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Thauvin-Robinet-Faivre Syndrome: A *FIBP* Variant in an Adolescent with Segmental Overgrowth and Thyroid Carcinoma

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What is already known on this topic?

Overgrowth syndromes are rare diseases seen in pediatric endocrinology clinics. Those affecting growth factor pathways may present with body parts or limb hypertrophy and a predisposition to malignancy, including endocrine organs. *Fibroblast growth factor 1 intracellular binding protein* gene variants cause Thauvin-Robinet-Faivre syndrome (TROFAS), a new disease defined in this pathway.

What this study adds?

The present case represents the ninth instance of TROFAS reported worldwide and the fifth from Türkiye, contributing to the literature in terms of clinical findings and clinical presentations in cases with suspected tumor predisposition.

ABSTRACT

Overgrowth syndromes are rare genetic disorders arising from alterations in the growth factors pathway. These syndromes can present as generalized overgrowth, characterized by macrosomia and excessive height compared to peers, or partial overgrowth syndromes, where specific body regions exhibit disproportionate growth often accompanied by vascular anomalies. Both forms are associated with an increased risk of tumor development. The *fibroblast growth factor 1 (FGF-1) intracellular binding protein (FIBP)* gene plays a critical role in cell proliferation and differentiation by interacting with growth factors. In this article, we present a case of Thauvin-Robinet-Faivre syndrome (TROFAS) in a 16-year-old girl, diagnosed with homozygous NM_004214.5 c.412-3_415dup (p. Asp139AlafsTer3) variant in the *FIBP* gene. This case exhibits

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phenotypic features and tumor development, including thyroid follicular carcinoma and parotid mucoepidermoid carcinoma, that have not been previously reported in association with this syndrome. Recent studies have implicated FIBP gene defects in overgrowth syndromes, with only a limited number of cases described globally. This case expands the known clinical and tumor spectrum associated with TROFAS, providing new insights into the pathophysiology of this rare disorder.

Keywords: Overgrowth syndromes, *FGF-1 intracellular binding protein (FIBP)* gene, cancer

Introduction

Overgrowth syndromes comprise a diverse group of disorders characterized by prenatal overgrowth, persistent postnatal overgrowth in both weight and height, segmental overgrowth, congenital malformations, intellectual disability, and, in some cases, an increased risk of neoplasia. Segmental overgrowth refers to excessive growth localized to specific body regions, such as a single digit, an extremity, one side of the face, or the head (macrocephaly). It often presents as asymmetrical overgrowth of musculoskeletal, adipose, or brain tissue, vascular malformations and/or overlying skin lesions. This phenotype is frequently associated with mosaic variants in the phosphoinositide-3-kinase-protein kinase B (PI3K/AKT)/mammalian target of rapamycin (mTOR) pathway, a key signaling cascade activated by growth factors such as insulin-like growth factor-1 (IGF-1), vascular endothelial growth factor (VEGF), and epidermal growth factor (EGF) (1,2,3).

The *fibroblast growth factor 1 (FGF-1) intracellular binding protein (FIBP)* gene, located on chromosome 11q13.1, is expressed in the membranes of microsomes and mitochondria within the cytoplasm and nucleus (4). The interaction between FGFs and their receptors (FGFRs) plays a crucial role in multiple biological processes, including embryogenesis, neuronal development, and the formation of bone, cartilage, and vascular structures (4,5).

Thauvin-Robinet-Faivre Syndrome (TROFAS) is a rare autosomal recessive disorder caused by biallelic pathogenic variants in the *FIBP* gene. TROFAS (OMIM: 617107) is characterized by a broad spectrum of phenotypic features, including mild to severe learning difficulties, distinctive facial dysmorphisms, enlarged hands and feet, generalized overgrowth, and a predisposition to various congenital anomalies affecting the heart, eyes, kidneys, and skeleton. Only eight cases of TROFAS have been reported worldwide, making it a largely under-recognized condition with limited available literature (6,7,8,9).

Here, we present the case of a 16-year-old female patient diagnosed with a homozygous likely pathogenic variant in the *FIBP* gene, born to consanguineous parents. This case is particularly notable due to the malignancies and distinct clinical features not previously described in other reported TROFAS cases.

Case Report

A 16-year-old female presented with a six-month history of swelling on the right side of her neck and chin. She had previously undergone surgery at another hospital, where a right hemithyroidectomy and excision of a right parotid mass were performed. Histopathological examination revealed thyroid follicular carcinoma and a low-grade mucoepidermoid carcinoma of the parotid gland.

The patient was born at full term to consanguineous parents, with a birth weight of 2800 g. Her neuromotor developmental history was notable for significant delays; she began walking at the age of 3 years and started speaking between 6 and 7 years of age. The family history included two healthy siblings, one sister with congenital heart disease, and two siblings who died in the neonatal period for unknown reasons.

On physical examination, her weight was 59.6 kg (66th percentile), height was 159.3 cm (29th percentile), and body mass index was 23.5 kg/m² (79th percentile). Her head circumference measured 56.8 cm (77th percentile). Sitting height was 85 cm, with a sitting height-to-height ratio of 0.53 (within the normal range). Her arm span was 163 cm, with an arm span-height difference of 4 cm [+1 standard deviation (SD) to +2 SD]. Pubertal assessment was consistent with Tanner stage 5; she had menarche at 12.5 years, and her menstrual cycles were regular.

Notable facial features included a flat midface, mild right ptosis, low palpebral fissures, deep-set eyes, thick lips, thick and broad eyebrows with mild synophrys, a pointed chin, a prominent nasal bridge, a narrow forehead, a high palate, posteriorly rotated ears, and raised earlobes. Skeletal examination revealed mild pectus excavatum and pes planus. In addition, she had enlarged hands and feet, broad port-wine stains on her arms and legs, and varicose veins in both legs. Syndactyly was present between the second and third toes on the left foot, and a flexion contracture was noted between the first and second phalanges of the third finger on the right hand, which developed following hemangioma surgery (Figure 1 and 2). Numerous nevi were also observed on her face and trunk. The patient had an intellectual disability.

Parental heights were 155 cm (-3.4 SD) for the father and 157 cm (-1.04 SD) for the mother. Her 25-year-old brother's height was 169 cm (-1.17 SD), and her 20-year-old sister's height was

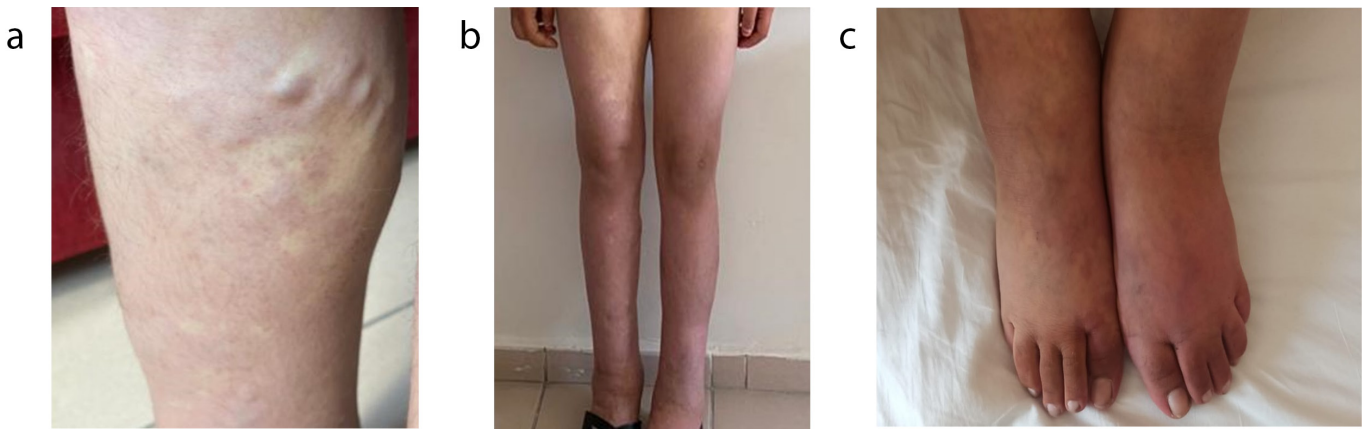


Figure 1. a) The patient's hand is noticeably larger than a normal hand. b) Capillary malformation (Port-wine Stain) on the hand extending to the arm and contracture of the fingers due to vascular anomaly

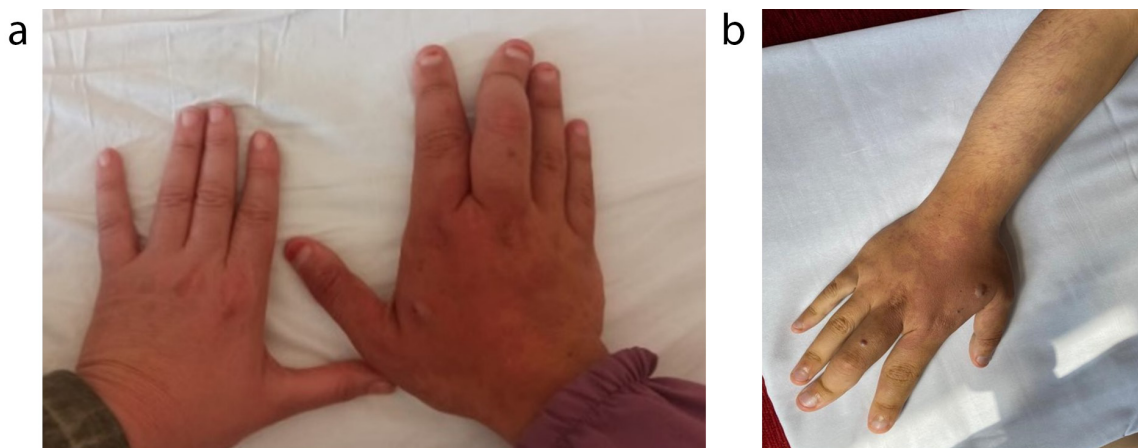


Figure 2. a) Varicose veins. b) Port-wine-Stain on legs and feet, wide ankles, and feet. c) Syndactyly between the second and third toes of the left foot

158 cm (-0.87 SD). It was reported that her 14-year-old sister has a normal height but is being followed at another center for a cardiac condition.

Laboratory tests, including a complete blood count and biochemistry, were unremarkable. Eye and hearing examinations were normal; her IQ score was 65. Radiographic imaging showed mild thoracic scoliosis and closed epiphyses. Abdominal and renal ultrasonography did not detect any anomaly. Anatomical variation was noted, with the hepatic artery originating directly from the abdominal aorta, while the splenic and left gastric arteries arose from a single branch. Venous Doppler imaging revealed multiple enlarged perforating veins with reflux, connected to the great saphenous vein, with numerous varicose veins in both legs. An echocardiogram detected mild mitral valve prolapse, trace mitral regurgitation, and a small atrial septal defect.

Surgery and pathology reports indicated that the 3x2.5 cm diameter nodule removed from the right lobe hemithyroidectomy material was non-invasive follicular carcinoma, and the 3.5x3 cm mass removed from the right parotid was low-grade mucoepidermoid carcinoma. No metastatic or residual mass was detected in further scans.

Methods and Results

We performed whole exome sequencing (WES) for the patient and her healthy father using the Illumina DNA Prep & TWIST Bioscience Exome 2.0 Plus Comprehensive Exome library preparation and the Illumina NovaSeq 6000 sequencing platform were utilized. The bioinformatics analysis pipeline included the Burrows-Wheeler Aligner (BWA 0.7.15) for read alignment, the Genome Analysis Toolkit (GATK 3.6) for variant calling, and the Variant Effect Predictor (VEP 89) for annotation. Variants were

filtered based on their frequency using public and in-house databases (e.g., gnomAD) (10,11,12,13).

Variant pathogenicity was assessed according to the 2015 American College of Medical Genetics and Genomics (ACMG) guidelines (14) Variants in the *FIBP* gene that met the analysis criteria, correlated with the clinical phenotype, and had variant allele frequency values consistent with the expected inheritance pattern were reported. WES revealed a homozygous c.412-3_415dup p.Asp139AlafsTer3 frameshift variant which was assessed to be likely pathogenic in the *FIBP* gene (NM_004214.5), classified according to ACMG criteria as PVS1-PM2. Family segregation analysis confirmed that the father was a heterozygous carrier of this variant, but maternal carrier status could not be verified due to the lack of available DNA samples. The *FIBP* gene is known to follow a loss-of-function disease mechanism, and the identified variant was absent in population databases, including gnomAD, ExAC, and the Turkish Variome.

Discussion

That the protein product of *FIBP* binds acidic FGF intracellularly, leading to morphological changes, differentiation, and stimulation of mitogenic activity in various cell types (4). The FGF receptor (FGFR) family, similar to IGF receptors, functions through tyrosine kinase signaling pathways. Activation of FGFRs initiates downstream signaling cascades, including the MAPK and PI3K/AKT pathways, which play central roles in cellular differentiation, proliferation, survival, and migration. These pathways are also involved in angiogenesis, vascular repair, and wound healing. Aberrations in FGFR genes such as translocations, amplifications, or activating variants have been implicated in various malignancies, including gastric, lung, breast, and ovarian cancers (15).

Thauvin-Robinet et al. (16) first described TROFAS in 2016 in a 23-year-old male of North African descent, born to consanguineous parents. The patient exhibited overgrowth, macrocephaly, intellectual disability, facial dysmorphism, bilateral retinal coloboma, ventricular septal defect, mitral valve prolapse bifid ureter, renal malrotation, transient neutropenia, and varicose venous anomalies. The biallelic variant identified in the *FIBP* gene suggested that loss of function in the intracellular domain of FGFRs may be associated with overgrowth syndromes. Further analysis of the patient's fibroblasts demonstrated significantly reduced *FIBP* cDNA expression but an increased proliferation rate compared to controls, indicating a potential link between *FIBP* dysfunction and overgrowth through the FGFR3 pathway (16,17).

In 2016, a homozygous indel variant (NM_198897.1 c.175_176insTAA, p.His59delinsLeuAsn) was identified in three siblings from an Arab family, each presenting with an overgrowth

syndrome accompanied by congenital abnormalities. Functional studies using *in vitro* and *in vivo* models demonstrated that patient-derived fibroblasts exhibited markedly increased proliferation rates, reinforcing the association of *FIBP* loss-of-function variants with humans overgrowth syndromes (18).

A recent case from Türkiye described a 9-year-old boy with distinct dysmorphic features, including macrosomia at birth, neuromotor delay, and intellectual disability. His facial features included a triangular face, midface retrusion, pointed chin, deep-set eyes, hypertelorism, long and wavy palpebral fissures, a prominent nasal bridge, thick and broad eyebrows with mild synophrys, a short philtrum, a high palate, and posteriorly rotated ears with upturned earlobes. Additional findings included chest narrowing, mild pectus excavatum, hyperextensible elbow joints, pes planus, telangiectasias, hypoplastic nipples, and unilateral cryptorchidism. Genetic analysis via exome sequencing identified a homozygous frameshift variant (NM_004214.5 c.412-3_415dup, p.Asp139AlafsTer3) in the *FIBP* gene, which was classified as likely pathogenic (6). Subsequently, three additional cases with a similar phenotype carrying the same homozygous NM_004214.5 c.412-3_415dup, p.Asp139AlafsTer3 variant were reported from Türkiye. All reported cases' clinical and genetic characteristics are summarized in Table 1 (6,7,8,9).

The same frameshift variant was also detected in our patient, introducing a premature stop codon in exon 4 of 10, leading to a truncated or non-functional *FIBP* protein compatible with TROFAS. The loss of *FIBP* function impairs its interaction with growth factors, contributing to dysregulated growth and tumorigenesis, fitting with the observed overgrowth and tumor predisposition in TROFAS. Unlike previously reported cases, our patient did not present with macrosomia, tall stature, macrocephaly, or renal anomalies. However, she shared several overlapping features, including facial dysmorphism, intellectual disability, skeletal abnormalities, and cardiac defects. Moreover, varicose venous enlargement, previously reported in the 23-year-old male described by Thauvin-Robinet et al. (16), was also present in our younger patient, although this feature has not been consistently observed in other cases. Further imaging revealed that our patient also had multiple hemangiomas scattered throughout the body.

Segmental overgrowth syndromes are associated with segmental mosaicism, where a postzygotic variant arising during embryogenesis disrupts tissue growth regulation. Overgrowth can result from either increased cell size (hypertrophy) or increased cell proliferation (hyperplasia), with the severity of symptoms depending on the affected gene and its role in cellular growth and survival (19).

Syndromes associated with somatic *PIK3CA* variants are classified under the *PIK3CA*-related overgrowth spectrum

Table 1. genetic characteristics of the nine reported cases of Thauvin-Robinet-Faivre Syndrome (TROFAS)

	Case 1	Case 2	Case 3	Case 4	Case 5
Age	17 years, male	14 years, female	10 years, male	3 years, female	4 years, male
Macrosomia	+	+	+	-	+
Tall stature	+	+	+	+	+
Intellectual Disability/developmental delay	+	+	+	+	+
Macrocephaly	+	-	-	-	+
Cardiovascular anomalies	VSD MVP Varicose veins	VSD Double chamber right ventricle	-	-	Small PDA Multiple telangiectasias
Genitourinary anomalies	Renal malrotation Left bifid ureter	-	-	Cystic dysplastic kidneys	Mild right renal pelvis dilatation Unilateral cryptorchidism
Hearing loss	-	+	-	-	-
Dysmorphic features	Occipital prominent aspect Long and down-slanting palpebral fissures Large and prominent ears Thick lips Mild macroglossia	Round face Widely spaced eyes Epicanthic folds Depressed nasal bridge Short, small nose Flat mid-face and full lips	Round face Widely spaced eyes Epicanthic folds Depressed nasal bridge Short, small nose Flat mid-face and full lips Strabismus	Round face Widely spaced eyes Epicanthic folds Depressed nasal bridge Short, small nose Flat mid-face and full lips	Triangular face Midface retrusion Pointed chin Deeply set eyes Hypertelorism Long and wave-shaped palpebral fissures Prominent nasal bridge Short philtrum High palate, Posteriorly rotated ears and uplifted earlobes Thick and broad eyebrows with mild synophrys Hypoplastic nipples
Skeletal abnormalities	Hands and feet were large with large thumbs and hallux Bilateral asymmetric limitation of thumb extension Spatulate digits Camptodactyly		Bilateral talipes equino varus, Rotation of both femurs and tibiae Spina bifida occulta		Narrow chest Mild pectus excavatum, Hyperextensible elbow joints Pes planus
Other	Bilateral retinal coloboma Transient neutropenia. Inguinal hernia Psychosis	Chronic benign neutropenia	Chronic benign neutropenia	Polyhydramnios	Multiple nevus Fibrotic changes in bilateral basal lung segments consistent with chronic pulmonary disease
Tumor/cancer predisposition	-	Wilms tumor	-	-	-
<i>FIBP</i> gene variant	Homozygous NM_004214.5 c.652C>T	Homozygous NM_198897.1 c.175_176insTAA	Homozygous NM_198897.1 c.175_176insTAA	Homozygous NM_198897.1 c.175_176insTAA	Homozygous NM_004214.5 c.415_416insCAGTTTG
Author	Thauvin-Robinet et al. (17) 2016	Akawi et al. (18) 2016	Akawi et al. (18) 2016	Akawi et al. (18) 2016	Duzenli et al. (6) 2023

Table 1. Continued			
Case 6	Case 7	Case 8	Present case
16 years, male	5 years, female		16 years, female
-	-	+	-
+	+		-
+	+	+	+
+	+		-
Minimal pericardial effusion	-	-	Varicose veins Hemangiomas Hepatic artery originating from the abdominal aorta Mild MVP Small ASD
Right renal atrophy	-	-	-
-	-	-	-
Macrocephaly Round face Widely-spaced deep-set eyes Prominent supraorbital ridges Thick eyebrows Down-slanting palpebral fissures Marked philtrum	Round face Flat midface Widely-spaced eyes Thick eyebrows Marked philtrum.	Hypertelorism Thick and arched eyebrows Pointed chin Down-slanting palpebral fissures Epicanthic folds Broad nasal bridge Small nose High palate	Flat midface Mild right ptosis Low palpebral fissures Deep-set eyes Thick lips Thick and broad eyebrows Mild synophrys Pointed chin Prominent nasal bridge Narrow forehead High palate Posteriorlyrotated ears and raised earlobes
Scoliosis Pectus deformity Thoracic asymmetry Large hand Large feet Talipes varus	-	Coxa valga	Mild pectus excavatum and pes planus Enlarged hands and feet
Parieto-occipital perivascular cystic gliotic changes Slightly decreased muscle strength in the lower extremities	Cerebral periventricular patchy hyperintense lesions	-	Port-wine stains on the arms and legs Widespread nevi on face and trunk
-	-	-	Parotis low-grade mucoepidermoid carcinoma Thyroid follicular carcinoma
Homozygous NM_004214.5 c.412-3_415dupCAGTTTG	Homozygous NM_004214.5 c.412-3_415dupCAGTTTG	Homozygous NM_004214 c.412-3_415dupCAGTTTG	Homozygous NM_004214.5 c.412-3_415dup
Kılıç and Koşukcu (7) 2023	Kılıç and Koşukcu (7) 2023	Yüksel Ülker et al. (8) 2024	
ASD: atrial septal defect, VSD: ventricular septal defect, MVP: mitral valve prolapse, PDA: patent ductus arteriosus			

(PROS), which encompasses conditions such as fibro adipose overgrowth, hemihyperplasia-multiple lipomatosis, dysplastic megalencephaly, congenital lipomatous overgrowth, vascular malformations, epidermal nevi, skeletal anomalies (CLOVES syndrome), Klippel-Trenaunay syndrome, macrodactyly, and others. These disorders share a common genetic origin but present a broad range of phenotypic expressions, from isolated macrodactyly to more complex cases. The variability in symptom severity is influenced by the timing and cellular context of the variant during development (20,21).

Other genes in the PI3K signaling pathway, including *PTEN* (a negative regulator), are also implicated in segmental overgrowth syndromes. Loss-of-function variants in *PTEN* increase PIP3 levels, activate the AKT/mTOR pathway, and lead to conditions such as germline and mosaic forms of *PTEN* hamartoma tumor syndrome (PHTS). For instance, SOLAMEN syndrome results from a mosaic loss of *PTEN* function and is characterized by localized hypertrophy, vascular malformations, macrodactyly, and an increased cancer risk. Furthermore, activating variants in genes, including *PIK3R2*, *AKT1*, *AKT2*, *AKT3*, and *CCND2* have been identified. Somatic activation of *AKT1* is associated with Proteus syndrome, which is marked by asymmetrical lesions, lipomatous tumors, lymphovascular malformations, and overgrowth of bones and connective tissues, also with a higher cancer risk (22,23).

The presence of a prominent hemangiomas lesion, along with intellectual disability, excessive soft tissue and bone overgrowth, skeletal abnormalities, and an increased risk of tumors, led us to evaluate the possibility of segmental overgrowth syndromes. TROFAS affects similar genetic pathways and presents with comparable clinical features, and may be included within the *PIK3CA*-PROS. Our patient was thoroughly evaluated for other overgrowth syndromes (21,24). Since *PIK3CA* mosaicism may be present in the affected tissues, it was impossible to completely rule out other diseases in this group for our patient.

Recent research suggests that *FIBP* regulates the expression of the mTOR/STAT3 pathway, a signaling molecule implicated in carcinogenesis, which provides a potential mechanism for the increased cancer risk observed in overgrowth syndromes (25). *FIBP* overexpression has been reported in tumors, such as colon carcinoma and head and neck cancers. Given that FGF1 is linked to several cancers, it is hypothesized that the interaction between *FIBP* and FGF1 plays a role in tumorigenesis (5,26,27,28).

The case described above is particularly notable due to the early onset of multiple malignancies. By the age of 16 years, the patient had developed two distinct cancers: thyroid follicular carcinoma and mucoepidermoid carcinoma of the parotid gland. Among the eight reported cases of TROFAS in the literature, only a single instance of Wilms' tumor has been documented (18). This case

suggests a broader diversity of malignancies in TROFAS and an increased spectrum of cancer predisposition associated with *FIBP* gene variants.

Conclusion

The presented case is significant for its distinct phenotypic features and the early development of malignancies, highlighting the role of *FIBP* variants in overgrowth syndrome and cancer predisposition. It should also be remembered that individuals with segmental overgrowth are not always macrosomic at birth, and their height and weight during childhood may remain within normal ranges.

Ethics

Informed Consent: Consent was obtained from the patient and her parents for the scientific use of genetic testing and photographs.

Footnotes

Authorship Contributions: Surgical and Medical Practices: Ülkü Gül Şiraz, Deniz Koçak Göl, Concept: Ülkü Gül Şiraz, Nihal Hatipoğlu, Design: Ülkü Gül Şiraz, Data Collection or Processing: Ülkü Gül Şiraz, Ekrem Ünal, Analysis or Interpretation: Meino Rohlf, Christoph Klein, Literature Search: Ülkü Gül Şiraz, Ekrem Ünal, Writing: Ülkü Gül Şiraz.

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References

1. Neylon OM, Werther GA, Sabin MA. Overgrowth syndromes. *Curr Opin Pediatr.* 2012;24:505-511.
2. Bertino F, Braithwaite KA, Hawkins CM, Gill AE, Briones MA, Swerdlin R, Milla SS. Congenital limb overgrowth syndromes associated with vascular anomalies. *Radiographics.* 2019;39:491-515.
3. Manor J, Lalani SR. Overgrowth syndromes-evaluation, diagnosis, and management. *Front Pediatr.* 2020;8:574857.
4. Kolpakova E, Frengen E, Stokke T, Olsnes S. Organization, chromosomal localization and promoter analysis of the gene encoding human acidic fibroblast growth factor intracellular binding protein. *Biochem J.* 2000;352 Pt 3:629-635.
5. Xie Y, Su N, Yang J, Tan Q, Huang S, Jin M, Ni Z, Zhang B, Zhang D, Luo F, Chen H, Sun X, Feng JQ, Qi H, Chen L. FGF/FGFR signaling in health and disease. *Signal Transduct Target Ther.* 2020;5:181.
6. Duzenli T, Sezer A, Kayhan G, Arslan AT, Percin FE. Expanding the phenotype and genotype in Thauvin-Robinet-Faivre syndrome: a new patient with a novel variant and additional clinical findings. *Am J Med Genet A.* 2023;191:2232-2239. Epub 2023 May 23
7. Kılıç E, Koşukcu C. New cases of recently described Thauvin-Robinet-Faivre syndrome with a novel homozygous *FIBP* gene variant. *Am J Med Genet A.* 2024;194:e63449. Epub 2023 Oct 24
8. Yüksel Ülker A, Uludağ Alkaya D, Çağlayan AO, Usluer E, Aykut A, Aslanger A, Vural M, Tüysüz B. An investigation of the etiology and follow-up findings in 35 children with overgrowth syndromes, including biallelic *SUZ12* variant. *Am J Med Genet A.* 2023;191:1530-1545. Epub 2023 Mar 15

9. Duzenli T, Sezer A, Percin FE. Letter to the Editor regarding “New cases of recently described Thauvin-Robinet-Faivre syndrome with a novel homozygous FIBP gene variant” by Kılıç and Koşukçu, “An investigation of the etiology and follow-up findings in 35 children with overgrowth syndromes, including biallelic SUZ12 variant” by Yüksel Ülker et al. and “Expanding the phenotype and genotype in Thauvin-Robinet-Faivre syndrome: A new patient with a novel variant and additional clinical findings” by Duzenli et al. *Am J Med Genet A*. 2024;194:e63507. Epub 2023 Dec 15
10. Li H, Durbin R. Fast and accurate short read alignment with Burrows-Wheeler transform. *Bioinformatics*. 2009;25:1754-1760.
11. McKenna A, Hanna M, Banks E, Sivachenko A, Cibulskis K, Kernytzky A, Garimella K, Altshuler D, Gabriel S, Daly M, DePristo MA. The Genome Analysis Toolkit: a MapReduce framework for analyzing next-generation DNA sequencing data. *Genome Res*. 2010;20:1297-1303. Epub 2010 Jul 19
12. McLaren W, Gil L, Hunt SE, Riat HS, Ritchie GR, Thormann A, Flicek P, Cunningham F. The ensembl variant effect predictor. *Genome Biol*. 2016;17:122.
13. Karczewski KJ, Francioli LC, Tiao G, Cummings BB, Alföldi J, Wang Q, Collins RL, Laricchia KM, Ganna A, Birnbaum DP, Gauthier LD, Brand H, Solomonson M, Watts NA, Rhodes D, Singer-Berk M, England EM, Seaby EG, Kosmicki JA, Walters RK, Tashman K, Farjoun Y, Banks E, Poterba T, Wang A, Seed C, Whiffin N, Chong JX, Samocha KE, Pierce-Hoffman E, Zappala Z, O'Donnell-Luria AH, Minikel EV, Weisburd B, Lek M, Ware JS, Vittal C, Armean IM, Bergelson L, Cibulskis K, Connolly KM, Covarrubias M, Donnelly S, Ferriera S, Gabriel S, Gentry J, Gupta N, Jeandet T, Kaplan D, Llanwarne C, Munshi R, Novod S, Petrillo N, Roazen D, Ruano-Rubio V, Saltzman A, Schleicher M, Soto J, Tibbetts K, Tolonen C, Wade G, Talkowski ME; Genome Aggregation Database Consortium; Neale BM, Daly MJ, MacArthur DG. The mutational constraint spectrum quantified from variation in 141,456 humans. *Nature*. 2020;581:434-443. Epub 2020 May 27. Erratum in: *Nature*. 2021;590:E53. Erratum in: *Nature*. 2021;597:E3-E4.
14. Richards S, Aziz N, Bale S, Bick D, Das S, Gastier-Foster J, Grody WW, Hegde M, Lyon E, Spector E, Voelkerding K, Rehms HL; ACMG Laboratory Quality Assurance Committee. Standards and guidelines for the interpretation of sequence variants: a joint consensus recommendation of the American College of Medical Genetics and Genomics and the Association for Molecular Pathology. *Genet Med*. 2015;17:405-424.
15. Touat M, Ileana E, Postel-Vinay S, André F, Soria JC. Targeting FGFR signaling in cancer. *Clin Cancer Res*. 2015;21:2684-2694.
16. Chen F, Hristova K. The physical basis of FGFR3 response to fgf1 and fgf2. *Biochemistry*. 2011;50:8576-8582. Epub 2011 Sep 16.
17. Thauvin-Robinet C, Duplomb-Jego L, Limoge F, Picot D, Masurel A, Terriat B, Champilou C, Minot D, St-Onge J, Kuentz P, Duffourd Y, Thevenon J, Rivière JB, Faivre L. Homozygous FIBP nonsense variant responsible of syndromic overgrowth, with overgrowth, macrocephaly, retinal coloboma and learning disabilities. *Clin Genet*. 2016;89:e1-e4. Epub 2016 Jan 20
18. Akawi N, Ben-Salem S, Lahti L, Partanen J, Ali BR, Al-Gazali L. A recessive syndrome of intellectual disability, moderate overgrowth, and renal dysplasia predisposing to Wilms tumor is caused by a mutation in FIBP gene. *Am J Med Genet A*. 2016;170:2111-2118. Epub 2016 May 17
19. Keppler-Noreuil KM, Sapp JC, Lindhurst MJ, Parker VE, Blumhorst C, Darling T, Tosi LL, Huson SM, Whitehouse RW, Jakkula E, Grant I, Balasubramanian M, Chandler KE, Fraser JL, Gucev Z, Crow YJ, Brennan LM, Clark R, Sellars EA, Pena LD, Krishnamurthy V, Shuen A, Braverman N, Cunningham ML, Sutton VR, Tasic V, Graham JM Jr, Geer J Jr, Henderson A, Semple RK, Biesecker LG. Clinical delineation and natural history of the PIK3CA-related overgrowth spectrum. *Am J Med Genet A*. 2014;164A:1713-1733. Epub 2014 Apr 29
20. Venot Q, Canaud G. Les syndromes de surcroissance segmentaire et les stratégies thérapeutiques [Segmental overgrowth syndromes and therapeutic strategies]. *Med Sci (Paris)*. 2020;36:235-242. Epub 2020 Mar 31
21. Mirzaa G, Graham JM Jr, Keppler-Noreuil K. PIK3CA Related Overgrowth Spectrum. In: Adam MP, Mirzaa GM, Pagon RA, Everman DB, et al., editors. *GeneReviews®* [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2023. [Accessed on: August 25th, 2022]. <https://www.ncbi.nlm.nih.gov/books/>
22. Yehia L, Keel E, Eng C. The clinical spectrum of *PTEN* Mutations. *Annu Rev Med*. 2020;71:103-116. Epub 2019 Aug 21
23. Kinross KM, Montgomery KG, Mangiafico SP, Hare LM, Kleinschmidt M, Bywater MJ, Poulton IJ, Vrahnas C, Henneicke H, Malaterre J, Waring PM, Cullinane C, Sims NA, McArthur GA, Andrikopoulos S, Phillips WA. Ubiquitous expression of the *Pik3ca*H1047R mutation promotes hypoglycemia, hypoinsulinemia, and organomegaly. *FASEB J*. 2015;29:1426-1434.
24. Hughes M, Hao M, Luu M. PIK3CA vascular overgrowth syndromes: an update. *Curr Opin Pediatr*. 2020;32:539-546.
25. Xu Y, Li J, Zhu K, Zeng Y, Chen J, Dong X, Zhang S, Xu S, Wu G. FIBP interacts with transcription factor STAT3 to induce EME1 expression and drive radioresistance in lung adenocarcinoma. *Int J Biol Sci*. 2023;19:3816-3829.
26. Tassi E, Henke RT, Bowden ET, Swift MR, Kodack DP, Kuo AH, Maitra A, Wellstein A. Expression of a fibroblast growth factor-binding protein during the development of adenocarcinoma of the pancreas and colon. *Cancer Res*. 2006;66:1191-1198.
27. Li W, Wang C, Juhn SK, Ondrey FG, Lin J. Expression of fibroblast growth factor binding protein in head and neck cancer. *Arch Otolaryngol Head Neck Surg*. 2009;135:896-901.
28. Colvin JS, Bohne BA, Harding GW, McEwen DG, Ornitz DM. Skeletal overgrowth and deafness in mice lacking fibroblast growth factor receptor 3. *Nat Genet*. 1996;12:390-397.