

Report of the Phenotype of a Patient with Roberts Syndrome and a Rare *ESCO2* Variant

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J Pediatr Genet 2020;9:58–62.

Abstract

Roberts syndrome is a rare autosomal recessive genetic disease. In this report, we report a Brazilian patient with a rare *ESCO2* variant. The patient manifested a broad range of clinical findings including the significant, bilateral shortening of the extremities. He deteriorated and passed away at 20 days of age. High-resolution GTG-banded karyotype showed lack of centromeric constriction in some chromosomes, premature centromere separation in others, and repulsion of the heterochromatin regions. Molecular analysis of the *ESCO2* gene revealed a deletion of 4 bp involving exon 4 in homozygosity (NM_00107420.2:c.875_878delACAG), which causes loss of *ESCO2* function. We describe the clinical presentation caused by a rare *ESCO2* variant.

Keywords

- ▶ Roberts syndrome
- ▶ phocomelia
- ▶ *ESCO2* variant

Introduction

Roberts syndrome (RS) (OMIM #268300) is a rare autosomal recessive genetic disorder, named after the report by John Roberts in 1919.^{1–4} There are earlier reports, however, in 1671 and 1737, which presented characteristics suggestive of RS.^{5,6} Herrmann et al also used, in 1969, the term “SC Phocomelia syndrome” or “thalidomide-like syndrome” (due to similarity of the clinical findings observed in patients exposed to thalidomide during pregnancy) in the description of two European families with clinical findings similar to RS.^{7–10}

Tomkins et al in 1979 described a cytogenetic alteration involving the centromere, which affected most of the chromosomes of RS patients, that was later called “premature

separation of the centromere” by James German.^{11,12} Furthermore, it was found that individuals with RS exhibit a lack of cohesion involving the heterochromatic region of some chromosomes, those around the centromere and in the distal portion of the Y chromosome long arm (called heterochromatin repulsion [HR]). Currently, it is known that RS is caused by biallelic variants in the *ESCO2* gene.³

Around 150 cases of RS have been described in the literature.¹³ The condition is characterized by a broad range of clinical findings with variable expressivity, such as growth retardation (of pre- and postnatal onsets), craniofacial abnormalities (such as microcephaly and cleft palate), symmetrical limb reduction (tetrachomelia), and varying degrees of intellectual disability.^{1,3,5,14} The prognosis of RS is generally

received

March 31, 2019

accepted after revision

July 22, 2019

published online

September 3, 2019

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 Verlag KG, Stuttgart · New York

DOI <https://doi.org/10.1055/s-0039-1696636>
 ISSN 2146-4596.

poor, and survival usually does not exceed the neonatal period.¹⁵

Here, we report a patient with a rare *ESCO2* variant.

Case Report

The patient was the first son of a young, healthy Caucasian couple, not known to be related to each other. There was no similar case in his family. The pregnancy was uneventful. A fetal ultrasound performed in the fifth month of gestation indicated a probable cleft lip and palate, and shortened long bones. The child was born at term by Cesarean section weighing 1,570 g, measuring 34 cm, with head circumference of 26 cm, and Apgar scores of 9 and 10 in the first and fifth minutes, respectively.

Physical examination on the second day of life evidenced a wide anterior fontanelle (extended from the base of frontal bone to the superior occipital region); orbital hypertelorism; blepharophimosis; shallow orbits with exophthalmos; a central facial cleft involving the nose, lip, and palate with the presence of a nasal rudiment; and small, low-set, and posteriorly rotated ears. The extremities exhibited dramatic bilateral shortening of humerus and apparent bilateral absence of forearms with small hands, absence of the left thumb, and presence of an appendicular thumb on the right. He had bilateral shortening of thighs and legs, with restricted knee movement, syndactyly between fourth and fifth toes. Hypertrichosis was noted in the sacral regions and thighs, as well as a hypertonic posture with neck extension. He had normal male genitalia (►Fig. 1).

Radiographic evaluation revealed the presence of marked deformity of the skull with reduction of its anteroposterior diameter; eleven ribs; shortened and deformed humerus; absent radius; right hand presenting two metacarpal bones with normal appearance, two fused metacarpals and four fingers with phalanges; left hand with three metacarpal bones

and four fingers with phalanges; femurs were bent, and tibia and fibula were absent bilaterally (►Fig. 2). Brain and abdominal ultrasound examinations were normal. Cardiological evaluation showed interventricular communication and tricuspid regurgitation. The patient was deteriorated and died at 20 days of age.

High-resolution GTG-banded karyotype made with peripheral lymphocytes showed a male chromosomal constitution (46,XY). There was also lack of centromeric constriction in some chromosomes, premature centromere separation (PCS) in others, and regions of HR (most evident on chromosomes 1, 9, 16, and Y). The total number of scored cells was 32. The percentage displaying PCS/HR phenotype was 100%, and the number of chromosomes with PCS/HR per cell was 8 (►Fig. 3). Molecular analysis from DNA extracted from peripheral blood of the patient through sequencing of all 10 coding exons and splice junctions of the *ESCO2* gene using BigDye Terminator chemistry and an ABI 3100 sequencer (Applied Biosystems, Foster City, California, United States) revealed a deletion of 4 bp involving exon 4 in homozygosity (NM_001017420.2:c.875_878delA-CAG, rs80359856) (►Fig. 4). Both parents were heterozygotes for this variant. This variant causes a frameshift and premature stop codon. Because of nonsense-mediated decay of the mutant mRNA, the predicted truncated protein p. (Asp292Glufs) is unlikely to be produced in significant quantities. A subsequent pregnancy of this couple was monitored, and the fetus was found to be homozygous for the normal allele (►Fig. 4). No other family members were tested for this variant.

Discussion

In our literature review in MEDLINE, we found only one Brazilian study involving individuals with RS. Barbosa et al evaluated the replication patterns of homologous aliphoid



Fig. 1 Patient at 2 days of age. Note orbital hypertelorism and facial cleft involving the nose, lip, and palate with the presence of a nasal rudiment (A); small, low-set, and posteriorly rotated ears; drastic shortening of the arm and forearm along with small hands with absence of thumb on the left side and presence of an appendicular thumb on the right, and remarkable shortening of thighs and legs (B).



Fig. 2 Radiographic evaluation showing 11 ribs and shortened and deformed humerus (A); marked deformity of the skull with reduction of its anteroposterior diameter (B), and femur bending, with absence of tibia and fibula (C).

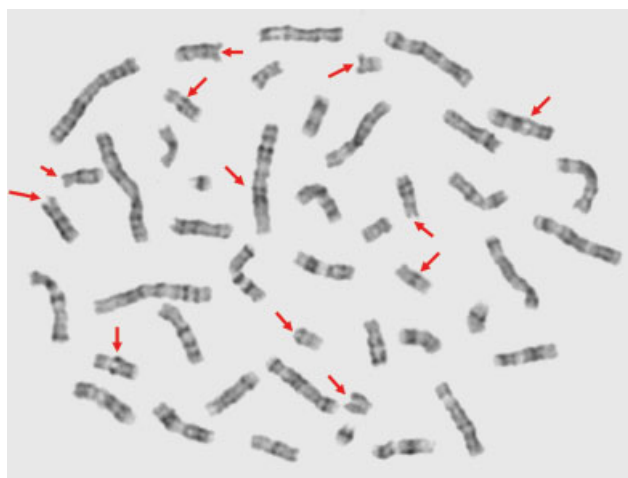


Fig. 3 High-resolution GTG-banded karyotype revealing lack of centromeric constriction in some chromosomes, premature centromere separation in others, and repulsion of the heterochromatin regions (see red arrows).

centromeric DNA of chromosomes 9, 11, 16, and 17 using fluorescence in situ hybridization analysis. They analyzed three patients. However, there was no description of their clinical features.¹⁵ Searching other databases from Latin America, we verified the presence of three studies, one from Brazil and the other two from Cuba and Venezuela. This Brazilian study by Gollop et al claimed that it was the first case of RS described in Brazil.¹⁶ The diagnosis was made by fetal ultrasound during pregnancy and was based on the findings of tetraphocomelia, marked retromicrognathia without lip–palatine fissures or premaxillary protrusions. These findings were confirmed later by autopsy. In both reports, there was no description of additional genetic evaluations, as the gene for RS was discovered only years later. Therefore, our case represents the first molecularly confirmed case of RS in Brazil.

Our patient also presented abnormalities identified through karyotype analysis that consisted of lack of centromeric constriction in some chromosomes, PCS in others, and

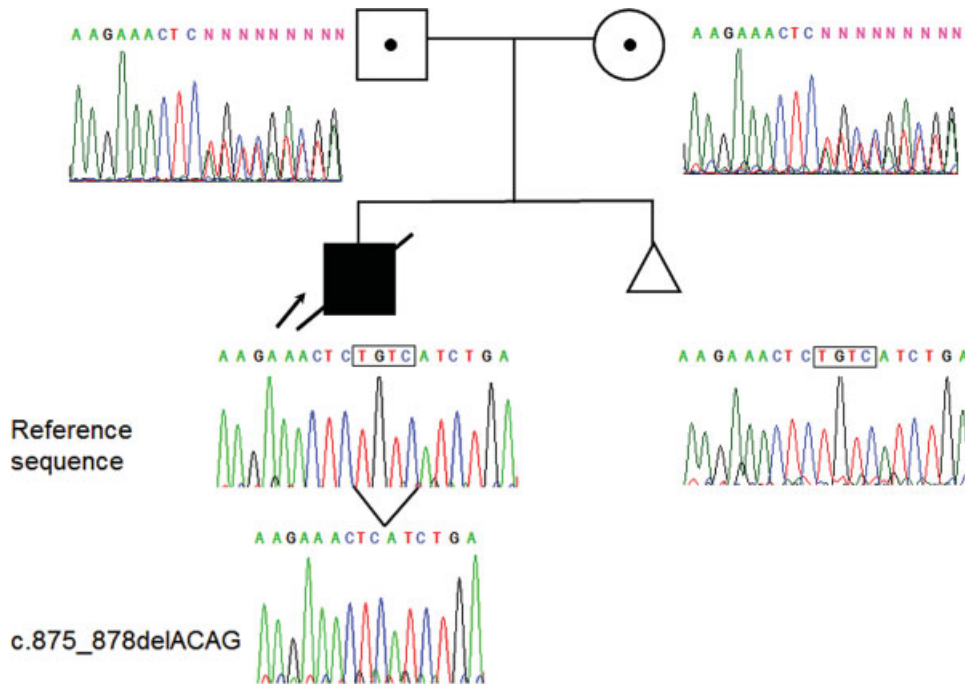


Fig. 4 Family tree with DNA sequence tracings of the reverse strand. Both parents are heterozygous for the deletion. The proband is homozygous for the c.875_878delACAG deletion. A subsequent pregnancy was monitored prenatally, and the fetus was homozygous for the normal allele (identical to the reference sequence shown).

repulsion of heterochromatin regions. These are considered pathognomonic findings for the diagnosis of RS.² These cytogenetic findings are described in 79.1% of the patients with RS.³ Studies have suggested that the lack of fusion of heterochromatin in RS would be caused by the variant in *ESCO2*. This ensures that the cohesin ring around the sister chromatids establishes cohesion during the S phase of replication along with *ESCO1*.^{2,3,10,16-19}

As for prenatal diagnosis, we observed, in our case, the ultrasonographic description in the fifth month of gestation of a probable cleft lip and palate. Moreover, shortening of the long bones was evident. Literature reports of prenatal diagnosis of RS most frequently describe reduced fetal movements, restriction of fetal growth, and shortening of the long bones.^{2,5,9,12,18,20-22} In the case of prenatal diagnosis reported by Gollop et al, those findings were observed during a standard ultrasound at 21 weeks of gestation.²³

The clinical manifestations observed in our patient did not differ from the majority of RS cases described in the literature with RS (►Table 1).²² This suggests that RS is not clinically variable. Few patients with RS also survive for more than 1 month, which is compatible with the clinical evolution observed in our case.²²

The *ESCO2* variant identified in our patient was described once before by Gordillo et al in a French family, although no clinical data were provided. This variant is not reported in the Genome Aggregation Database (gnomAD).²⁴ In ClinVar, it is described once.²⁵ In the Exome Aggregation Consortium (ExAC), the variant allele was also reported only once, in the Asian population (minor allele frequency: 8.299e-06).²⁶ According to the College of Medical Genetics and Genomics and the Association for Molecular Pathology Standards and

Table 1 Clinical features of Roberts syndrome described in the literature in comparison with our patient (based on Van Den Berg and Francke²²)

Clinical features	Literature %	Our patient
Consanguinity	40.8	—
Sex	54 M/39 F/7 U	M
Neurological		
Mental retardation	79.4	X
Growth retardation	97.5	+
Craniofacial anomalies		
Microcephaly	80.4	
Hemangioma	29	
Exophthalmia/prominent eyes	69.4	+
Hypertelorism	86.7	+
Cloudy cornea	68.1	
Hypoplastic nostrils	32	
Cleft palate	60.6	+
Cleft lip	49	+
Highly arched palate	16	
Prominent maxilla	30	
Micrognathia	13	
Abnormal ears	75.9	+

(Continued)

Table 1 (Continued)

Clinical features	Literature %	Our patient
Limb anomalies		
Phocomelia	100	+
Ulnar aplasia	96.5	+
Radial aplasia	97.8	+
Femoral aplasia	64.9	
Tibial aplasia	74.1	+
Fibular aplasia	80.5	+
Leg bone synostosis	7	
Syndactyly	42.5	+
Flexion contractures	51	+
Other anomalies		
Heart defects	13	+
Kidney anomalies	50	
Enlarged phallus	53.2	+
Cryptorchidism	51.2	

Abbreviations: F, female; M, male; U, undetermined sex; X, unknown.

Guidelines, the variant was classified as pathogenic based on the following criteria: causing a frameshift of extremely low frequency in ExAC and not described in the 1000 Genomes project; the protein length changes (normal protein: 602 aa; mutated protein: 339 aa); and other support computational evidences that the variant is deleterious, and it has a relationship with the observed phenotype.²⁷

Thus, the variant found in our patient is considered a rare *ESCO2* variant, whose phenotype was not previously described. This variant causes loss of function of *ESCO2*, and the clinical findings and clinical course of our patient were typical for RS cases. This represents the first Brazilian report of a patient with RS describing his clinical features.

Conflict of Interest

None declared.

References

- McDaniel LD, Prueitt R, Probst LC, et al. Novel assay for Roberts syndrome assigns variable phenotypes to one complementation group. *Am J Med Genet* 2000;93(03):223–229
- Schüle B, Oviedo A, Johnston K, Pai S, Francke U. Inactivating mutations in *ESCO2* cause SC phocomelia and Roberts syndrome: no phenotype-genotype correlation. *Am J Hum Genet* 2005;77(06):1117–1128
- Vega H, Waisfisz Q, Gordillo M, et al. Roberts syndrome is caused by mutations in *ESCO2*, a human homolog of yeast *ECO1* that is essential for the establishment of sister chromatid cohesion. *Nat Genet* 2005;37(05):468–470
- Roberts JB. A child with double cleft lip and palate, protrusion of the intermaxillary portion of the upper jaw and imperfect development of the bones of the four extremities. *Ann Surg* 1919;70:252–254
- Bates AW. A case of Roberts syndrome described in 1737. *J Med Genet* 2001;38(08):565–567

- Oostra RJ, Baljet B, Dijkstra PF, Hennekam RC. Congenital anomalies in the teratological collection of Museum Vrolijk in Amsterdam, The Netherlands. I: syndromes with multiple congenital anomalies. *Am J Med Genet* 1998;77(02):100–115
- Herrmann J, Feingold M, Tuffli GA, Opitz JM. A familial dysmorphic syndrome of limb deformities, characteristic facial appearance and associated anomalies: the pseudothalidomide or SC-syndrome. *Birth Defects Orig Art Ser* 1969;5:81–89
- Herrmann J, Opitz JM. The SC phocomelia and the Roberts syndrome: nosologic aspects. *Eur J Pediatr* 1977;125(02):117–134
- Al Kaissi A, Csepan R, Klaushofer K, Grill F. Femoral-tibial-synostosis in a child with Roberts syndrome (pseudothalidomide): a case report. *Cases J* 2008;1(01):109
- Sánchez-Segura M, Marsán-Suárez V, Macías-Abraham C, et al. Roberts syndrome associated with immunodeficiency. *Rev Cubana Hematol Inmunol Hemoter* 2012;28(02):185–191
- Tomkins D, Hunter A, Roberts M. Cytogenetic findings in Roberts-SC phocomelia syndrome(s). *Am J Med Genet* 1979;4(01):17–26
- Schulz S, Gerloff C, Ledig S, et al. Prenatal diagnosis of Roberts syndrome and detection of an *ESCO2* frameshift mutation in a Pakistani family. *Prenat Diagn* 2008;28(01):42–45
- Zhou J, Yang X, Jin X, Jia Z, Lu H, Qi Z. Long-term survival after corrective surgeries in two patients with severe deformities due to Roberts syndrome: a case report and review of the literature. *Exp Ther Med* 2018;15(02):1702–1711
- Gordillo M, Vega H, Trainer AH, et al. The molecular mechanism underlying Roberts syndrome involves loss of *ESCO2* acetyltransferase activity. *Hum Mol Genet* 2008;17(14):2172–2180
- Barbosa AC, Otto PA, Vianna-Morgante AM. Replication timing of homologous alpha-satellite DNA in Roberts syndrome. *Chromosome Res* 2000;8(07):645–650
- German J. Roberts' syndrome. I. cytological evidence for a disturbance in chromatid pairing. *Clin Genet* 1979;16(06):441–447
- Louie E, German J. Roberts's syndrome. II. Aberrant Y-chromosome behavior. *Clin Genet* 1981;19(01):71–74
- Vega H, Trainer AH, Gordillo M, et al. Phenotypic variability in 49 cases of *ESCO2* mutations, including novel missense and codon deletion in the acetyltransferase domain, correlates with *ESCO2* expression and establishes the clinical criteria for Roberts syndrome. *J Med Genet* 2010;47(01):30–37
- Goh ES-Y, Li C, Horsburgh S, Kasai Y, Kolomietz E, Morel CF. The Roberts syndrome/SC phocomelia spectrum—a case report of an adult with review of the literature. *Am J Med Genet A* 2010;152A(02):472–478
- Petrinelli P, Antonelli A, Marcucci L, Dallapiccola B. Premature centromere splitting in a presumptive mild form of Roberts syndrome. *Hum Genet* 1984;66(01):96–99
- Parry DM, Mulvihill JJ, Tsai SE, Kaiser-Kupfer MI, Cowan JM. SC phocomelia syndrome, premature centromere separation, and congenital cranial nerve paralysis in two sisters, one with malignant melanoma. *Am J Med Genet* 1986;24(04):653–672
- Van Den Berg DJ, Francke U. Roberts syndrome: a review of 100 cases and a new rating system for severity. *Am J Med Genet* 1993;47(07):1104–1123
- Gollop TR, Eigier A, Hauschild D, Guidugli J, Moron AF. Prenatal ultrasound diagnosis of Roberts syndrome at 21 weeks. *Braz J Genet* 1990;13(03):607–612
- The Genome Aggregation Database (gnomAD). Retrieved from <https://gnomad.broadinstitute.org/>. Accessed April 25, 2019
- ClinVar. Retrieved from <https://www.ncbi.nlm.nih.gov/clinvar/>. Accessed September 4, 2018
- ExAC Browser (Beta)|Exome Aggregation Consortium. Retrieved from <http://exac.broadinstitute.org/>. Accessed September 4, 2018
- Richards S, Aziz N, Bale S, et al; 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(05):405–424