

Rare case of Fanconi anemia associated with polydactyly

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Introduction

Fanconi anemia (FA) is a rare progressive congenital bone marrow failure syndrome. It is inherited in an autosomal recessive manner and is associated with a constellation of clinical findings. Lymphocytes and skin fibroblasts in patients with FA are characterized by hypersensitivity to DNA-cross linking agents such as Diepoxybutane (DEB) and Mitomycin C (Chromosome Breakage Test) resulting in chromosomal aberrations. Mutations in at least 20 genes have been associated with FA. Most frequent gene mutations in FA patients worldwide is FANCA which account for 60- 65% of the patients.

The epidemiology and frequency of different subtypes of FA in our population are not well known. In one study of the molecular patterns of FA in ten Saudi patients showed that BRIP1 (FANCI) was the most frequent compared to only 2% in Europeans. In Saudi Arabia, FA is one of the most frequent indications for bone marrow transplantation among patients with inherited bone marrow failure syndromes [1]. Fanconi anemia can present with skeletal anomalies, bone marrow failure, and increase in the risk of malignancy. In the first decade of life, FA patients usually present with pancytopenia caused by progressive bone marrow failure [2].

Case report

A four years old Saudi girl with a history of failure to thrive, pancytopenia for six months presented with fevers for two days as well as vomiting for one day. She was admitted as a case of pancytopenia for investigations. On physical examination, her height was 92 cm (<5th %) consistent with short stature. Her weight was 12.8 kg (<5th %). She was also found to have bilateral microcoria as well as partial ptosis. She has hyperpigmented lesions (Café-au-lait spots). She has also been noted to have polydactyly in the right hand. She has a small extra digit next to her right thumb. Other systemic examination was unremarkable.

Laboratory investigations showed Hemoglobin of 4.2 g/fl. White cell count 1.9. Platelets count of 6. Neutrophil count was 0.4 (Absolute Neutrophil Count was 400). Renal function tests and liver enzymes were all normal. She received packed red blood transfusions and platelets transfusion. Subsequently, she was admitted on average every 2-3 weeks with pancytopenia requiring blood and platelets transfusion. She also had episodes of severe febrile neutropenia requiring admission and intravenous (IV) antibiotics.

She also had abdominal and renal ultrasound imaging which showed left ectopic kidney. Echocardiogram was done and it was unremarkable.

On 29/2/2017 bone marrow aspiration and biopsy was performed. It showed significantly hypocellular marrow with all three cell lines affected. There was no fibrosis and no blasts.

We also sent for Chromosome Breakage Tests on peripheral lymphocytes using Mitomycin C and Diepoxybutane twice and were both normal. In light of the strong clinical suspicion of Fanconi anemia, we decided to send for FA gene panel. It showed homozygous mutation in Fanconi anemia G subtype (FANCI-G). Therefore, she was confirmed to have as Fanconi anemia.

Discussion

Fanconi anemia is the most common cause of inherited bone marrow failure syndromes. It is characterized by genomic instability which leads to the observed clinical manifestations. Fanconi anemia proteins are involved in cell division checkpoints and DNA repair mechanisms. They are responsible for maintaining genomic stability and cell division integrity. Defects in these genes lead to chromosomal breakage, cell cycle disturbances and increased rate of somatic mutations. Clinically, it manifests commonly as growth retardation, bone marrow failure causing pancytopenia, increased risk of malignancy, skin pigmentation, and skeletal malformations. Less commonly, it may affect eyes, gastrointestinal tract, genitourinary tract, heart, oral cavity, and central nervous system. Up to twenty FA genes have been identified so far. The more common genes are FANCA, FANCB, FANCI, FANCD1 (BRCA2), FANCD2, FANCE, FANCF, FANCG, FANCI, and FANCI (BRIP1). The less common FA genes include; FANCL, FANCM, FANCN (PALB2), FANCO (RAD51C), FANCP (SLX4), FANCI (ERCC4), FANCI (RAD51), FANCI (UBE2T), FANCI (XRCC2), and FANCI (MAD2L2) [3,4].

FANCI-G has been localized to the short arm of chromosome p923. It was cloned by Liu et al in 1997. It was also observed that cells from patients with FANCI-G show resistance to Mitomycin C [5]. Later, Van der Heijden *et al.* in 2003 showed that both somatic and inherited FANCI-G mutations are associated with young-onset pancreatic cancer [6].

The age of onset of hematologic abnormalities may vary with the mean age reported around 8 years. An early diagnosis of FA is important as an initial step to assess the need for Hematopoietic stem cell transplant which is ideally performed before the need for more frequent regular blood transfusions [7]. In our case the patient was discovered early at 4 years of age which was an advantage to initiate treatment. She presented with classical signs of a fanconi anemia; short stature, suboptimal weight, and pancytopenia. She also has bilateral

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microcoria with partial ptosis and polydactyly in the right hand. Also, she has hypoplasia of the thumb of the other hand. After analyzing the data of 700 patients with FA in the literature, almost 300 patients had thumb anomalies such as an absent thumb or hypoplastic thumb, floating or bifid thumb, polydactyly, as well as triphalangeal thumbs. The association of thumb abnormalities and FA has been well established [8,9].

After BM aspiration was done it showed decrease in all cell lines. Then gene studies showed FANCG gene mutation in our patient. This gene mutation affects about 10% of all FA cases compared to the more detected mutation which is FANCA. FANCA is the most frequent mutation in patients with fanconi anemia accounting for about 70% of FA cases [10].

Patients with FANC-G gene mutation had more severe pancytopenia and a higher incidence of malignancy. Our case started to have pancytopenia at the age of 6 months which required frequent monitoring and early intervention to avoid the poor outcome [11].

Conclusions

In conclusion, this is a rare presentation for a FA patient having polydactyly and thumb hypoplasia, with a non-common subtype of gene mutation FANCG was detected early, such cases must be reported from Saudi Arabia, to avoid the poor hematologic outcome and to identify the etiology of this association between specific subtypes of gene mutations and their outcome, further genetic mutation analysis for FA is needed.

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