

DGAT1 mutation is linked to a congenital diarrheal disorder

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Congenital diarrheal disorders (CDDs) are a collection of rare, heterogeneous enteropathies with early onset and often severe outcomes. Here, we report a family of Ashkenazi Jewish descent, with 2 out of 3 children affected by CDD. Both affected children presented 3 days after birth with severe, intractable diarrhea. One child died from complications at age 17 months. The second child showed marked improvement, with resolution of most symptoms at 10 to 12 months of age. Using exome sequencing, we identified a rare splice site mutation in the DGAT1 gene and found that both affected children were homozygous carriers. Molecular analysis of the mutant allele indicated a total loss of function, with no detectable DGAT1 protein or activity produced. The precise cause of diarrhea is unknown, but we speculate that it relates to abnormal fat absorption and buildup of DGAT substrates in the intestinal mucosa. Our results identify DGAT1 loss-of-function mutations as a rare cause of CDDs. These findings prompt concern for DGAT1 inhibition in humans, which is being assessed for treating metabolic and other diseases.

Introduction

Congenital diarrheal disorders (CDDs) are rare and heterogeneous enteropathies, often with severe clinical manifestations (1, 2). Those whose cause has been identified typically result from autosomal recessive mutations. Affected genes include those related to disaccharidase deficiency, ion or nutrient transport defects, pancreatic insufficiency, or lipid trafficking (2). Some types of CDDs can be treated with dietary modification, but many present challenging clinical conditions, often requiring chronic nutritional support.

Here, we identified and characterized a rare DGAT1 mutation in a family with CDD. DGAT1 encodes 1 of 2 acyl CoA:diacylglycerol acyltransferases (DGATs), which catalyze the final step in triglyceride (TG) synthesis (3). DGAT1 is expressed ubiquitously, with highest expression in human intestine (4). Mice lacking DGAT1 have normal fat absorption, although absorption is delayed and more fat reaches distal intestinal regions (5). Because of the favorable metabolic phenotype of DGAT1-knockout mice (6), DGAT1 inhibitors have been developed (7–9) and proven efficacious in animal studies (8, 10). Several are being evaluated in clinical trials (11–13). However, mutations in human DGAT1 have not been reported, and information about human DGAT1 deficiency is limited.

Results and Discussion

Clinical summary. The affected family is a nonconsanguineous couple of Ashkenazi Jewish descent with 3 children from full-term, uncomplicated pregnancies. The first child, a boy, was unaffected. The second child (case 1), a girl, weighed 3.18 kg at birth and was fed with breast milk and cow’s milk formula. Three days after birth, she developed vomiting, colicky pain, and nonbloody, watery diarrhea, 8–10 times daily. She was treated with oral rehydration solution, and her formula was changed to soy-based formula, but diarrhea continued. Cultures for bacterial pathogens, rotavirus, and adenovirus were negative. She exhibited protein-losing enteropathy, with stool α1 antitrypsin of 8 to 20 mg/g (normal, <3 mg/g stool) and hypoalbuminemia. She required total parenteral nutrition and intermittent infusions of albumin. Stomach, duodenum, and colon biopsies were negative for chronic granulomatous disease, autoimmune enteropathy, food protein-induced enterocolitis, microvillus inclusion disease, and tufting enteropathy (Supplemental Figure 1; supplemental material available online with this article; doi:10.1172/JCI64873DS1). Neuroendocrine cells were present in intestinal biopsies. Congenital lymphangiectasia, a cause of protein-losing enteropathy, was excluded by CT scan and histology. There was evidence of dystrophic microvilli in the duodenum. Immunological tests were unremarkable except for slightly decreased IgG (275 mg/dl) with normal subclasses.

The child exhibited hyperlipidemia, with a fasting serum TG level of 325 mg/dl at 1 month of age (subsequently, 81–631 mg/dl; mean, 264 mg/dl [n = 55]) (Table 1). The father had elevated fasting TG (118–481 mg/dl) and total cholesterol levels (140–260 mg/dl), with a HDL cholesterol level of 33 to 39 mg/dl. The mother also had elevated fasting TG (144–229 mg/dl) and total cholesterol levels (174–220 mg/dl), with a HDL cholesterol level of 39 to 42 mg/dl.

At 14 months of age, the child was below the first percentile for weight, despite parenteral nutrition and feeding per gastrosomy tube. She had recurrent episodes of sepsis, presumably

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related to a venous catheter. Eventually, she tolerated tube feedings (80 kcal/kg/d) with amino acid–based formula containing maltodextrin and medium chain TGs but did not gain weight and continued to lose protein in her stool. She died at 17 months of age from complications of malnutrition and sepsis.

The third child (case 2), a boy, weighed 3.7 kg at birth. He initially tolerated breast milk and soy formula but developed diarrhea 3 days after birth. His stools were nonbloody, 4–6 times daily, and watery. Six days after birth, he was admitted for dehydration, metabolic acidosis, and hyponatremia. Stool cultures were negative for bacteria pathogens, rotavirus, and adenovirus. Stool sodium and potassium levels were normal. Serum IgG was normal, but fecal IgA was increased to 4.7 to 7.9 mg/g. He required intravenous and predicted to be deleterious by PolyPhen2 (16). A single loss of function (nonsense, splice, frameshift) or were nonsynonymous and predicted to be deleterious by PolyPhen2 (16). A single candidate gene (DGAT1) was identified. The child was homozygous (and both parents heterozygous) for a splice variant (chromosome 8, 145541756 A → G) in the splice donor site 3’ of exon 8, altering the invariant GT to GC. We identified 3 individuals in approximately 12,500 control exomes as carriers for this mutation and estimate probability of homozygosity of 1 in approximately 50 to 100 million births, revealing a novel and severe recessive disorder.

Using PCR and RFLP, we confirmed both affected children as homozygous carriers of the mutation. The maternal grandmother, paternal grandfather, both parents, and the unaffected child were heterozygous (Figure 1B). The known founder effect likely explains why two unrelated families from the Ashkenazi Jewish subpopulation carry such a rare mutation (17).

The predicted result of the mutation is skipping of exon 8 (18), yielding an in-frame deletion of 75 bp (Figure 1C). Using RT-PCR with primers flanking exon 8 (Figure 1D), we confirmed cDNA products corresponding to the wild-type and exon

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<th>Table 1 Blood lipids</th>
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Case 2

<table>
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<tr>
<th>Age (mo)</th>
<th>TG (mg/dl)</th>
<th>Total cholesterol (mg/dl)</th>
<th>LDL-C (mg/dl)</th>
<th>HDL-C (mg/dl)</th>
<th>VLDL-C (mg/dl)</th>
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*Cholestyramine initiated. LDL-C, LDL cholesterol; HDL-C, HDL cholesterol; VLDL-C, VLDL cholesterol.
in that they occur distal to TG synthesis, whereas, with DGAT1 deficiency, the defect is in TG synthesis itself.

The phenotype of DGAT1 deficiency in humans differs from that of DGAT1-KO mice, which exhibit reduced body fat, increased energy expenditure, improved glucose tolerance, resistance to diet-induced obesity, and extended longevity (6, 22). The mice also have delayed fat absorption and decreased postprandial excursions of plasma TG but not excess fecal fat or diarrhea (5). These findings suggest species-specific differences for DGAT1 deficiency. In mammals, DGAT1 and DGAT2 account for nearly all TG synthesis (20). Using quantitative PCR, we found that both Dgat1 and Dgat2 are expressed in murine intestine, but only DGAT1 is expressed highly in human intestine (Figure 2E), as we reported (6, 23). Thus, human intestine may be more sensitive to DGAT1 inhibition, owing to lack of DGAT2 expression.

How human DGAT1 deficiency causes diarrhea is unclear, but buildup of DGAT1 lipid substrates in the intestinal mucosa or lumen likely contributes. Excess diacylglycerols or fatty acids could become toxic by acting as bioactive signaling lipids or by detergent-like behavior of fatty acid moieties. Toxicity to enterocytes could result in protein-losing enteropathy. Alternatively, bile acid malabsorption can cause diarrhea, and DGAT1 deficiency could affect bile acid metabolism. Unfortunately, no information on fecal bile acid levels in the affected individuals was available.

The reasons for the different clinical outcomes of the affected children remains unclear. Case 2 suggests that there may be aging-associated increases in intestinal DGAT2 expression or other adaptation to DGAT1 deficiency. Without complications from sepsis, the case 1 individual might also have improved. The two children were different sexes, but relative levels of DGAT2 expression in male and female children are unknown. Treatment with a bile acid–binding resin might have helped in case 2. However, timing of this treatment did not correspond with clinical improvement.

Moderate hyperlipidemia, including hypertriglyceridemia, occurred in the affected children and parents. Whether this is causally linked to the DGAT1 mutation or an independent trait is unclear. Mice lacking DGAT1 have normal fasting levels of serum TGs (6) and reduced TGs after fat intake (5), thus hypertriglyceridemia linked to DGAT1 deficiency would be specific to humans. If hypertriglyceridemia is associated, overcompensation from hepatic DGAT2 may contribute via lipoprotein-mediated secretion of TGs. Alternatively, interruption of bile acid absorption in the distal small intestine is associated with VLDL overproduction and hypertriglyceridemia (24). The identification of more individuals with homozygous DGAT1 deficiency could shed light on this issue.

Notably, recent studies of DGAT1 inhibitors in humans reported dose-related adverse effects of mild-to-moderate diarrhea in some patients (25). Our findings suggest this effect results from DGAT1 inhibition, rather than being off target. A better understanding of DGAT1’s role in the intestine and how DGAT1 deficiency causes diarrhea will be useful for developing or improving therapies with DGAT1 inhibitors.
Methods

Experimental procedures are described in detail in Supplemental Methods.

Genetic screening and verification. Exome capture was performed with the Agilent Whole Exome SureSelect v2 Kit. Single-nucleotide-variant search parameters are described in the text (see Mutation identification and characterization). For RFLP analysis, PCR products were digested with Fnu4-HI. Bands are 165, 32, and 3 bp (wild type) and 122, 43, 32, and 3 bp (mutant).

Molecular characterization. MEFs were isolated and immortalized (26). DGAT activity was assayed (4) with conditions specific for DGAT1 (50 mM MgCl2). Total RNA was isolated with TRIzol (Invitrogen) and reverse-transcribed with the iScript cDNA Synthesis Kit (Bio-Rad). Antibodies were FLAG antibody M2 (1:2,000, Sigma-Aldrich), DGAT1 antibody NB110-41487 (1:1,000, Novus), and HSP90 (1:2,000, BD Biosciences). HRP secondary antibodies (1:5,000, Amersham) were used with ECL (Pierce).

Statistics. The probability of Δ8 mutation causing CDD was empirically determined from the frequency of any homozygous or compound heterozygous, deleterious mutations occurring in the reference exome sets (14, 15).

Study approval. Written informed consent was obtained from all adult participants and from the parents of the children. These genetic studies were approved by the Partners Institutional Review Board, Boston, Massachusetts, USA.

Acknowledgments

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Figure 2

Δ8 mutant allele yields an unstable protein, resulting in loss of DGAT1 activity. DGAT1 cDNAs were expressed in DGAT1-KO or DGAT1,2-KO MEFs. (A) DGAT1 activity (accumulation of [14C]-TG) was absent in lysates from Δ8-expressing cells, despite the presence of mRNA. All lysates synthesized [14C]-diacylglycerol ([14C]-DAG) (internal control). (B) Intact Δ8-expressing cells do not accumulate TG, as measured by thin-layer chromatography in DGAT1,2-KO cell lines after incubation with oleic acid (200 mM). Free cholesterol (FC) was used as a loading control. (C) Immunoblot showing that proteasome inhibition (MG132) rescues Δ8 protein expression in DGAT1,2-KO MEFs. (D) MG132-treated DGAT1,2-KO MEFs expressing Δ8 do not accumulate [14C]-TG after a 2-hour treatment with 200 mM [14C]-oleic acid. (E) qPCR showing that humans lack DGAT2 expression in the small intestine. Act., activity; Duo., duodenum; Jeju., jejunum; Ile., ileum.
brief report