Letter to the Editor

Diagnostic Genetics



Ann Lab Med 2019;39:105-108 https://doi.org/10.3343/alm.2019.39.1.105 ISSN 2234-3806 elSSN 2234-3814

ANNALS OF LABORATORY MEDICINE

Compound Heterozygous Pathogenic Variants of the 15-Hydroxyprostaglandin Dehydrogenase Gene in a Patient With Hypertrophic Osteoarthropathy: First Case in Korea

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Dear Editor,

Hereditary or primary hypertrophic osteoarthropathy (PHO) is an autosomal recessive disorder characterized by excessive proliferation of skin and bone cells at the distal parts of the extremities, resulting in clubbing of fingers and toes, periostitis of long tubular bones, and arthritis [1]. Pathogenic variants in the 15-hydroxyprostaglandin dehydrogenase (*HPGD*) gene and solute carrier organic anion transporter family member 2A1 gene (*SLCO2A1*) have been identified in hypertrophic osteoarthropathy, primary, autosomal recessive 1 (PHOAR1; Online Mendelian Inheritance in Man [OMIM] #259100) and hypertrophic osteoarthropathy, primary, autosomal recessive 2 (PHOAR2; OMIM #614441), respectively [2, 3]. Some differences exist between the clinical phenotypes of PHOAR1 and PHOAR2: 1) pachydermia and cutis gyrate are more frequent and severe in PHOAR2 than in PHOAR1;

Received: February 6, 2018 Revision received: April 26, 2018 Accepted: August 27, 2018

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Co-corresponding author: Chang-Seok Ki, M.D. https://orcid.org/0000-0001-7679-8731 Green Cross Genome, 107 Ihyeon-ro, Giheung-gu, Yongin 16924, Korea Tel: +82-31-260-9600, Fax: +82-31-260-9087 E-mail: changski.md@gmail.com 2) a certain proportion of PHOAR2 exhibits gastrointestinal hemorrhage, which is not observed in PHOAR1; 3) higher urinary prostaglandin E2 (PGE2) levels are observed in PHOAR2 (approximately 10-fold increase) than in PHOAR1 (approximately 4-fold increase); and 4) onset is usually around puberty in PHOAR2, whereas onset is often around birth in PHOAR1 [4]. In Korea, several cases of PHO have been reported, all harboring *SLCO2A1* mutations [5, 6]. We report the first case in Korea of a patient with PHO carrying compound heterozygous pathogenic variants in the *HPGD* gene.

A 21-year-old military soldier was referred to the outpatient clinic of the Armed Forces Capital Hospital, Seongnam, Korea, with a suspected diagnosis of acromegaly in February 2011. He had abnormal thickening of both hands and feet, as well as digital clubbing, which is an unusual finding in patients with acro-

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megaly. The digital clubbing required him to visit a tertiary hospital when he was six years old, but the cause of clubbing was not found. He denied any medical history, including that of gastrointestinal hemorrhage, any affected family members, or a sudden increase in height. Upon physical examination, digital clubbing of the fingers (Fig. 1A) and abnormal thickening of the feet (Fig. 1B) were noted; however, forehead pachydermia was not evident. Laboratory test results were within the reference range (blood chemistry, complete blood cell count, erythrocyte sedimentation rate, and C-reactive protein) as were hormonal test results (total-triiodothyronine, free-thyroxine, thyroid-stimulating hormone, and insulin-like growth factor 1). The chest X-ray showed no abnormalities, whereas the hand X-ray showed soft tissue swelling of the fingers without apparent acroosteolysis (Fig. 1C). No other long bones showed abnormalities. Bone scintigraphy and sella magnetic resonance imaging did not reveal any remarkable findings. Secondary causes of hypertrophic osteoarthropathy were ruled out using echocardiography, pulmonary function test, ultrasonography of the abdomen, and esophagoduodenoscopy. Urinary PGE2 was 2,626 ng, approximately 4-fold

higher than the upper normal range (reference range: 400–620 ng/24-hour urine).

The patient was suspected to have PHOAR1 rather than PHOAR2, and we decided to perform sequence analysis of the HPGD gene. After obtaining informed consent, genomic DNA was extracted from peripheral blood leukocytes of the patient and both parents. All seven coding exons and their flanking intronic regions were amplified and sequenced using an ABI 3730xl genetic analyzer (Applied Biosystems, Foster City, CA, USA) and the Big-Dye Terminator v3.1 Cycle Sequencing Kit (Applied Biosystems) with primers designed by the authors (available upon request). HPGD sequences were analyzed using Sequencher software (Gene Codes Corp., Ann Arbor, MI, USA) and compared with the reference sequence (NM 000860.5). The patient was compound heterozygous for a 2-bp deletion (NM_000860.5:c.310_ 311delCT) in exon 3 and a 2-bp deletion (c.527_528delTG) in exon 6, and both parents were heterozygous carriers of each variant (Fig. 2).

The c.310_311delCT variant has been predicted to cause a truncated protein (p.Leu104Alafs*3) and was reported as a patho-



Fig. 1. Clinical and radiological findings of a 21-yearold male Korean patient: (A) digital clubbing, (B) abnormal thickening of the feet, and (C) hand X-ray showing soft tissue swelling of the fingers.

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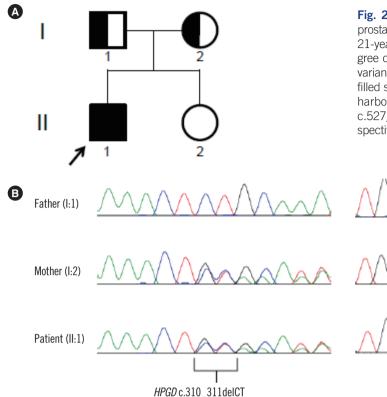
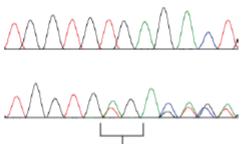


Fig. 2. Pedigree and sequence chromatogram of the 15-hydroxyprostaglandin dehydrogenase (*HPGD*) gene variants identified in a 21-year-old male Korean patient and his parents. (A) Family pedigree of the patient. The patient's younger sister does not have any variant. Male, square; female, circle; filled symbols, affected; halffilled symbols, heterozygous carriers; arrow, patient. (B) The patient harbors compound heterozygous variants, c.310_311delCT and c.527_528delTG, which are carried by his mother and father, respectively.



HPGD c.527_528delTG

genic variant in two pairs of Turkish siblings [7, 8], two Chinese siblings [9], and seven sporadic Chinese patients [9]. The c.527_ 528deITG variant is predicted to lead to a truncated protein (p.Val176Glufs*11) by Mutalyzer 2.0.28. It is absent from the Genome Aggregation Database (http://gnomad.broadinstitute. org) and the Korean Reference Genome database (http://152. 99.75.168/KRGDB/), but it can be classified as a pathogenic variant according to the guidelines of the American College of Medical Genetics and Genomics and the Association for Molecular Pathology [10].

Previously reported patients carrying c.310_311delCT homozygous variants presented common clinical PHOAR1 findings: digital clubbing, periostosis, acroosteolysis, joint swelling, and palmoplantar hyperkeratosis. Tüysüz *et al.* [7] described that the Turkish boys' father and grandfather also had mild digital clubbing and palmar hyperkeratosis. Carriers of *HPGD* gene mutations have been known to present some signs of PHO [1]. However, Erken *et al.* [8] showed that the Turkish siblings' relatives who were carriers did not exhibit signs of PHO. Neither parent of our patient showed any signs of PHO.

Although our patient had abnormal thickening of both hands and feet before school age, he neither experienced hand pain nor exhibited apparent acroosteolysis. Variable clinical phenotypes between PHOAR1 and PHOAR2, as well as within the same disease subtype, have not been completely elucidated; thus, further studies are necessary to elucidate the genotype-phenotype correlation.

Authors' Disclosures of Potential Conflicts of Interest

No potential conflicts of interest relevant to this article were reported.

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