

# Novel Fructose-1,6-bisphosphatase Gene Mutation in Two Siblings

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Fructose-1,6-bisphosphatase (FBPase) deficiency is an autosomal, recessively inherited disease that progresses with severe hypoglycemia, and metabolic attacks result in a defect in gluconeogenesis. If not appropriately treated, and if fructose is not excluded from the diet, the outcome could be fatal. Two Turkish children with FBPase deficiency were diagnosed based on mutation of the FBP1 gene. The first, a 2-year-old girl, was referred to our clinic because of lactic acidosis, uncorrectable hypoglycemia, and increased transaminases. FBPase deficiency was suspected in the patient, who recovered dramatically after a high-dose glucose infusion and adequate bicarbonate replacement. The second patient, a five-and-a-half-year-old male sibling of the patient, was also hospitalized, twice, because of hypoglycemic attacks and metabolic acidosis. Different from previous analyses, a homozygous c.658delT mutation was detected at exon 5 of the FBP1 gene in the two siblings. As a result of this mutation, there was a TGA (stop codon) at exon 6. There was first-degree consanguinity between the parents. These two cases were the first FBP1 gene mutations reported in our country.

## Introduction

**F**RUCTOSE-1,6-BISPHOSPHATASE (FBPase) is a key enzyme in gluconeogenesis, and if there is a deficiency, prominent hypoglycemia and metabolic acidosis can develop. FBPase converts fructose 1,6 bisphosphate to fructose 6-phosphate and inorganic phosphate. The enzyme is most active in the intestine and liver (Steinmann *et al.*, 2001). FBPase deficiency is caused by FBP1 gene mutation. FBPase coded in two distinct genes which are called FBP1, FBP2. In muscle FBPase is coding by FBP2 gene (Asberg *et al.*, 2010). When liver glycogen stores are depleted, the body uses gluconeogenetic precursors, such as lactate, glycerol, and pyruvate, to maintain blood glucose levels. During periods of low food intake or infection, an FBP1 defect can result in hypoglycemia, ketonuria, elevated blood lactate, and metabolic acidosis. The deficiency can be fatal in neonates, but tolerance to fasting generally improves with age. Relatively few cases of FBPase deficiency have been described, and it has been suggested that patients with deficient FBPase have been misdiagnosed with sudden infant death syndrome or Reye's syndrome (Emery *et al.*, 1988; Zammarchi *et al.*, 1995). If this autosomal, recessively inherited disease is not diagnosed and treated early, it could result in a fatal outcome (Baker and Winegrad, 1970; Baerlocher *et al.*, 1971). In treatment, the main principle is to decrease fructose intake and to abstain from long-term fasting. Herein we present the clinical and genetic results of the cases of two siblings diagnosed with FBPase deficiency that progressed with hypoglycemia and metabolic acidosis.

## Materials and Methods

### Patient reports

**Patient 1.** A 2-year-old girl was referred to the emergency unit because of recurrent vomiting and hypoglycemia attacks. There were no drugs or chronic disease in the medical history; however, the information showed that the patient had been hospitalized 1 year prior with the same signs and symptoms. In addition, there was first-degree consanguinity between the parents. The physical examination at admittance revealed a height of 89 cm (50–75 p) and weight of 11 kg (10–25 p), and dry mucosa, tachypnea, and hyperperic respiratory, accompanied by confusion and hepatomegaly. The laboratory examination results were as follows: glucose 14 mg/dL (70–100 mg/dL); uric acid 12.41 mg/dL (2–8 mg/dL); alanine transaminase (ALT) 196 U/L (5–40 U/L); aspartate transaminase (AST) 150 U/L (5–42 U/L); ketone positivity in urine (+++); prominent metabolic acidosis in the blood gases examination—pH 7.10 (7.35–7.45), pCO<sub>2</sub> 11.5 mmHg (35–45 mmHg), HCO<sub>3</sub> 5.8 mM (21–28 mM), BE –24.6 mM (–5 to +3 mM), and anion gap 28.8 mM (10–14 mM); and lactate 93.6 mg/dL (4.5–20 mg/dL). A glucose infusion was started at a rate of 10 mg/kg/min, and bicarbonate support was provided. The glucose infusion was increased, and insulin therapy was added due to low blood glucose levels. Approximately 15 h later, the hepatomegaly was recovered. The ALT level was corrected in 36 h. In addition, the blood gases parameters were recovered; the blood glucose was regulated; and the lactate level was decreased to

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TABLE 2. GENOTYPING OF FBPI GENE

Exon	Forward	Reverse	Product size (bp)
Promoter	5'-ACAGTGCCGGGTGGAGGGCAC-3'	5'CGCAGCGCTCGTGGTGAAC-3'	274
1	5'-CAGTGCCTACTGCCCTCTCTT-3'	5'-AGGCTCCCCAGGCAGGCAGACAGACA-3'	313
2	5'-CTACATGTTCTGGTGGTCATGG-3'	5'-CTGGGAAGAAGACCGGCTACAT-3'	270
3	5'-CTCCTTTAGTGTATCTTGCT-3'	5'-CTTCTGTCCCCAAACCAAGTG-3'	193
4	5'-TTGAGAATGCCTCCTGTTAAT-3'	5'-TCATTGCTCACAGACACCAG-3'	281
5	5'-ATCCAGGCCTGGGGACCCAG-3'	5'-CCCAGAACCTGCACCACCCTC-3'	238
6	5'-CACAGAACTTAGGAGACACC-3'	5'-ATCTGCTCCTCACTCCCTCTC-3'	236
7	5'-TTGGAAGTCCCACCAGCTCT-3'	5'GAATGTAAGGTGCACAGCAGG-3'	385

Primers and PCR product for FBPI gene exons and promoter regions (el-Maghrabi *et al.*, 1995).

sequencing results, FBPI gene c658T deletion in exon 5 was determined. Because of this deleted T nucleotide, there was a frame shift mutation; all of the codon had changed, and in exon 6 TGA (stop codon), the codon had disappeared, and FBPI protein could not be expressed effectively (Fig. 3).

### Discussion

Fructose-1,6-bisphosphatase (EC 3.1.3.11) is known as the regulatory enzyme that hydrolyzes fructose-1,6-bisphosphate to inorganic phosphate and fructose 6-phosphate. FBPI deficiency, which is related to hypoglycemia and metabolic acidosis, is encountered less frequently than hereditary fructose intolerance; the incidence of FBPI deficiency was reported as 1:350,000 in the Netherlands (Visser *et al.*, 2004). While approximately half of subjects with FBPI deficiency show signs during the newborn period, they are symptomatic during long-term fasting due to febrile diseases in more advanced ages. Patients experience generally normal health between attacks; however, they may show signs after high-dose fructose intake in this disease. The attacks decrease with age, and the majority of cases exhibit normal somatic and

psychomotor development (Baerlocher *et al.*, 1971). The disease was first defined in a 5-year-old child in 1970, and many patients were reported in the following years (Baker and Winegrad, 1970; Pagliara *et al.*, 1972). Kikawa *et al.* (1997) presented data of 13 children from 11 Japanese families, and reported that there was consanguinity in three families. In our cases, there was first-degree consanguinity between the parents. Contrary to hereditary fructose intolerance, gastrointestinal signs are few, and there is no restriction of fructose needed. In our two cases, there were no gastrointestinal signs, but they had severe acidosis and hypoglycemia. Both cases were life threatening, and the patients were treated under intensive care unit conditions. In cases of hypoglycemia and lactic acidosis, FBPI deficiency should be considered in the differential diagnosis, even though it is rarely encountered. In addition, fructose should be excluded from the diet and long-term fasting should be prevented.

The FBPI gene, which is thought to be responsible for the disease, was defined on chromosome 9q22.2-q22.3 in 1995 by el-Maghrabi *et al.* (1995). The gene consists of seven exons and spans over 31 kb (el-Maghrabi *et al.*, 1995). The FBPI gene has three transcripts (FBPI-201, FBPI-001 and FBPI-002

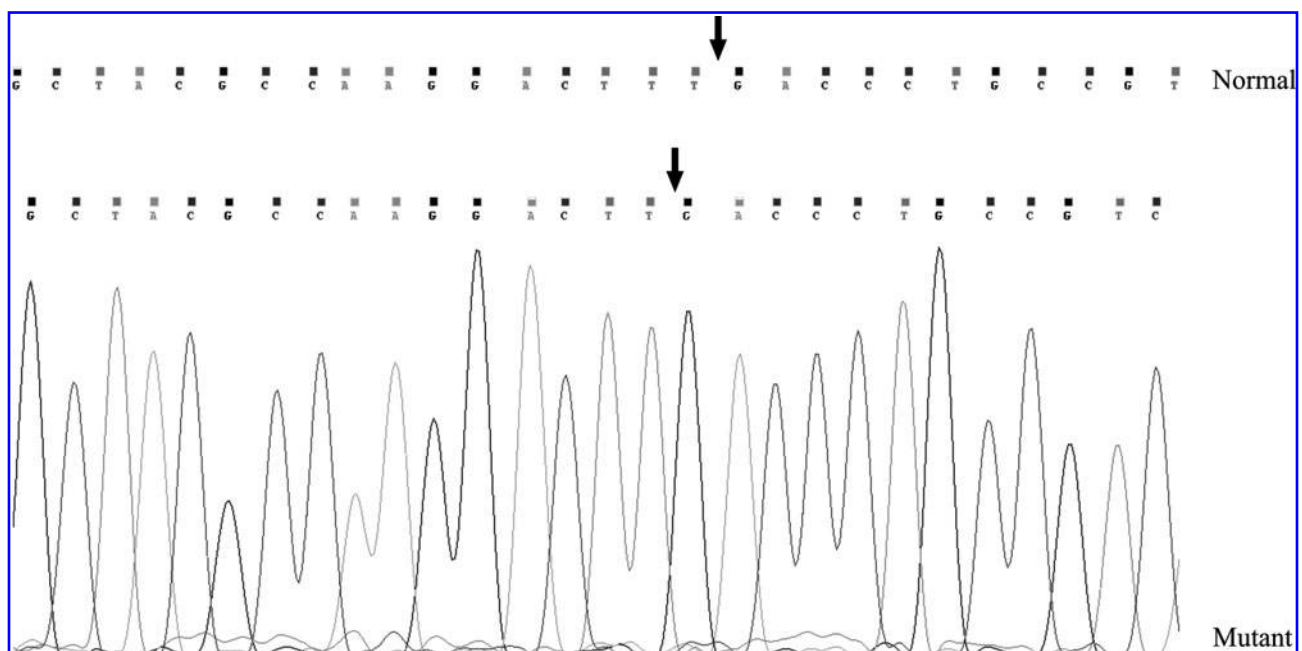


FIG. 2. Mutation analyses of patients' FBPI gene c.658 del T novel mutation. Arrows indicate deleted position.

FIG. 3. The GAC (D) codon T nucleotide was deleted, resulting in frame shift mutation. The 1-bp T deletion (bp position 658) results in a shift of the reading frame and a stop codon after 56 amino acids.

C658Tdel Mutation										
GAC (D)	TTG (L)	ACC (T)	CTG (L)	CCG (P)	TCA (S)					Exon5
CTG (L)	AGT (S)	ACA (T)	TCC (S)	AGA (A)	GGA (G)	AGA (A)				
(A) AGT (S)	TCC (S)	CCC (P)	CAG (Q)	ATA (I)	ATT (I)					Exon6
(I) CAG (Q)	CTC (L)	CTT (L)	ATG (M)	GGG (G)	CCC (P)					
GGT (G)	ATG (M)	TGG (W)	GCT (A)	CCA (P)	TGG (W)	TGG (W)				
(W) CTG (L)	ATG (M)	TTC (F)	ATC (I)	GCA (A)	CTC (L)					
TGG (W)	TCT (S)	ACG (T)	GAG (Z)	GGA (G)	TAT (Y)	TTC (F)				
(F) TGT (C)	ACC (T)	CCG (P)	CTA (L)	ACA (T)	AGA (R)					
AGA (R)	GCC (A)	CCA (P)	ATG (M)	GAA (Z)	AGC (S)					
TGA (STOP CODON)										

splice variants), and the c.658delT mutation is not encoded across a splice junction. We think that FBP1 protein is not expressed clearly in these patients because of the c.658delT mutation, which results in FBPase deficiency. Homozygous and heterozygote mutations were determined in the study performed in Japan, and a 1 bp insertion (960insG) on the FBP1 gene was defined as the one most commonly encountered (10 out of 16 mutant alleles). In another study performed in Japan, two mutations, which were heterozygote on the FBP1 gene, were detected in a girl (Matsuura *et al.*, 2002). Eleven different FBP1 mutations were also defined in that study. Herzog *et al.* (1999) detected a homozygous mutation in exon 7 of 960-961insG on the FBP1 gene; in addition, mutations of 35delA in exon 1, 778GA and 966delC in exon 7 were detected for the first time. According to the study performed on Japanese patients, the c.961 ins G mutation was defined as 46% (Kikawa Y *et al.*, 1997). Other homozygous and heterozygous mutations belonging to the FBP1 gene were also defined in different studies performed on other patients with FBPase deficiency (Faiyaz-Ul-Haque *et al.*, 2009; Moon *et al.*, 2011). As such, different mutations were defined in FBPase deficiency diseases in different populations (Table 3).

TABLE 3. MUTATIONS THAT HAVE BEEN FOUND IN FBP1 GENE IN DIFFERENT POPULATIONS

Mutation	Population	Reference
1-bp insertion (960insG) in exon 7	Japan	Kikawa <i>et al.</i> (1995)
490G>A transition in exon 4	Japan	Kikawa <i>et al.</i> (1997)
530C>A transversion in exon 4	Japan	Kikawa <i>et al.</i> (1997)
88G>T transversion in exon 1	Japan	Kikawa <i>et al.</i> (1997)
581T>C transition in exon 5	Japan	Matsuura <i>et al.</i> (2002)
six nucleotide repetitive insertion, c114_119dup CTGCAC	Saudi Arabia	Faiyaz-Ul-Haque <i>et al.</i> (2009)
c.841G>T		
c.778G>A c.881G>A G164S or 838delT in exon 7	Sweden South Korea	Asberg <i>et al.</i> (2010) Moon <i>et al.</i> (2011)

## Conclusion

Different from other studies, homozygous c.658delT mutation, which was not previously defined, was determined by FBP1 gene analysis of our patients with FBPase deficiency. Homozygous c.658delT mutation caused a stop codon in exon 6. Because of that, FBP1 gene expression is blocked at exon 6. Defective FBP1 protein is caused to FBPase deficiency in these patients. The present study is important because it is the first to report FBP1 gene novel mutation in our country.

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## Disclosure Statement

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