

Hutchinson–Gilford Progeria Syndrome with G608G LMNA Mutation

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Hutchinson–Gilford progeria syndrome (HGPS) is a rare condition originally described by Hutchinson in 1886. Death result from cardiac complications in the majority of cases and usually occurs at average age of thirteen years. A 4-yr old boy had typical clinical findings such as short stature, craniofacial disproportion, alopecia, prominent scalp veins and sclerodermatous skin. This abnormal appearance began at age of 1 yr. On serological and hormonal evaluation, all values are within normal range. He was neurologically intact with motor and mental development. An echocardiogram showed calcification of aortic and mitral valves. Hypertrophy of internal layer at internal carotid artery suggesting atherosclerosis was found by carotid doppler sonography. He is on low dose aspirin to prevent thromboembolic episodes and on regular follow up. Gene study showed typical G608G (GGC- > GGT) point mutation at exon 11 in LMNA gene. This is a rare case of Hutchinson–Gilford progeria syndrome confirmed by genetic analysis in Korea.

Key Words: Hutchinson Gilford Progeria Syndrome; LMNA; Progeria

INTRODUCTION

Progeria is a genetic disorder rarely encountered and is characterized by features of premature aging. It is also known as “Hutchinson-Gilford Progeria syndrome” (1). Although signs and symptoms vary in age of onset and severity, they are remarkably consistent overall. Children with HGPS usually appear normal at birth. Profound failure to thrive occurs during the first year (2). Characteristic facies, with receding mandible, narrow nasal bridge and pointed nasal tip develop. Motor and mental development is normal. Death occurs as a result of complications of severe atherosclerosis, either cardiac disease (myocardial infarction) or cerebrovascular disease (stroke), generally between ages 6 and 20 yr. Average life span is approximately 13 yr (2).

We report here a 4-yr-old boy with an apparently typical Hutchinson–Gilford progeria syndrome with G608G LMNA mutation.

CASE DESCRIPTION

A 4-yr-old boy was referred to the department of pediatrics with short stature and sclerodermatous skin on 2 September 2010. He was the first child born to non-consanguineous parents with no significant family history. He was born at full term with birth weight of 3.35 kg by spontaneous vaginal delivery. His abnormal appearance was not found at birth. At age of 1 yr, growth retar-

ation, hair loss, and alteration of skin color on the abdominal region began.

On physical examination, his length was 88 cm and weight 11.5 kg, both less than 3rd percentile and head circumference was 52 cm (ca. 77th percentile) (Fig. 1). He had generalized indurated and shiny skin associated with decreased subcutaneous fat, especially on the abdomen. His hair was fine and sparse and his scalp veins were easily visible (Fig. 1). His anterior fontanelle was still patent (horizontal diameter 1.7 cm, vertical diameter 1.4 cm). He had craniofacial disproportion for his age due to micrognathia, prominent eyes, scant eyelashes and small nose. His bone age was 3 yr. He was neurologically intact with motor and mental development. On serological and hormonal evaluation, all values are within normal range. An echocardiogram did not show concentric left ventricular hypertrophy nor increased left ventricular pressure but showed calcification of aortic and mitral valves (Fig. 2). Hypertrophy of internal layer at internal carotid artery suggesting atherosclerosis was found by carotid doppler sonography (Fig. 2). He is on low dose aspirin to prevent thromboembolic episodes and on regular follow up.

We obtained his DNA from white blood cell in peripheral blood and sequencing was performed by such method as follows; Samples were extracted using a MG™ Tissue Kit. We designed polymerase chain reaction (PCR) primers using the Primer 3 program. The primers Tms ranged between 59°C and 62°C as far as possible. The PCR reaction was performed with 20 ng of genom-



Fig. 1. (A) General appearance of the patient with characteristic prematurely aged appearance, retarded growth. (B) Lateral and top view of the head showing macrocephaly, alopecia, prominent scalp veins.

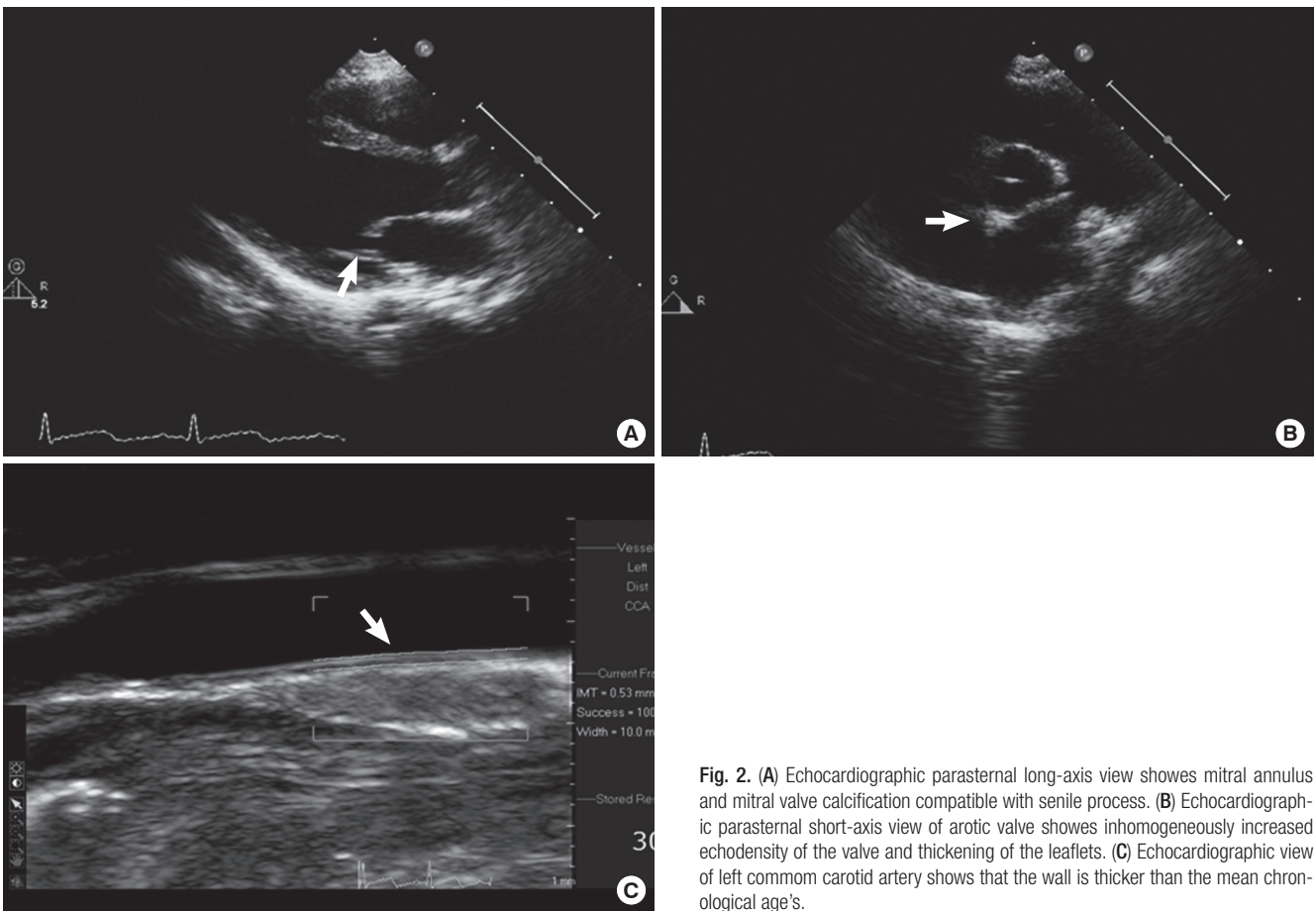


Fig. 2. (A) Echocardiographic parasternal long-axis view shows mitral annulus and mitral valve calcification compatible with senile process. (B) Echocardiographic parasternal short-axis view of aortic valve shows inhomogeneously increased echodensity of the valve and thickening of the leaflets. (C) Echocardiographic view of left common carotid artery shows that the wall is thicker than the mean chronological age's.

ic DNA as the template in a 30 μ L reaction mixture by using a EF-Taq (SolGent, Korea) as follows: activation of Taq polymerase at 95°C for 2 min, 35 cycles of 95°C for 1 min, 55°C-63°C, and 72°C

for 1 min each were performed, finishing with a 10 min step at 72°C. The amplification products were purified with a multi-screen filter plate (Millipore Corp., Bedford, MA, USA). Sequenc-

anticancer target when it was discovered that the oncoprotein Ras, which has been estimated to be involved in up to 30% of all human cancers, required farnesylation for its function. Two of these drugs (lonafarnib_SCH66336 from Schering–Plough, Kenilworth, NJ, and tipifarnib_R115777 from Johnson & Johnson, New Brunswick, NJ) have entered phase III trials and have been well tolerated, including in trials involving children (16). Lonafarnib competes with protein substrates for binding to the FTase enzyme (17). Similar to Ras, the Lamin A precursor is also farnesylated, with farnesylation serving as a required step to insert prelamin A into the nuclear membrane as well as to allow for the two downstream cleavage steps which complete the processing of Lamin A (18). For this patient, we are waiting for FTIs for clinical trials.

With the knowledge that the single C-to-T base change seen in nearly all cases of HGPS created a cryptic splice site and, thus, deleted the normal second endoproteolytic cleavage site in the Lamin A processing pathway, it was hypothesized that progerin was forced to retain its farnesyl group and, therefore, could not dissociate itself from the nuclear membrane. With other members of the nuclear lamina also potentially becoming trapped in complexes with the mislocalized progerin, a mechanistic connection between this permanently farnesylated state and the striking nuclear blebbing and disrupted nuclear architecture seen in HGPS cells was proposed, and the possibility of preventing or reversing this phenotype through FTIs was raised. Detailed study of HGPS and LMNA mutations may also advance our understanding of the process of aging. Why do LMNA mutant cells enter senescence earlier than normal cells? (19).

In summary, we found a new patient with typical Hutchinson–Gilford Progeria syndrome with mutation of the G608G in the LMNA gene. This is the first case diagnosed as Hutchinson–Gilford Progeria syndrome by gene study in Korea.

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