



RESEARCH ARTICLE

Ehlers-Danlos syndrome kyphoscoliotic type 2 caused by mutations in the FKBP14 gene: an analysis of five cases

[version 1; peer review: 2 approved]

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Abstract

Background. This study deals with a rare (orphan) monogenic connective tissue disorder - Ehlers-Danlos syndrome kyphoscoliotic type 2 (EDSKS2). Kyphoscoliotic type 2 Ehlers-Danlos syndrome is an autosomal recessive disorder caused by mutations in the FKBP14 gene (7p14.3), which encodes the FKBP22 protein. According to the 2017 classification, this type is in group seven - collagen spatial structure and cross-linking defects. We present results of clinical examination and molecular genetic analysis for five patients with age varying from two to fifteen years.

Methods. Five patients were examined using clinical and laboratory methods. DNA samples used for the analysis were extracted from whole blood samples using a Wizard® Genomic DNA Purification Kit (Promega, USA) according to the manufacturer's protocol.

Results. The major clinical findings were kyphoscoliosis, early motor development delay, muscular weakness, hypotonia and hearing loss. Molecular genetic analysis detected a homozygous c.362dupC duplication in exon 3 of the FKBP14 gene in all five patients. This mutation is common in various countries. Differential diagnostics were carried out to exclude other Ehlers-Danlos syndrome types and myopathies.

Conclusions. Literature analysis and examination of

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1

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report



report

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five EDSKS2 patients demonstrated the involvement of major organs and systems, such as joints, spine, muscles, cardiovascular system, respiratory system, hearing, and vision, into the pathological process. Kidney mobility increases and nephroptosis seems to be secondary caused by muscular weakness. During molecular genetic analysis, to verify EDSKS2 it is recommended to initially search for the c.362dupC duplication, which appears to be common in European countries, including Russia.

Keywords

children, rare (orphan) disorders, monogenic connective tissue disorders, clinical findings, FKBP14 gene, c.362dupC duplication

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The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

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Introduction

Ehlers-Danlos syndrome (EDS) is commonly encountered by various medical specialists. It is mainly characterized by skin hyperelasticity, joint hypermobility, easy bruising, and hypertrophic scarring. The disorder is genetically heterogeneous: according to the 2017 classification, there are 13 clinical genetic types,¹ divided into seven groups depending on the pathogenesis. The kyphoscoliotic type belongs to the B group – disorders of collagen folding and collagen cross-linking – or type VI in the 1997 classification. Patients with kyphoscoliotic type come to the attention of pediatricians, neurologists, and orthopaedists.

EDS kyphoscoliotic type is an autosomal recessive disorder with estimated prevalence of one per 100,000 newborns.² Its main symptoms (major clinical diagnostics criteria) are severe muscle hypotonia at birth (“floppiness”), early-onset kyphoscoliosis (usually progressive), hypermobility of joints, dislocations/subluxations (especially of knee joints).¹ Additional (minor) diagnostic criteria include skin hyperelasticity, easy bruising, arterial ruptures/aneurisms, osteopenia/osteoporosis, bluish sclerae, umbilical/inguinal hernia, thorax deformations, marfanoid habitus, equinovarus feet deformities, myopia.

EDS kyphoscoliotic type is currently subdivided into two subtypes. EDS kyphoscoliotic type 1 (EDSKS1) is caused by mutations in the *PLOD1* gene (1p36.22), which encodes lysyl hydroxylase, a catalyst essential for the stability of the intermolecular collagen crosslinks.^{3,4} EDS kyphoscoliotic type 2 (EDSKS2) is caused by mutations in the *FKBP14* gene (7p14.3), which encodes the FKBP22 protein. FKBP22 is a highly conservative peptidyl-prolyl *cis-trans* isomerase (PPIase), which catalyzes collagen folding and acts as a chaperone to collagen types III, VI, X.⁵ Thus, both genes are tied to collagen spatial structure formation.

These two subtypes are clinically similar, but slightly different. Mutations in the *PLOD1* gene lead to moderate scarring and bruisability, microcornea, scleral and ocular ruptures, and facial abnormalities (low set ears, epicanthal folds, downslanting eyes, synophrys, high-arched palate). EDSKS2 is less explored because of its recent discovery and lower frequency. Some patients show signs of hearing loss (sensorineural, conductive or mixed) that were not present at birth, as well as follicular hyperkeratosis, muscular atrophy, and bladder diverticulum.

Because this EDS type is less frequent and not fully explored, in this study we present the clinical data of five patients with EDSKS2.

Methods

Clinical examination

Five patients were examined using clinical and laboratory methods in the clinical genetics department of Veltishev Research and Clinical Institute for Pediatrics, Moscow, Russia. Molecular genetic analysis was carried out in the DNA diagnostics laboratory of Research Centre for Medical Genetics, Moscow, Russia.

DNA extraction

DNA samples used for the analysis were extracted from whole blood samples using a Wizard[®] Genomic DNA Purification Kit (Promega, USA) according to the manufacturer's protocol.

FKBP14 variant detection

The sequence of the *FKBP14* gene ((Accession number NG_032173.1 (genomic), Accession number NM_017946.4 (mRNA)) was analyzed for possible mutations via direct automated Sanger sequencing. Primer sequences, MgCl₂ concentrations and primer annealing temperatures are presented in Table 1. PCR products were sequenced using the ABI PRISM Big Dye Terminator (v 3.1) Cycle Sequencing Kit (Applied Biosystems, Foster City, CA, USA) on an ABI3130xl Genetic Analyzer (Applied Biosystems, Foster City, CA, USA).

Sequencing results were analyzed using BLAST (Basic Local Alignment Search Tool) (<http://www.ncbi.nlm.nih.gov/blast>) to compare a subject nucleotide sequence with the database. In our research we worked with search database Human genomic plus transcript (Human G+T) using blastn (https://www.ncbi.nlm.nih.gov/Class/MLACourse/Modules/BLAST/nucleotide_blast.html) (Optimized for somewhat similar sequences).

Case descriptions

We examined five patients with EDSKS2: three boys and two girls with age ranging from two to fifteen years (Table 2). In four families, the marriage did not appear to be consanguineous. However, in family number (no.) 3 (girl, 11 years) both parents had distant relatives originating from the same small village, which could potentially mean kinship.

Table 1. Primer sequences and PCR conditions for *FKBP14* exons.

DNA fragment	Primer sequence	Fragment length, bp	MgCl ₂ concentration, mM	Primer annealing temperature, °C (cycle)
Exon 1	F-GTCGAGGGACCTTTCGCTGC	163	4	63 (32)
	R-GCTGGCATAAGTGAGTGGATTCC			
Exon 2	F-CACTTACTGGTGGGAAAATGCAC	263	4	63 (32)
	R-CTGTCTCCTAATCCAGAGAACAA			
Exon 3	F-CATATATGACAATCTTAGGAAGGCTC	240	2	65 (32)
	R-GGAGTAGGAAGAAGGAAAGGTC			
Exon 4	F-GCTCAATGTGGGTATCTTATGAATCC	690	1.6	67 (32)
	R-GCCCTCTCTTGAAGATGAGTGC			

Ethical approval

The study was approved by the Ethics committee of the Research and Clinical Institute of Pediatrics (approval number #2, 2021) named after Yuri Veltischev of the Pirogov Russian National Research Medical University of the Ministry of Health of the Russian Federation. A written informed consent was obtained from the participants and parents of participants under the age of 18 to take part in the study. The study was done in accordance with the principles outlined in the Helsinki Declaration (1964).

Results

All patients had moderate to severe kyphoscoliosis, early motor development delay, muscular weakness, hypotonia, and hearing impairment (Table 2). Figure 1 and Figure 2 (a, b) show main phenotypic characteristics of patients no. 3 and 5 with EDSKS2. Physical development of the children at the time of examination varied. Probands no. 1 and 4 (three and fifteen years respectively) had very low, harmonic development (all values below third percentile). Patients no. 3 and 5 (eleven and two years respectively) had disharmonic physical development (no. 3 - average body length 25–50 percentile and very high mass >97 percentile; no. 5 - above average body length 50–75 percentile and below average mass 10–25 percentile). Patient no. 2 had high, harmonic physical development: all values above 90–97 percentile.

The age of kyphoscoliosis formation varied from eight to twenty-four months. Clinical examination and X-ray showed thoracolumbar kyphoscoliosis in all five cases. Three children (aged two, three, eight years) had kyphoscoliosis of the 2nd degree, while the older patients (11 and 15 years) had kyphoscoliosis of the 3rd and 4th degree respectively. Probands no. 2 and 4 (eight and fifteen years), aside from kyphoscoliosis, had *pectus excavatum* of the 2nd degree. Patients no. 1, 3, 4, and 5 had *pes planovalgus*. X-ray showed signs of osteoporosis in patients no. 1, 3, and 5. In addition to this, patient 1 had *spina bifida sacralis dorsalis* S3–S5.

Joint hypermobility was evaluated as eight on the Beighton scale for four patients and as six for one patient (Patient no. 5). Four patients did not reach the maximum score of nine because of the rigid spine with restricted flexibility.

History analysis for all five probands showed early motor development delay, decrease of tendon reflexes, “floppiness”, and positive Gowers' sign. Myopathy did not progress with age. Intellectual development was normal in four children, and patient no. 3 (girl, 11 years) had a slight developmental delay – her IQ was 80 (normal values 85–115).

Apprehensive analysis of internal organs showed pathological alterations, the most notable were cardiovascular and bronchopulmonary problems. ECG detected sinus arrhythmia of varying degree in three patients (no. 1, 3, and 4). Echocardiography results revealed open *ductus arteriosus* in two patients (no. 1 and 2), and borderline (17mm) narrowed aorta at sinotubular junction (Patient no. 3). According to spirometry results, combined respiratory pulmonary function impairments were found in three probands. It was impossible to carry out spirometry for two patients due to their age (two and three years). Bilateral nephroptosis was detected in two children (Patient no. 1 and 2), left side nephroptosis was noted in one patient (no. 5) as well as midshaft hypospadias.

Hearing loss, another main feature of EDSKS2, was diagnosed in three patients (three, eleven, and fifteen years). Mild to moderate bilateral sensorineural hearing loss was revealed in the 11-year-old girl (Patient no. 3) and 15-year-old boy

Table 2. The main clinical, laboratory and functional indicators in patients with kyphoscoliotic type 2 Ehlers-Danlos syndrome.

	1	2	3	4	5
Patient number	1	2	3	4	5
Sex	M	F	F	M	M
Age of diagnosis, years	3	8	11	15	2
Intelligence Quotient	90	95	80	100	93
Physical development	Very low, harmonic, <3 percentile	Very high, harmonic; body length and mass >75-90 percentile	Disharmonic; body length 25-50 percentile, mass >97 percentile	Very low, harmonic, <3 percentile	Disharmonic; body length 50-75 percentile, mass 10-25 percentile
Kyphoscoliosis	+	+	+	+	+
Age of kyphoscoliosis onset, months	17	12	24	24	8
X-ray data	Combined thoracolumbar kyphoscoliosis of the 2nd degree. <i>Spina bifida sacralis dorsalis</i> S3-S5	<i>Pectus excavatum</i> of the 2nd degree. S-shaped thoracolumbar scoliosis of the 3d degree. (post-surgery)	Thoracolumbar dextroscoliosis of the 3d degree. Osteoporosis.	<i>Pectus excavatum</i> of the 2 nd degree. S-shaped thoracolumbar scoliosis of the 4th degree	Thoracolumbar levokyphoscoliosis of the 2nd degree. Osteoporosis.
Hypermobility Beighton score	8	8	8	8	6
Motor development impairment	+	+	+	+	+
Gowers' sign	+	+	+	+	+
Tendon reflexes decreased	+	+	+	+	+
Cardiovascular impairments	+	+	+	+	+
Congenital heart defect	Open <i>ductus arteriosus</i>	Open <i>ductus arteriosus</i> , bicuspid aortic valve	Borderline narrowed aorta – 17 mm at sinotubular junction	-	-
Electrocardiographic abnormalities	sinus arrhythmia	-	sinus arrhythmia	sinus tachyarrhythmia	-
Extrinsic respiratory restrictions (spirometry results)	Spirometry not performed	Profound combined respiratory pulmonary function impairment	Profound combined respiratory pulmonary function impairment	Profound combined respiratory pulmonary function impairment	Spirometry not performed
Ultrasound imaging of abdominal organs and kidneys	Bilateral nephroptosis	Bilateral nephroptosis			Nephroptosis (left side)
Hearing impairment	-	-	Mild to moderate bilateral sensorineural hearing loss	Bilateral sensorineural hearing loss	Mild bilateral mixed hearing loss
Vision impairment	Mild bilateral hypermetropia	-	-	Progressive high myopia with astigmatism, choroiditis (right side) without signs of inflammation	Mixed bilateral astigmatism



Figure 1. Proband no. 3: a) thoracolumbar dextroscoliosis of the 3rd degree; b) joint hypermobility, obesity of the 1st/2nd degree.



Figure 2. Proband no. 5: a) thoracolumbar levokyphoscoliosis of the 2nd degree; b) epicanthic fold, *genu valgum*, *pes planovalgus*

(Patient no. 4), mild bilateral mixed hearing loss was detected in a three-year-old boy (Patient no. 5). Ophthalmologic problems were diagnosed in three children, and the most severe alterations were in the 15-year-old boy (Patient no. 4). He had progressive high myopia with astigmatism and right side chorioretinitis without signs of inflammation. Patient

no. 1 (three-year-old boy) had mild bilateral hypermetropia and Patient no. 5 (two-year-old boy) had mixed bilateral astigmatism.

Blood and urine tests, as well as biochemical analysis indicating basic metabolism levels, were normal in all five patients.

A homozygous c.362dupC pathogenic variant in exon 3 of the *FKBP14* gene was detected in all five patients (Accession number VCV000279809.13) (Figure 3).⁶

We present a brief highlight of a child's clinical history (no. 4).

Z., a 15-year-old boy, was admitted to a clinic with complaints of progressive thoracolumbar kyphoscoliosis, *pectus excavatum*, muscular weakness, fatigue impairing his ability to engage in physical activities and to walk independently for a moderate period of time, vision and hearing impairments.

The proband was born to young healthy non-related parents (Figure 4). His mother is currently 45-years-old, and his father is 41-years-old. The proband was from the third pregnancy, second delivery. It was established that the firstborn child was also male with the same clinical picture. He underwent multiple surgical operations to correct stage 4 kyphoscoliosis, but the last operation at the age of nine years was fatal.

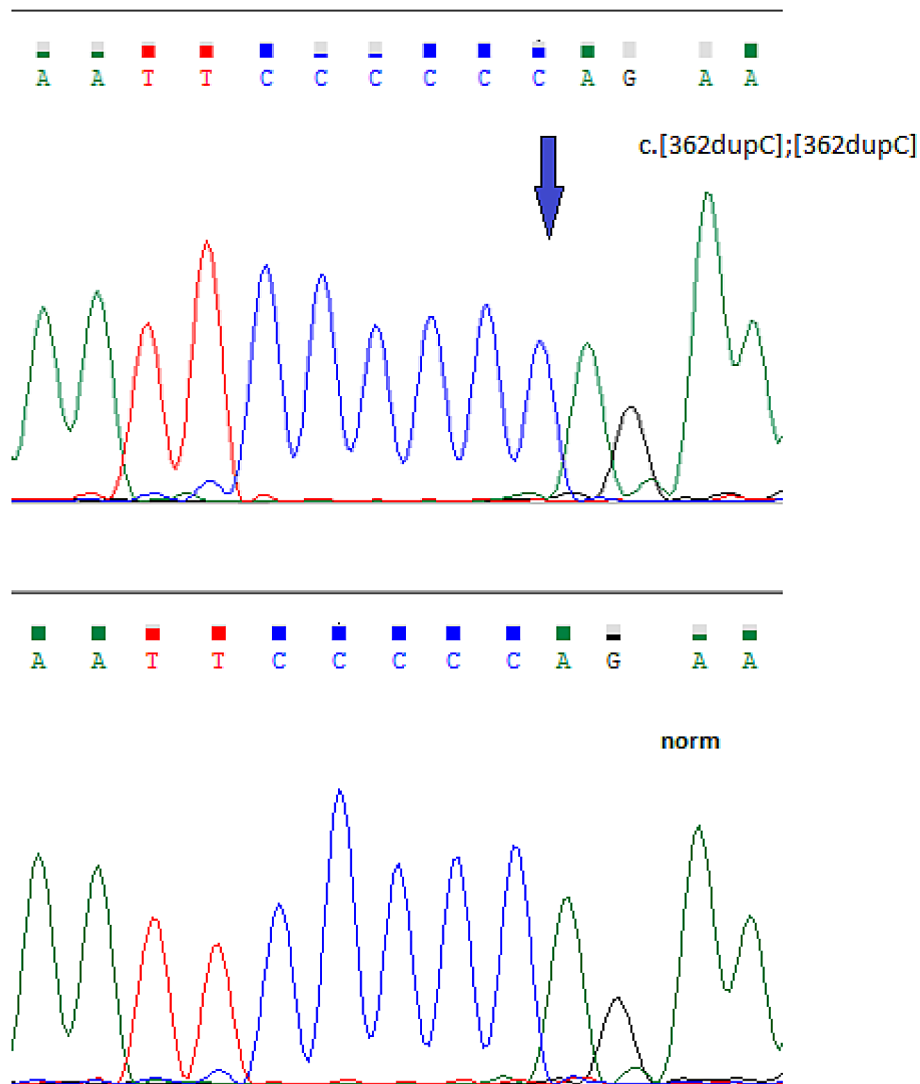
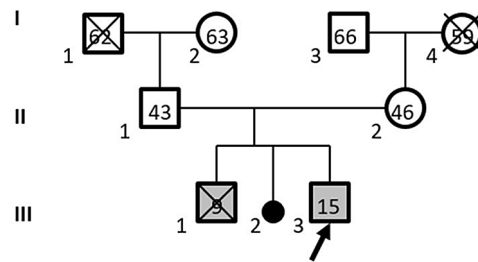


Figure 3. A homozygous c.362dupC pathogenic variant in exon 3 of the *FKBP14* gene.



III,1; III,3 – Ehlers-Danlos syndrome
 kyphoscoliotic type 2
 I,1 – myocardial infarction
 I,4 – stomach cancer

Figure 4. Pedigree of Proband no. 4, 15 years old. Roman numerals - generation number, Arabic numerals - family member number, numbers inside figures - family member's age, crossed out figures - deceased family member.

The second pregnancy was aborted medically on the mother's request. The third pregnancy was complicated by gestosis with risk of termination in trimesters I and II, but resulted in timely physiological delivery. The newborn's mass was 3000 g, body length 51 cm, APGAR score 6/7. Neonatologists stated the child's limbs were in the position of adduction. Early motor development was delayed: he started holding his head around the age of six months, sitting up at 12 months, walking independently at three years. At the age of two years progressive thoracolumbar kyphoscoliosis and *pectus excavatum* were noted. The child was under medical observation at the place of residence, received symptomatic therapy, including sanatorial treatment. However, the disorder kept progressing, and its origin was unclear to specialists. Some form of progressive muscular dystrophy was suggested; in order to obtain a clearer diagnosis, the patient was referred to the clinical genetics department of Veltischev Research and Clinical Institute for Pediatrics, Moscow, Russia.

Upon admission, the proband's condition was evaluated to be moderate to severe, in accordance with his main disorder. His physical development was very low, harmonic: body length - 140 cm, mass - 31 kg respectively (below third percentile). The following phenotypic features were the most notable (Figure 5): thoracolumbar kyphoscoliosis of the 4th degree, *pectus excavatum* of the 2nd degree, *pes planovalgus*, joint hypermobility (eight on the Beighton scale), fatigue, muscular weakness resulting in Gowers' maneuvers. Tendon reflexes were absent. Intellectual development corresponded to age.

Cardiac functional examination showed moderate sinus arrhythmia with periods of tachycardia, heart rate was 115–88 bpm. Echocardiography showed no heart defects: the chambers were not enlarged, the valves were intact, the contractory function of the myocardium was satisfactory, the diastolic function was normal, false chordae were detected in the left ventricle.

External ventilation function analysis showed profound combined respiratory pulmonary function impairment. Ultrasound examination of abdominal organs and kidneys did not reveal any pathology.

X-ray examination of the thoracolumbar spine region (dorsal-plantar and lateral) confirmed the presence of S-shaped stage 4 thoracolumbar scoliosis.

The electroneuromyography data indicated the muscular type of lesion: a slight decrease in the amplitudes of M-waves, a decrease in the amplitudes and duration of the motor unit action potentials (MUAPs), accelerated recruitment of the MUAPs; interference pattern analysis (IPA) indicators were below the standard limits.

Pure tone threshold audiometry showed mild to moderate bilateral sensorineural hearing loss. An ophthalmologist's examination showed progressive high myopia with astigmatism, as well as OD chorioretinitis without signs of inflammation.

Blood and urine tests, as well as biochemical analysis indicating basic metabolism levels, were normal. Creatine kinase level was 167 upl, reference values 15–190 upl.



Figure 5. a) Proband no. 4: very low height and body mass (<3 percentile), thoracolumbar kyphoscoliosis of the 4th degree, *pectus excavatum* of the 2nd degree; b) Proband no. 4: thoracolumbar kyphoscoliosis of the 4th degree, long thin limbs, *pes planovalgus*.

Clinical data suggested kyphoscoliotic type Ehlers-Danlos syndrome; it was also necessary to exclude congenital muscular dystrophy. The latter was suggested based on the congenital nature of the disorder, motor development delay, absence of tendon reflexes, muscular strength decrease, Gowers' maneuvers, and electroneuromyography results. However, the slow muscular pathology progression with fast kyphoscoliosis progression (stage 4), normal CK levels, severe hypermobility syndrome (eight on the Beighton scale), and mild to moderate hearing loss called the diagnosis of primary muscular pathology into question. Molecular genetic analysis results showed a c.362dupC mutation in a homozygous state in exon 3 of the *FKBP14* gene. The boy's parents had this mutation in a heterozygous state. These acquired results allowed us to confirm the EDSKS2 diagnosis.

The proband's family was given medical genetic counselling, according to which the risk of an affected child was 25%. The couple was inclined towards preimplantation diagnostics.

Discussion

In 2012, Baumann *et al.*⁵ used genetic mapping to describe the *FKBP14* gene as a cause of Ehlers-Danlos syndrome kyphoscoliotic type. Since that moment, less than 30 such patients have been described in the medical literature.^{5,7–10} To date, according to The Human Gene Mutation Database (HGMD), eight pathogenic variants are described in the *FKBP14* gene. The most common variant is the c.362dupC duplication, detected in 11 out of 17 patients by Giunta *et al.* (2018).⁹ The majority of patients with this variant are Caucasian from various countries: UK, Austria, Croatia, Poland, etc. One Columbian patient has been reported.¹⁰ The c.362dupC duplication was detected in a homozygous state in all five of our non-related patients from different regions of Russia. Baumann *et al.* (2012)⁵ suggested the presence of a founder effect. However, a recurrent mutation caused by replication slippage in the polycytidine tract cannot be excluded.

The *FKBP14* gene encodes the FKBP22 protein, which is endoplasmic reticulum (ER) resident and controls post-translational modification of polypeptide chains containing hydroxyproline. FKBP22, a peptidyl-prolyl *cis-trans* isomerase, is a collagen folding catalyst and interacts with collagen types III, VI, and X.^{5,11} *FKBP14*-deficient skin fibroblast examination showed normal proportion and electrophoretic mobility of type I, III, and V collagen α -chains. Immunofluorescence revealed disorganisation of extracellular matrix proteins – collagen types I, III, VI, fibronectin, tenascins. Electronic microphotography of skin fibroblasts showed ER cisternae enlargement, fragmentation of elastic

fibers. Symptoms of EDSKS1 and EDSKS2 are very similar: progressive kyphoscoliosis, severe congenital muscular hypotonia, hypermobility syndrome, skin hyperelasticity, medium arterial ruptures, osteopenia/osteoporosis. Diagnostic differentiation criteria: *PLOD1*-associated kyphoscoliotic type 1 - marfanoid phenotype, profound skin bruisability, scleral and ocular ruptures; *FKBP14*-associated kyphoscoliotic type 2 - hearing loss, follicular hyperkeratosis, muscular atrophy, bladder diverticulum. In addition, lysyl pyridinoline (LP) and hydroxylysyl pyridinoline (HP) excretion analysis may be useful - LP/HP ratio is increased in the case of kyphoscoliotic type 1.

Clinical findings in our patients corresponded to those described in the literature.^{2,5} The main symptom was progressive kyphoscoliosis in varying stages depending on the age. Three out of five patients had hearing loss, which is described in 73% of kyphoscoliotic type 2 cases.⁴ The exact pathogenesis of this impairment is unclear. Three patients had osteoporosis; according to the literature, it is one of the minor diagnostic criteria, leading to fractures in 13% of cases.^{4,9} Three children had congenital malformations affecting three systems: cardiovascular (open *ductus arteriosus*, borderline narrowed aorta), skeleto-muscular (*spina bifida sacralis dorsalis* S3-S5), and genital (midshaft hypospadias). Karyotype analysis and chromosomal microarray analysis did not show any abnormalities.

As shown in the literature, there are various types of Ehlers-Danlos syndrome with vascular complications (vascular, classic, classic-like, musculocontractural), which should be considered in differential diagnostics.^{1,7,12} These complications in kyphoscoliotic type 2 patients could be caused by a pathogenetic link of the *FKBP14* gene with type III collagen, which is a significant component of vascular structure.¹³

Kyphoscoliotic type 2 with symptoms of severe muscular atrophy has to be differentiated from Ullrich and Bethlem myopathy, which is caused by type VI collagen defects. The latter does not cause skin hyperelasticity, ecchymoses, hearing loss, but does cause striae, atrophic scars, respiratory abnormalities, and major joint contractures, uncharacteristic of EDSKS2.

Without any doubt, with such similarities in clinical pictures of EDSKS2 and the above-mentioned disorders, the final diagnosis has to be verified by molecular genetic analysis.

Conclusion

Literature analysis and examination of five EDSKS2 patients demonstrated the involvement of major organs and systems, such as joints, spine, muscles, cardiovascular system, respiratory system, hearing, and vision, into the pathological process. Kidney mobility increases and nephroptosis seems to be secondary, caused by muscular weakness. During molecular genetic analysis, to verify EDSKS2 it is recommended to initially search for the c.362dupC duplication, which appears to be common in European countries, including Russia.

Many questions regarding the disorder's clinical polymorphism and progressive course remain unanswered. Some of them might be solved by a more detailed analysis of the *FKBP14* gene functions. The obtained information would improve our understanding of the disorder's pathogenetic mechanisms and aid in target therapy development.

Data availability

Underlying data

ClinVar: NM_017946.4(FKBP14):c.362dup (p.Glu122fs). Accession number VCV000279809.13; variation ID 279809; <https://identifiers.org/clinvar:279809>.

NCBI Nucleotide: Homo sapiens FKBP prolyl isomerase 14 (FKBP14), RefSeqGene (LRG_454) on chromosome 7. Accession number NG_032173.1; https://identifiers.org/ncbiprotein:NG_032173.1.

NCBI Nucleotide: Homo sapiens FKBP prolyl isomerase 14 (FKBP14), transcript variant 1, mRNA. Accession number NM_017946.4; https://identifiers.org/ncbiprotein:NM_017946.4.

4TU.ResearchData: Underlying data for: Ehlers-Danlos syndrome kyphoscoliotic type 2 caused by mutations in the *FKBP14* gene: an analysis of five cases. <https://doi.org/10.4121/14705859.v1>.⁶

This project contains the following underlying data:

- Picture file 1. Gel of PCR fragments from patient no. 3. *FKBP14* gene, exons 1–4.

- Picture file 2. Electropherogram of the exon 3 of the *FKBP14* gene. A homozygous c.362dupC pathogenic variant in exon 3 of the *FKBP14* gene.
- Data file 1. Readme.pdf.

Data are available under the terms of the Creative Commons Zero “No rights reserved” data waiver ([CC0 1.0 Public domain dedication](#)).

Consent

Consent for publication

A written informed consent for the publication of this manuscript including identifying images and other personal and clinical details was obtained from the participants and parents or legal guardians of all participants under the age of 18.

References

- Malfait F, Francomano C, Byers P, *et al.*: **The 2017 international classification of the Ehlers-Danlos syndromes.** *Am J Med Genet C Semin Med Genet.* 2017; **175**(1): 8–26.
[PubMed Abstract](#) | [Publisher Full Text](#)
- Rohrbach M, Vandersteen A, Yiş U, *et al.*: **Phenotypic variability of the kyphoscoliotic type of Ehlers-Danlos syndrome (EDS VIA): clinical, molecular and biochemical delineation.** *Orphanet J Rare Dis.* 2011; **6**: 46.
[PubMed Abstract](#) | [Publisher Full Text](#) | [Free Full Text](#)
- Kivirikko KI, Myllylä R: **Posttranslational Enzymes in the Biosynthesis of Collagen: Intracellular Enzymes.** *Methods Enzymol.* 1982; **82**: 245–304.
[PubMed Abstract](#) | [Publisher Full Text](#)
- Lim PJ, Lindert U, Opitz L, *et al.*: **Transcriptome Profiling of Primary Skin Fibroblasts Reveal Distinct Molecular Features Between PLOD1- and FKBP14-Kyphoscoliotic Ehlers-Danlos Syndrome.** *Genes.* 2019; **10**(7): 517.
[PubMed Abstract](#) | [Publisher Full Text](#) | [Free Full Text](#)
- Baumann M, Giunta C, Krabichler B, *et al.*: **Mutations in FKBP14 Cause a Variant of Ehlers-Danlos Syndrome with Progressive Kyphoscoliosis, Myopathy, and Hearing Loss.** *Am J Hum Genet.* 2012; **90**(2): 201–216.
[PubMed Abstract](#) | [Publisher Full Text](#) | [Free Full Text](#)
- Semyachkina AN, Nikolaeva EA, Galeeva NM, *et al.*: **Underlying data for 'Ehlers-Danlos syndrome kyphoscoliotic type 2 caused by mutations in the FKBP14 gene: an analysis of five cases.** *4TU. ResearchData.* 2021.
[Publisher Full Text](#)
- Dordoni C, Ciaccio C, Venturini M, *et al.*: **Further delineation of FKBP14-related Ehlers-Danlos syndrome: A patient with early vascular complications and non-progressive kyphoscoliosis, and literature review.** *Am J Med Genet A.* 2016; **170**(8): 2031–2038.
[PubMed Abstract](#) | [Publisher Full Text](#)
- Bursztejn AC, Baumann M, Lipsker D: **Ehlers-Danlos syndrome related to FKBP14 mutations: detailed cutaneous phenotype.** *Clin Exp Dermatol.* 2017; **42**(1): 64–67.
[PubMed Abstract](#) | [Publisher Full Text](#)
- Giunta C, Baumann M, Fauth C, *et al.*: **A cohort of 17 patients with kyphoscoliotic Ehlers-Danlos syndrome caused by biallelic mutations in FKBP14: expansion of the clinical and mutational spectrum and description of the natural history.** *Genet Med.* 2018; **20**(1): 42–54.
[PubMed Abstract](#) | [Publisher Full Text](#) | [Free Full Text](#)
- Ruiz-Botero F, Ramírez-Montaño D, Pachajoa H: **FKBP14 kyphoscoliotic Ehlers-Danlos Syndrome in adolescent patient: the first Colombian report.** *Arch Argent Pediatr.* 2019; **117**(3): e274–e278.
[PubMed Abstract](#) | [Publisher Full Text](#)
- Ishikawa Y, Mizuno N, Holden P, *et al.*: **The novel missense mutation Met48Lys in FKBP22 changes its structure and functions.** *Sci Rep.* 2020; **10**(1): 497.
[PubMed Abstract](#) | [Publisher Full Text](#) | [Free Full Text](#)
- Murray ML, Yang M, Fauth C, *et al.*: **FKBP14-related Ehlers-Danlos syndrome: Expansion of the phenotype to include vascular complications.** *Am J Med Genet Part A.* 2014; **164**: 1750–1755.
[PubMed Abstract](#) | [Publisher Full Text](#)
- Germain DP: **Ehlers-Danlos syndrome type IV.** *Orphanet J Rare Dis.* 2007; **2**: 32.
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Tomoki Koshi 

Department of Medical Genetics, Shinshu University School of Medicine, Nagano, Japan

The reviewer has evaluated a paper by Dr. Semyachkina, describing five patients with EDSKS2 caused by the same common variant in *FKBP14*. The disorder is very rare, and detailed clinical information would be valuable to understand the whole picture of the disorder including the natural history. Several issues have been found to be considered as follows:

1. X-ray images of the spine, hands, and feet had better be considered to be included.
2. How about the intellectual status of these patients, normal or subnormal?
3. How about craniofacial features of these patients, some similarities among the series?
4. How about skin involvement, hyperextensibility, fragility, and bruisability?

Is the work clearly and accurately presented and does it cite the current literature?

Yes

Is the study design appropriate and is the work technically sound?

Yes

Are sufficient details of methods and analysis provided to allow replication by others?

Yes

If applicable, is the statistical analysis and its interpretation appropriate?

Not applicable

Are all the source data underlying the results available to ensure full reproducibility?

Yes

Are the conclusions drawn adequately supported by the results?

Yes

Competing Interests: No competing interests were disclosed.

Reviewer Expertise: Clinical genetics

I confirm that I have read this submission and believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard.

Reviewer Report 12 July 2021

<https://doi.org/10.5256/f1000research.55519.r88243>

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Matthias Baumann

Division of Pediatric Neurology, Department of Pediatrics I, Medical University of Innsbruck, Innsbruck, Austria

The manuscript presents in detail the findings in five additional patients with the rare kyphoscoliotic Ehlers-Danlos syndrome type 2 caused by mutations in the FKBP14 gene. They carried the most common c362dupC duplication. The patients of different age illustrate very well the spectrum of clinical symptoms. An interesting aspect, not well described so far, is the increased kidney mobility and nephroptosis.

The paper helps to increase the awareness for this rare connective tissue disease, which is probably still underdiagnosed.

Is the work clearly and accurately presented and does it cite the current literature?

Yes

Is the study design appropriate and is the work technically sound?

Yes

Are sufficient details of methods and analysis provided to allow replication by others?

Yes

If applicable, is the statistical analysis and its interpretation appropriate?

Not applicable

Are all the source data underlying the results available to ensure full reproducibility?

Yes

Are the conclusions drawn adequately supported by the results?

Yes

Competing Interests: No competing interests were disclosed.

Reviewer Expertise: Neuromuscular disorders, Ehlers-Danlos-Syndrome

I confirm that I have read this submission and believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard.

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