of growth plate chondrocytes, followed by an impairment in endochondral bone growth. The incidence rate of ACH is ~1/15-40,000 live births (1). In total, between 80 and 90% of ACH cases are sporadic (2). Newborn infants with ACH typically present with disproportionate shortening of the limbs, a long and narrow trunk, a large head with frontal bossing and midfacial hypoplasia. The hands are short and broad, and frequently exhibit a three-pronged (trident) configuration. Furthermore, numerous joints, with the exception of the elbow, are hyperextensible (3). The disease shows an autosomal dominant inheritance and is caused by mutations in the gene encoding the transmembrane receptor, fibroblast growth factor receptor 3 (FGFR3), which is an important regulator of linear bone growth. FGFR3 is expressed in various tissues including the cartilage, brain, kidneys and the intestines at different stages of development. FGFR3 mutations generate deficient proteins that affect chondrocyte proliferation and calcification, and hinder cartilage growth and development, thereby resulting in an external phenotype of ACH (4). The human FGFR3 gene is located on chromosome 4q16.3. In total, >90% of patients with ACH have a G1138A mutation in the transmembrane domain of the FGFR3 gene (5,6). Research on ACH began later in China than in Europe and the US. Currently, almost 60 clinical cases have been reported around the country and there have been no studies, to the best of our knowledge, on the incidence rate of ACH in China. In the present study, the clinical characteristics of a Chinese male child diagnosed with ACH were analyzed, and tests for the FGFR3 gene mutation were performed on the patient and patient's family.

Case report

Patient characteristics and clinical observations. A four-year-old male was admitted to the First Hospital of Lanzhou University (Lanzhou, China) with growth retardation since the age of three years. The patient was an only child, and was born at full term via vaginal delivery with a birth weight of 3,900 g. Teething began between seven and eight
months, and the patient was walking at one year. After one year of age, it was noted that the patient's growth and development were slow, and that his height was lower compared with other children of a similar age. The annual increase in height was <5 cm; however, the weight and intelligence level were normal. A physical examination at the time of admission revealed the following characteristics: Body temperature, 36°C; pulse rate, 90 bpm; respiratory rate, 20 breaths/min; height, 85 cm; and weight, 16 kg. The patient had a large head with a prominent forehead. In addition, there was disproportionate shortening of the upper arms and legs, and the patient had short hands with fingers that assumed a three-pronged (trident) configuration.

Laboratory results were as follows: Serum calcium, 2.41 mmol/l (normal range, 2.10–2.80 mmol/l); serum phosphorus, 1.66 mmol/l (normal range, 0.70–1.60 mmol/l); intact parathyroid hormone, 24.20 pg/ml (normal range, 14–72 pg/ml); 25-hydroxy vitamin D, 72.6 mmol/l (normal range, 47.7–144 mmol/l); osteocalcin, 30.7 pg/ml (normal range, 18.8–55 ng/ml); bone-specific alkaline phosphatase, 89.7 ng/ml (normal range, 7.3–22.4 ng/ml); urine calcium, 1.30 mmol/24 h (normal range, 2.5–7.5 mmol/24 h); urine phosphorus, 13.75 mmol/24 h (normal range, 23–48 mmol/24 h); thyroid-stimulating hormone (TSH), 1.26 µIU/ml (normal range, 0.55–4.78 µIU/ml); triiodothyronine (T3), 1.69 ng/ml (normal range, 0.60–1.81 ng/ml); thyroxine (T4), 10.0 µg/dl (normal range, 4.50–10.9 µg/dl); free T3, 4.19 pg/ml (normal range, 2.3–4.2 pg/ml); and free T4, 1.34 ng/dl (normal range, 0.89–1.76 ng/dl). Test for antibodies against the TSH receptor, thyroid peroxidase and thyroglobulin were negative. In addition, the levels of sex hormones, cortisol and adrenocorticotropic hormone (ACTH) were normal. The patient's karyotype was 46,XY. A radiograph of the upper limbs and hands revealed the hands to be short and broad, with a trident configuration, and that the estimated age of the wrist bones was lower than the actual age of the patient (Fig. 1A). A radiograph of the lower limbs revealed disproportionate shortening of the limbs, bilateral trumpet-like enlargement of the distal femoral metaphyses and blurring of the epiphyseal shape (Fig. 1B). Furthermore, a magnetic resonance imaging (MRI) scan revealed a decreased pituitary volume and a hyperintense and spot-like corpus callosum, indicative of malacia, on the T1- and T2-weighted images. No similar phenotype was identified in the parents or other family members (grandfather, grandmother, aunts, uncles and cousins) of the patient.

**Growth hormone provocation tests.** Growth hormone (GH) responses to provocation tests (0.1 IU/kg insulin-induced hypoglycemia, 10 mg/kg L-dopa and 0.5 g/kg arginine) and the levels of GH during exercise (brisk walking for 15 minutes followed by running for 5 min with the heart rate reaching >120 beats/min; GH levels obtained 20 min after exercise initiation) were evaluated. Serum GH concentrations were determined using an immunoradiometric assay (Tianjin Nine Tripods Medical & Bioengineering Co., Ltd., Tianjin, China). The intra-assay variation was <5.8% and the inter-assay variation was <9.3%. The responses of cortisol and ACTH to insulin-induced hypoglycemia were evaluated simultaneously.

Peak GH levels of <10 ng/ml on GH provocation tests with insulin, L-dopa and arginine, and a GH level of <5 ng/ml during exercise were defined as blunted GH secretion, according to the criteria of the Foundation for Growth and Science in China.

**DNA amplification and mutation detection.** Exon 10 of the FGFR3 gene was amplified using polymerase chain reaction (PCR; Roche Diagnostics Co., Ltd., Shanghai, China). The sequence of the forward primer was 5'-AGGCCGGTGCTGAGTTCTGAG-3' and the sequence of the reverse primer was 5'-GGGATCTTTGTGCACCTGG-3' (Sangon Biotech Co., Ltd., Shanghai, China). All PCR amplifications were performed in a total volume of 50 µl, which contained 2 µl extracted DNA, 20 pmol each of the forward and reverse primers, 19 µl 2X Taq PCR Master mix (Takara Biotechnology Co., Ltd.) and 25 µl deionized water. Thermal cycling conditions were as follows: Initial activation of DNA polymerase at 95°C for 5 min, followed by 30 cycles of 95°C for 30 s, 58°C for 30 s and 72°C for 45 s, and a final extension at 72°C for 5 min. The PCR products were purified by a UNIQ-10 Column kit (Sangon Biotech Co., Ltd.) and sequenced directly on a ABI3730XL genetic analyzer (Applied Biosystems Life Technologies, Foster City, CA, USA) to detect the gene mutations. The sequence results were contrasted with the normal sequence of the FGFR3 gene obtained from GenBank. All mutations were confirmed through forward and reverse sequencing.

**Ethics statement.** Written informed consent was obtained from the patient's family. The study was approved by the Human Ethics Review Committee of the First Hospital of Lanzhou University and complied with the Declaration of Helsinki for Experimentation on Humans, 1975 (revised in 1983).

**Results**

**GH provocation tests.** GH responses to the provocation tests are shown in Fig. 2. The patient showed normal responses to the GH provocative tests using L-dopa (peak GH concentration, 42.38 ng/ml) and insulin (peak GH concentration was increased to 23.29 ng/ml during hypoglycemia). However, a slightly blunted response was observed for the GH provocation test with arginine (peak GH concentration, 7.31 ng/ml <10 ng/ml), and the GH level during exercise was low [4.8 ng/ml (<5 ng/ml)]. The cortisol and ACTH responses to insulin-induced hypoglycemia were normal.

**Gene mutation analysis.** Gene mutation analysis revealed that the patient had a G→A mutation at nucleotide 1,138 within exon 10 of the FGFR3 gene. This missense mutation caused the substitution of glycine with arginine at amino acid posi-
Figure 1. Radiographs of the upper and lower limbs. (A) Hands are short and broad, and exhibit a three-pronged configuration. The estimated age of the wrist bones is lower than the actual age of the patient. (B) Lower limbs show disproportionate shortening, bilateral trumpet-like enlargement of the distal femoral metaphyses and blurring of the epiphyseal shape.

Figure 2. GH response to the provocation tests. (A) GH concentrations following the provocation tests with L-dopa and arginine, and during exercise. (B) Response of cortisol, GH and ACTH to insulin-induced hypoglycemia. GH, growth hormone; ACTH, adrenocorticotropic hormone.

Figure 3. Partial sequencing results of the polymerase chain reaction products of exon 10 in the fibroblast growth factor receptor 3 gene. (A) Normal control (forward sequencing), homozygote GG. (B) Normal control (reverse sequencing), homozygote CC. (C) Patient with achondroplasia (ACH; forward sequencing), heterozygous AG. (D) Patient with ACH (reverse sequencing), heterozygous TC. The arrows indicate mutated bases.
tion 380 (G380R) in the FGFR3 protein. The mutation was heterozygous, and the results of the forward and reverse sequencing were consistent (Fig. 3). No mutation within exon 10 of the FGFR3 gene was observed in the samples taken from the patient's parents or from seven healthy controls (aged 10-35 years) from the First Hospital of Lanzhou University.

**Treatment and follow-up.** A diagnosis of ACH with subnormal GH secretion was considered on the basis of the clinical and laboratory results. Oral therapy with recombinant human growth hormone (rhGH; 0.1 IU/kg/day) and levothyroxine (12.5 μg/day) was initiated. After six months, the patient's height had markedly improved (to 93.5 cm). Furthermore, the arm span-to-height ratio and lower limb length-to-height ratio increased during treatment, while the head circumference decreased. The serum GH concentration was 60.37 ng/ml. No abnormality was found on the thyroid function and sex hormone tests.

**Discussion**

ACH, an autosomal dominant disorder, is the most common form of human dwarfism. The main clinical manifestation is an abnormally large head with a prominent forehead and flat nasal bridge, short upper arms and legs (rhizomelic dwarfism), an unusually prominent abdomen and buttocks, and short hands with fingers that assume a trident or three-pronged position during extension. In 1994, Francomano et al located the pathogenic gene of ACH to chromosome 4p16.3 through genetic linkage analysis (8). Soon after, Shiang et al reported a missense mutation at codon 380 in the transmembrane domain of the FGFR3 gene in patients with ACH (5, 6). A G → A displacement at nucleotide 1,138 in the FGFR3 gene is present in 90% of patients with ACH, while a G → C transversion is present in only a minority of patients (9, 10). Similar results have been reported in Chinese populations (11). G375C and G346E mutations on the transmembrane domain of the FGFR3 gene have also been observed in patients with ACH (12, 13). These results indicate a strong association between the transmembrane domain of the FGFR3 gene and ACH. Furthermore, Zhang et al reported a Ser217Cys mutation in the Ig II domain of the FGFR3 gene in a Chinese family with ACH (14).

FGFR3, a type of tyrosine receptor, comprises 806 amino acid residues and plays an important role in skeletal development. The length of the FGFR3 gene is ~15 kb, with 19 exons and 10 introns. Exon 10 encodes the transmembrane domain of the FGFR3 gene. As a membrane receptor, the structure of FGFR3 is comprised of three parts, including an intracellular region, a transmembrane domain and an extracellular region, the latter of which functions as a binding domain for numerous ligands, including three typical Ig-like structural domains (Ig I-III) (15). The extracellular region includes a near-nascent region, two conservative tyrosine kinase functional domains and an autophosphorylated C-terminal. The ligand, fibroblast growth factor (FGF), may attach to acetylated proteins on the surface of the cells and induce receptor dimerization and transautophosphorylation of the tyrosine kinase in the cytoplasm (16). The residual phosphate cellulose may be used as a docking site for proteins, subsequently resulting in the activation of various signaling pathways that mediate the effects of the FGF receptor with regard to the regulation of proliferation, differentiation and apoptosis in a number of cells (16).

In the present case report, the patient demonstrated clinical manifestations similar to those described in the literature on ACH, including a large head, prominent forehead, short upper arms and legs, and short hands with fingers that assumed a trident position. The results of the imaging examinations were also consistent with a diagnosis of ACH. Finally, a G → A displacement at nucleotide 1,138 within exon 10 of the FGFR3 gene was identified, which further confirmed that nucleotide 1,138 within the FGFR3 gene is a common site of mutations leading to ACH. The main reason that the FGFR3 mutation leads to ACH may be the suppression of the proliferation and differentiation of cartilage cells. The FGFR3 mutation causes dimerization of cell membrane proteins, and may lead to the continuous activation of intracellular tyrosine kinases in the absence of a ligand, which ultimately activates intracellular signaling pathways and suppresses the proliferation and differentiation of cartilage cells (17, 18). Matsuhashita et al reported that the extracellular signal-regulated kinase/mitogen-activated protein kinase signaling pathway in cartilage cells plays an important role in the development of ACH (19).

Patients with ACH do not generally have a GH deficiency (20). In the current study, the patient demonstrated a normal response to GH provocation tests with L-dopa and insulin; however, a blunted response was observed in the GH provocation test with arginine (peak GH concentration, <10 ng/ml). In addition, low GH concentrations were observed during exercise (<5 ng/ml). Therefore, a subnormal GH secretion was suspected. Furthermore, an MR scan revealed that the pituitary volume was decreased. Therefore, therapy with rhGH (0.1 IU/kg/day) was initiated, and marked results were observed within six months, which is consistent with previous studies (21-24). A large-scale study in Japan demonstrated that rhGH therapy promoted bone growth in patients with ACH by improving the Z scores for growth rate and height, and also had a dose- and time-dependent effect on ACH (25). A possible mechanism underlying the effects of GH treatment on ACH may be through the epiphyseal growth plate, which is the area of bone formation. GH stimulates local cartilage cells in this area to produce insulin-like growth factors, which promotes cartilage cell proliferation and thereby promotes growth (26).

In the present study, no mutation was observed in the FGFR3 gene of the patient's parents, indicating that the patient had a de novo mutation. ACH has been associated with advanced paternal age (2). As sperm cells are produced constantly throughout life, the risk of mutations in the sperm cells increases with age (21). This suggests that factors influencing DNA replication and repair during spermatogenesis may predispose to the occurrence of ACH-associated mutations. Therefore, the mutation in the current patient was likely attributable to mutations in the father's sperm cells.

In conclusion, the results of the present study further support the hypothesis that the G1138A mutation within the FGFR3 gene is the most common mutation causing ACH in various populations. For certain ACH patients with subnormal GH secretion, GH therapy may be beneficial for the treatment of short stature.
References


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