Case report

# A New Variant of the IER3IP1 Gene: The First Case of Microcephaly, Epilepsy, and Diabetes Syndrome 1 from Turkey

# Söbü E et al. Microcephaly, Epilepsy, and Diabetes Syndrome 1

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#### What is already known on this topic?

MEDS1 manifests as microcephaly with simplified gyral pattern in combination with severe infantile epileptic encephalopathy and early-onset permanent diabetes.

#### What this study adds?

This is the first case reported from Turkey, differs from other cases due to the absence of a typical simplified gyral pattern on early brain MRI, the late onset of diabetes, and the presence of a new genetic variant.

#### Abstract

Microcephaly, Epilepsy, and Diabetes Syndrome 1 (MEDS1) is a rare autosomal recessive disorder and caused by defects in the IER3IP1 (Immediate Early Response 3 Interacting Protein 1) gene. Only 9 cases have been described in the literature. MEDS1 manifests as microcephaly with simplified gyral pattern in combination with severe infantile epileptic encephalopathy and early-onset permanent diabetes. A simplified gyral pattern has been described in all cases reported to the date. Diagnosis is made by de nonstration of specific mutations in the IER3IP1 gene. In this study, we present an additional case of a patient with MEDS1 who is homozygous for the c.53C >T p.(Ala18Val) variant. The case, the first to be reported from Turkey, differs from other cases due to the absence of a typical simplified gyral pattern on early brain MRI, the late onset of diabetes, and the presence of a rew genetic variant. The triad of microcephaly, generalized seizures and permanent neonatal diabetes should prompt screening for mutations in IER3IP1.

Keywords: Developmental delay, diabetes mellitus, epilepsy, IER3IP1 MEDS1

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## Introduction

The term monogenic diabetes (infantile-onset diabetes) refers to diabetes associated with monogenic defect and diagnosed usually in the first six months of life (1). Recent developments in the field of molecular genetics indicate that diabetes occurring very early in life is mostly caused by underlying monogenic defects (2,3). In fact, current studies indicate that monogenic diabetes should be considered in cases of diabetes diagnosed in the first two years of life, and these studies report that approximately 1%–6% of pediatric diabetes cases are actually neonatal diabetes (2,3.4). Microcephaly, epilepsy, and diabetes syndrome 1 (MEDS1) is an autosomal recessive neurodevelopmental disorder that was first described by Poulton et al. in 2011 and is characterized by microcephaly, simplified gyral pattern, severe epilepsy, and infantile diabetes (5). The disease is known to result from homozygous and compound heterozygous mutations in the *IER3IP1* gene, and it has so far been reported in a total of nine cases (5,6,7,8,9).

In this study, we present an additional case of a patient with MEDS1 who is homozygous for the c.53C >T p.(Ala18Val) variant. The case, the first to be reported from Turkey, differs from other cases due to the absence of a typical simplified gyral pattern on early brain MRI, the late onset of diabetes, and the presence of a new genetic variant. The triad of microcephaly, generalized seizures and permanent neonatal diabetes should prompt screening for mutations in *IER3IP1*.

# Case Report

The patient was born of the first live birth from the first pregnancy of a healthy father and mother, who were first cousins. He had no siblings, was born at term, and had no complications during pregnancy or the perinatal period. He had a birth weight of 3.400 kg (-0.24 SD) with a height of 49 cm (-0.37 SD) and a head circumference of 34 cm (-0.42 SD). At the age of 11 weeks, he presented to the pediatric neurology outpatient clinic with complaints of spasms, crying, and restlessness. Neurological examination performed at presentation revealed that he had no eye contact, object tracking, and head control and that he had central hypotonia and flexor spasms. Physical examination at presentation revealed that he weighed 4.700 kg (-0.83 SD), measured 55 cm (-1.31 SD), and had a head circumference of 35.5 cm (-2.46 SD) and a conspicuous microcephaly. He had hypertelorism, depressed nasal bridge and micrognathia (Figure 1). Laboratory tests revealed blood amino acid, cerebrospinal fluid (CSF) amino acid, acyl-carnitine, and urine organic acid levels within normal limits. Electroencephalogram (EEG) showed a burst suppression pattern. Brain MRI performed at the age of 2 months revealed normal cortical sulci and gyri with normal widths for age as well as normal ventricular system with normal width for age (Figure 2). Diagnosed with epilepsy, he was started on vigabatrin and pyridoxal 5 phosphate therapy. Owing to first-degree consanguineous marriage in the family and the co-occurrence of epilepsy and dysmorphic features, whole-exome sequencing (WES) was performed on DNA obtained from the proband and parents. He was found to have a homozygous missense variation (c.53C>T/p.Ala18Val) in *IER3IP1*. We could not find any phenotype-genotype study about this mutation in the literature. Homozygous pathogenic variants of this gene have been associated with autosomal recessive microcephaly, epilepsy, and diabetes syndrome type 1 (OMIM: 614231), and *IER3IP1* is a highly conserved protein with marked

expression in the cerebral cortex and in beta cells. As the patient's seizures did not respond to the initial treatment, other drugs were introduced in the following order: levatiracetam, topiramate, clonazepam, and clobazam. Brain Computed tomography (CT) performed at the age of 9 months revealed increased distance in the CSF space, particularly remarkable in the frontal lobe (Figure 3).

While there were no clinical signs of hypogonadism that might occur with MEDS1, thyroid function tests revealed thyroid stimulating hormone (TSH) level of 0.89 III/ml (normal range: 0.27–4.2) and free T4 level of 0.96 ng/ml (normal range: 0.87–1.76). At the age of 18

hormone (TSH) level of 0.89 IU/ml (normal range: 0.27–4.2) and free T4 level of 0.96 ng/ml (normal range: 0.87–1.76). At the age of 18 months, he presented with rapid breathing, and further tests revealed a blood glucose level of 311 mg/dl, insulin level of 4.9 mIU/L, C-peptide level of 0.43 µg/L, and HbA1c level of 7.09%; upon establishing a diagnosis of diabetes, insulin therapy was started at a dose of 0.4 U/kg. Diabetes antibodies (anti-glutamic acid decaboxylase, anti-insulin, and islet antibodies) were negative. There was no acidosis or ketonuria. Medical history was negative for polydipsia, polyuria, and significant weight loss. This variant has not been previously reported. Both parents were found to be heterozygous for the mutation.

The patient's parents provided informed consent for publication of this case report.

# Preparation for Genetic Analysis

Genomic DNA extraction was performed according to manufacturer's instruction (Maxwell RSC Blood DNA kit, Promega, USA) using Maxwell RSC Instrument (Promega, USA). 30 µl of Proteinase K (PK) Solution was added into 200 µl blood sample. 300 µl of Lysis Buffer was added to blood and PK mix and incubate at 56°C for 20 minutes. After this step, each blood lysate sample was transferred to the cartridges. At the end of assay in the instrument, 50 µl DNA was eluted. The concentration of DNA was determined spectrophotometrically by measurement of the absorbance at 260/280 nm using a Nanodrop 1000 apparatus (Thermo Fisher Scientific). The concentration of DNA samples for libraries were determined by using Qubit 3.0 (Thermo Fisher Scientific). The sequencing libraries for exome sequencing were prepared according to Twist Human Core Exome Kit protocol (Twist Bioscience, USA). Paired-end 150 bp read sequencing was performed on a NovaSeq system (Illumina, USA).Ana

#### Results

Raw data were uploaded to the Sophia DDM (Sophia Genetics-Lausanne-Switzerland) platform and for further analysis, which detected homozygous c.53C>T p.(Ala18Val) (NM\_016097) variation in IER3IP1. This detected variant could not be found in any literature report or in the healthy population database (gnomAD; https://gnomad.broadinstitute.org/). However, in silico prediction databases (MutationTaster, PROVEAN, SIFT) stated, in consensus, that the variation was "deleterious." The American College of Medical Genetics (ACMG) 2015 criteria qualified the variant as "Class 3 - variant of uncertain clinical significance" (10). The segregation analysis for the variation was performed with Sanger sequencing and Integrative Genomics Viewer (IGV) using samples collected from the patient's parents, and both parents were found to be heterozygous carriers of the mutation (Figure 4, Figure 5). Because the patient's clinical findings were similar to the expected symptoms of the "Microcephaly, epilepsy, and diabetes syndrome 1 (OMM: 614231)" phenotype caused by homozygous pathogenic variants in IER3IP1, this mutation was thought to account for the patient's phenotypic features.

## Discussion

This case report presented a male patient with homozygous variation in *IER31P1*, this is the first case reported from Turkey and the 10th case in the literature. It differs from other cases due to absence of a typical simplified gyral pattern on brain MRI and later onset of diabetes compared with other reported cases.

MEDS1 syndrome was first reported by Poulton et al. in 2011 in two cases from two unrelated families. Common findings in these cases were microcephaly with simplified gyral pattern in combination with severe infantile epileptic encephalopathy and early-onset permanent diabetes. An autopsy specimen from one patient showed increased apoptosis in the cerebral cortex and pancreas beta cells, implicating premature cell death as the pathogenetic mechanism (5).

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Microcephaly, epilepsy, and diabetes syndrome type L (MED S1) (OMIM: 614231), which shows an autosomal recessive pattern of inheritance, results from a defect in the production of immediate early response 3 interacting protein 1 (IER3IP1) expressed in beta cells of the cerebral cortex and pancreas. IER3IP1 is localized to the endoplasmic reticulum (ER) and is thought to play a role in the transport of proteins between the ER and Golg apparatus and to be involved in the ER stress response (5). The association of neonatal diabetes with IER3IP1 mutations suggests that IER3IP1 regulates β-cell survival and/or function. Increased apoptosis in the cerebral cortex and pancreatic beta cells in autopsy samples with IER3IP1 mutation points at early apoptosis as the pathogenic mechanism (5,7).

Neonatal diabetes refers to diabetes that is associated with a monogenic defect and is usually diagnosed in the first 6 months of life. The age at diagnosis of diabetes in the reported MEDS1 cases ranges from 14 days to 2 months (Table 1). In our case, however, diabetes emerged at age 18 months of age, later than in other reported cases (6,7,9). Although current studies have shown that monogenic diabetes usually occurs in the first 6 months, event studies have shown that it can rarely occur at the age of 12 or even 24 months (3,11,12). The important characteristics of the cases reported in the literature are summarized in Table 1.

The detection of c.62 T > G and c.233 T > C variants in all but one of the cases reported to date, and the fact that most of the cases are in Middle Eastern and North African countries or in countries receiving immigration from these regions, indicate that these variants are probably not mutational hotspots, but rather are rare ancestral variants unique to these regions. In the case of Shalev et al, the common c.62 T > G missense variant and the novel c.79delT frameshift variant were compound heterozygous, and although this novel variant was a frameshift variant, the patient was more mildly affected than previously reported ones and survived to 8 years of age (8). This shows that variants other than two common mutations can cause different phenotypes. The c.62 T > G (p.Val21Gly) variant affects the first transmembrane hydrophobic domain of the protein, and the c.233T>C (p.Leu78Pro) variant affects the second transmembrane hydrophobic domain of the protein's expression and/or function. The variant in our patient affects amino acid at position 18 in the first transmembrane hydrophobic domain, probably its mechanism of action is similar to c.62T>G, which affects amino acid at position 21. In addition, the c.62T>G variant is adjacent to the protein cleavage site. The milder phenotype of our case may be due to the fact that our variant is not so close to this cleavage site (5). Another piece of evidence supporting the pathogenicity of the variant in our patient is that the residues affected by both our patient's variant (18th residue) and the c.62T>G (21st residue) variant are located within a highly conserved 12-residue region among the species as shown below.

11 18 21
AALLCVNAIAVLHEERFLKNIG
AALLCVNAIAVLHEERFLKNIG
TAILFTNAIAVLHEERFLSKIG
AALLCVNAIAVLHEERFLKNIG

human mouse zebrafish cow In differential diagnosis, Wolcott-Rallison syndrome has been reported to be the most common cause of neonatal diabetes in families with consanguine marriages (13). This syndrome, which results from a homozygous mutation in EIF2AK3, is characterized by insulin-dependent diabetes mellitus before 6 months of age, skeletal dysplasia after 6 months of age, and liver failure. This syndrome manifests as renal failure, microcephaly, epilepsy, and central hypothyroidism and it must be ruled out in the differential diagnosis of MEDS1 (14). Increased ER stress, and thus beta-cell death, constitutes the pathogenesis of the disease, and management requires insulin replacement (15). This syndrome may also include episodes of liver failure and skeletal anomaly in later ages, indicating the importance of early genetic diagnosis. A study that presented 4 cases of MEDS1 reported that three of the patients had skeletal findings, including osteoporosis, metaphyseal changes, osteopenia, pathological fractures, and poor modeling of long bones (6). Our patient had no skeletal anomaly. IER3IP1 has an unclear role in the development of the cortex and in the pathogenesis of epilepsy and diabetes, but it is thought to be required

during early stages of neural development, for instance, during neural progenitor proliferation. Presence of microcephaly with simplified gyration has a distinctive role in differential diagnosis and already exists during gestation. Severe infantile epileptic encephalopathy is highly unusual in primary microcephaly and has been reported only in patients with WDR62 mutations (16). In addition, a rare combination of primary microcephaly and severe infantile epilepsy in patients with PNKP mutations has been reported (17).

Because our patient had refractory epilepsy, microcephaly, and axial hypotonia at the time of presentation, no specific complication at birth and was born of parents who were first cousins, he underwent WES early and was diagnosed with MEDS1. The patient's family was asked to be vigilant about symptoms of potential diabetes with blood glucose being monitored. Diabetes emerged later than reported in other cases in the literature, but early genetic analysis allowed for diagnosing diabetes before acidosis developed. The patient did not present with microcephaly at birth, but with increasing age, it became evident. It was remarkable that the simplified gyral pattern, which was detected all other cases, was absent on early MRI. Continued apoptosis in the postnatal period was thought to be the cause of magnetic resonance imaging (MRI) findings and microcephaly. This hypothesis was supported by increased distance in the CSF space detected on brain CT performed at the age of 9 months.

In conclusion, this is the first case of MEDS1 reported from Turkey and shows a variant that has not been previously described in the literature. Although the simplified gyral pattern, which co-occurs with the triad of microcephaly, epilepsy, and diabetes, may guide the diagnosis of MEDS1, manifestation of the symptoms may sometimes take time. Early genetic counseling should be considered in families where consanguine marriage is accompanied by epilepsy and microcephaly.

## Acknowledgements

None

## **Ethics Informed Consent**

The patient's parents provided informed consent for publication of this case report.

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**Table 1.** The important characteristics of the cases reported in the literature.

iporum cha	Poulton et al (2011)  Case 1   Case 2		reported in the literature.  Abdel-Salem et al. (2012)				Shalev	Valenzu	Rjiba	Curre
			Family 1251		Family 1578		et al(2015 ) Case 7	ela et al (2017)	et al (2021)	nt case
			Case 3 Case 4		Case 5   Case 6					
										10
Gender Consan	M +	<u>M</u>	M +	F +	F +	F +	M -	M -	M	M
guinity	'	'	'	'	'	'	_			'
MRI	simplif ied gyral pattern	simplifie d gyral pattern	simplified gyratio n, cortical atroph y, hypopl astic corpus callosu m, cerebel lar vermis hypopl asia	simplified gyral pattern and agenesi s of corpus callosu m with mild cerebel lar vermis hypopl asia without atrophy	cerebra l atroph ywith simplif ied gyral pattern , and agenesi s of the corpus callosu m	simplified gyral pattern and agenesi s of corpus callosu m with normal cerebel ium	simplifi ed gyral pattern	simplifie d gyral pattern, ventricul omegaly and hypoplast ic corpus callosum	atrophy of the supra- tentorial level without pituitary anomal y	Norma I
EEG	High voltage asymme tric multifoc al activity with abnorma l backero und	hypsarr ythmia	polyspi kes and slow waves with burst suppre ssion	general ized epilepti c abnorm alities with sharp and slow- waves	burst suppre ssion pattern	burst suppre ssion pattern	hypsarr ythmia	Low- amplitud e backgrou nd in the theta- delta range, suggestin g diffuse neuronal dysfuncti on, with no epileptifo rm discharge s.	NA	burst suppre ssion pattern
Genetic analysis	c.62 T > G p.Val21 Gly Homozy gous	c.233 T > C p.Leu7 8Pro Homoz ygous	c.233 T > C p.Leu7 8Pro Homoz ygous	c.233 T > C p.Leu7 8Pro Homoz ygous	c.233 T > C p.Leu7 8Pro Homoz ygous	c.233 T > C p.Leu7 8Pro Homoz ygous	c.62 T > G/ p.Val21 Gly and c.79del T/p. Phe27fs Ser*25 Compo und heteroz ygous	c.233 T > C C p.Leu78P ro Homozy gous	c.62 T > G p.Val21 Gly Homoz ygous	c.53C> T p.(Ala 18Val) Homoz ygous
Diabete	+	+	+	+	+	+	+	+	+	+
Age at diabetes onset	NA	NA	NA	NA	40 days	14 days	NA	5 Weeks	2 months	18 months
Hypogo	+	NA	bilatera	NA	NA	NA	retractil	NA	unilater	-

nadism			l undesc ended testes				e testes		al cryptorc hidism and small genitali a	
Skeletal findings	NA	NA	osteop orosis, metaph yseal change s	patholo gical fractur e	poor modeli ng of long bones and osteop enia	NA	NA	-	NA	-
Death	18 months	27 months	5 1/2 years	26 months	31/2 years	-	8 years	7 weeks	1 year	-

Table 1. General characteristics of MEDS patients
MRI: magnetic resonance imaging, EEG: electroencephalogram, NA: not available, M: Male, F:Female.
Figures
Figure 1. Patient's facial appearance.



Figure 2. Brain MR images of the patient at the age of three months old.

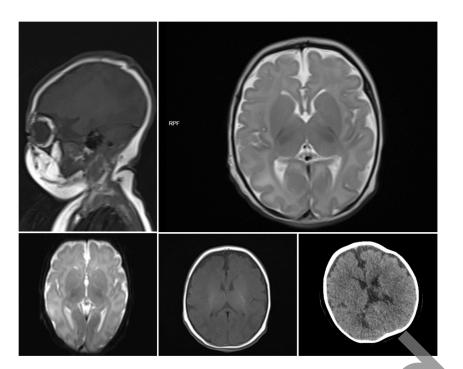
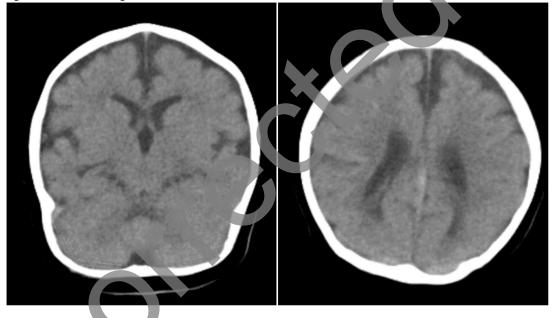


Figure 3. Brain CT at the age of 9 months old.



 $Figure\ 4. Results\ of\ Next-generation\ sequencing\ at\ the\ mutation\ locus\ c.53C>T\ p. (Ala18Val).$ 

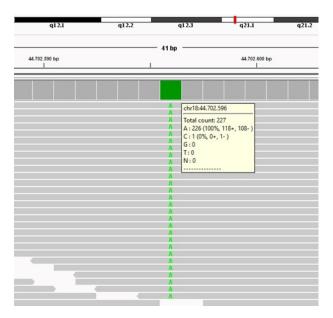


Figure 5. Integrative Genomics Viewer were used to analyse the characteristics of the mutated MEDS1 protein.

