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## Myelodysplastic Syndrome, Acute Myeloid Leukemia, and Cancer Surveillance in Fanconi Anemia

## Sharon A. Savage, MD<sup>1</sup> and Michael Walsh, MD, FAAP, FACMG, DABMG<sup>2</sup>

<sup>1</sup>Chief, Clinical Genetics Branch Division of Cancer Epidemiology and Genetics, National Cancer Institute

<sup>2</sup>Assistant Member, Departments of Medicine and Pediatrics, Divisions of Solid Tumor and Clinical Cancer Genetics, Memorial Sloan Kettering Cancer Center

## Synopsis

Fanconi anemia (FA) is a DNA repair disorder associated with high risk of cancer and bone marrow failure. Patients with FA may present with certain dysmorphic features such as radial ray abnormalities, short stature, typical facies, bone marrow failure, or certain solid malignancies. Some patients may be recognized due to exquisite sensitivity after exposure to cancer therapy. FA is diagnosed by increased chromosomal breakage after exposure to clastogenic agents. It follows both autosomal recessive and X-linked inheritance depending on the underlying genomic alteration(s). Nineteen genes encode proteins comprising the Fanconi DNA damage response complex. To date, pathogenic variants in at least 15 FA pathway genes and at least 5 related genes have been identified as causative of FA. Recognizing patients with FA is important for therapeutic decisions, genetic counseling, and optimal clinical management.

## Keywords

Bone Marrow Failure; Myelodysplastic Syndrome; Leukemia; Exquisite Therapeutic Sensitivity; DNA repair; BRCA1/2; Fanconi complex; Fanconi anemia; head and neck squamous cell cancer

## INTRODUCTION TO FANCONI ANEMIA

Fanconi anemia (FA, MIM 607139) is a rare, cancer-prone inherited bone marrow failure syndrome with a wide range of clinical presentations including radial ray anomalies, short stature, microcephaly, café au lait spots, and other medical problems. The condition is eponymous for Dr. Guido Fanconi, who originally described the syndrome in 1927. At present there are more than 2000 patients reported in the literature.(1) Advances in supportive care including hematopoietic cell transplantation (HCT) have improved the lifespan for patients with FA but cancer, HCT-related complications, and other complex

Corresponding Authors: Sharon A. Savage, 9609 Medical Center Dr., Rm. 6E456, MSC 9772, Bethesda, MD 20892-9772, savagesh@mail.nih.gov; Michael Walsh, 222 70th Street Rm 412, NY, NY, 10021, walshm2@mskcc.org.

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medical problems remain as significant causes of mortality. Herein we review the underlying biology and clinical manifestations of FA as well as the current recommendations for cancer surveillance and management.

#### The Biology of Fanconi Anemia

FA is a chromosomal instability disorder caused predominantly by autosomal recessive inheritance of pathogenic variants in key components of the DNA damage response(2, 3). There is one gene, FANCB, associated with X-linked recessive inheritance (4). Germline mutations (i.e., pathogenic variants) in at least 22 genes are associated with FA. There is consensus in the field regarding 18 genes, (FANCA, B. C. D1, D2, E. F. G. I. J. L. N. P. O. T, U, V and W) (4–26) (Table 1) (27). An additional four genes (FANCM, FANCO, FANCR and FANCS) are considered FA-like, as they have not been described in patients with bone marrow failure (25, 28, 29). FANCM is not considered a bonafide FA gene, as only it has only occurred in a patient also reported to also harbor bi-allelic FANCA mutations. Biallelic pathogenic variants in FANCA are the most common cause of FA, with 65% of patients harboring mutations in this gene. This is followed by FANCC(14%) and FANCG(9%), with the remaining 14% covered by the other genes (30). (Table 1). The proteins of the FA pathway create a biochemical circuit that function in DNA repair, DNA damage response, and other cellular processes. As DNA is replicated, nucleotide incorporation and processing of the replication fork are prone to errors including wrong nucleotides, damaged bases within the DNA, abnormal DNA-protein complexes, creation of DNA-RNA hybrids (Rloops) and aberrant DNA structures, such as G quadraplexes (31). The specific role of proteins in the Fanconi pathway is removal of DNA interstrand cross-links (ICLs). Interstrand cross-links may arise from endogenous and exogenous compounds such as aldehydes and platinum drugs, respectively. Interstrand cross-links prevent DNA strand separation and can act to block the DNA replication process and/or transcription. These aberrant DNA strands, if left intact, promote cell death (32).

#### **Clinical Manifestations**

Clinical characteristics of FA may vary significantly from patient to patient. Typical FA findings include radial ray abnormalities with missing or unusual thumbs, café au lait macules, and typical "Fanconi" facies. Abnormal thumbs are an important differentiating feature of FA and differentiate this syndrome from thrombocytopenia absent radius syndrome (TAR) in which thumbs are present(33). Other significant FA-associated anomalies include: kidney and urinary tract malformations, vertebral anomalies, esophageal atresia, hydrocephalus, short stature, and small eyes. Almost all organ systems may be involved with the highest number of congenital malformations associated with the FANCD1/ BRCA2 genotype(34). Clinical features of FA may significantly overlap with those of the VACTERL-H association: VACTERL-H is the acronym for vertebral anomalies, anal atresia, cardiac anomalies, trachea-esophageal fistula, esophageal or duodenal atresia, renal structural anomalies, limb deformities and hydrocephalus(35). Additional manifestations of FA may include metabolic abnormalities, endocrinopathies, and hearing impairments. While patients with FA may exhibit the VACTERL-H phenotype, it is not specific to FA. This led Alter and Giri to propose PHENOS (Pigmentation, small Head, small Eyes, central Nervous system [not hydrocephalus], Otology, and Short stature) as an acronym that includes the

Patients with FA are at very high risk of bone marrow failure (BMF), myelodysplastic syndrome (MDS), acute myeloid leukemia (AML), head and neck squamous cell carcinoma (HNSCC), and other malignancies. In some patients, BMF or cancer may be the first presenting sign of FA. It should be suspected in patients who have higher than expected toxicities related to cancer treatment.

#### **Diagnosing Fanconi Anemia**

FA is diagnosed by chromosome breakage studies on patient-derived cells. In this assay, the patient's peripheral blood T lymphocytes are exposed to diepoxybutane or mitomycin C and the number of resultant abnormal chromosomes are counted. False negatives may occur in samples obtained from individuals with hematopoietic somatic mosaicism and in such instances testing of skin fibroblasts is required to diagnose FA (37).

Genetic testing for FA now includes multiplex (*i.e.*, gene panel) testing, as well as whole exome or whole genome sequencing. Identification of the causative pathogenic variants is helpful in confirming the diagnosis of FA and essential for genetic counseling of the patient and their family members. The availability of multiplex genetic testing has led to widespread testing of patients in whom FA may not have been considered. In such instances, chromosomal breakage studies have been particularly important in confirming whether the genetic variants are contributory to disease.

## CONSEQUENSES AND MANAGEMENT OF HEMATOPOIETIC STEM CELL ATTRITION

#### Bone marrow failure and leukemia

BMF in FA is attributed to attrition of the hematopoietic stem and progenitor cell compartment due to elevated DNA damage response and apoptosis.(38) The bone marrow of patients with FA has been shown to be proinflammatory; specifically, TNF- $\alpha$  may contribute to apoptosis of FA hematopoietic stem cells (HSCs) through activation of the extrinsic apoptotic pathway.(38) Moreover, functional studies have illustrated that aberrantly activated oligoclonal T-cell populations suppress hematopoiesis by releasing cytokines (including INF- $\gamma$  and TNF- $\alpha$ ). (39, 40) This cascade of events is cytotoxic to HSCs and subsequently to lymphoid populations. (41) This is thought to result in a stem cell compartment favoring the evolution of somatically mutated cells with the capacity to resist the attack of T-cells and the evolution of covert leukemic clones (Figure 1) (42–45).

In addition to adverse effects of INF- $\gamma$ , TNF- $\alpha$  and other cytokines in FA patients, there is evidence that BMF may be exacerbated by endogenous aldehyde-induced toxicity and /or DNA damage-induced p53 activation, both of which result in HSC attrition.(46, 47) One report found that FA patients with *ALDH2* mutations have accelerated progression of bone marrow failure and murine models harboring *ALDH2* mutations independent of FA genetics have exhibited bone marrow failure. (48)

In the absence of typical congenital anomalies, cytopenias may be a presenting sign of FA. Additionally, a clue to an IBMFS may also be elevated hemoglobin F and high mean corpuscular volume for age without another cause. Most patients with FA manifest clinically significant cytopenias during their lifetimes. The cumulative incidence of severe BMF of 70% by age 50 and a peak hazard rate of 4% at 12 to 15 years of age (49). The age of BMF onset varies amongst patients with FA, even among siblings. (50)

Patients with FA and BMF should be monitored for the development of MDS and progression to AML by following both cytogenetic and morphologic abnormalities. Long term studies of FA have shown that the risk of developing MDS or AML within three years of observing an abnormal clone was roughly 1 in 3 (35%) compared to 1 in 30 (3.3%) for those without an identifiable clone.(49, 51) Cytogenetic findings in the bone marrow samples of FA patients may reveal recurring chromosomal changes of 1, 3 and 7 and these findings need to be followed closely given the risk of AML in FA. A notable finding in bone marrow of FA patients is the presence of aberrations of chromosome 3 (52, 53). In these patients with identifiable chromosome 3 abnormalities, the 3-year risk of MDS/AML was 90% and 17% respectively. In general, the classification scheme for AML is evolving and becoming more molecularly centric.(54) How applicable this evolving schema will be to FA patients is still to be determined and needs to be further analyzed in larger numbers of FA patients.

MDS and AML occur in a significant percentage of patients with Fanconi anemia. (51) In a recent report from Alter et al., interrogating the National Cancer Institute's registry between 1945-2014, one hundred and thirty families have been reported to date with FA, with approximately one hundred and sixty individuals who are presently living. Leukemia had a cumulative incidence of 5% in Fanconi anemia by the age of 30. The cumulative incidence of MDS in FA patients by the age of 50 was 50% (CI 35-65%).(49)

Of note, AML is significantly more common in FA than acute lymphoblastic leukemia and lymphomas. The latter hematopoietic malignancies have been reported in patients with FA harboring *FANCD1/BRCA2* gene mutations (51, 52, 55–57).

#### **Clinical management**

In general, patients with BMF require treatment for when BMF becomes severe, which is generally defined as hemoglobin <8 g/dL, platelets <30,000/mm<sup>3</sup>, absolute neutrophil count <500/mm<sup>3</sup>.(2) The decision to undergo hematopoietic cell transplant (HCT) is a serious one, which requires a multidisciplinary approach and assessment. The complexities of decision making in the context of FA-associated co-morbidities led to the development of an interesting mathematical decision model to estimate event free-survival (EES) conditional on age and per-year cause specific hazard rates, based on the assumption that bone marrow transplant eliminates the risk of bone marrow failure and acute myeloid leukemia but not the risk of solid tumors (2, 49, 58). This approach provides perspective in deciding when or if a patient with FA should undergo HCT and suggests that transplantation at a younger age offers a benefit that is not as clear in older patients.

Non-myeloablative HCT conditioning regimens are recommended for patients with FA undergoing HCT for BMF without MDS or AML due to their exquisite sensitivity to chemotherapy and ionizing radiation(56, 59–64). Matched sibling donors are recommended, when available, and mutation identification and testing is essential in this context. It is important for patients and their families to understand that HCT can cure BMF, but it does not cure non-hematopoietic complications of FA (49).

All patients with FA in whom HCT is considered should be evaluated and treated at a center specializing in FA(2). This is important to ensure the optimal supportive care and clinical expertise in the management of the complex FA co-morbidities. Patients with FA who evolved to MDS or AML require a highly specialized approach due to their sensitivity to standard cancer therapeutics (2, 65).

Androgens are an option for BMF in patients with FA who cannot undergo HCT (66, 67). There are multiple well recognized side effects of androgens including but not limited to: virilization, premature closure of growth plates, behavioral changes, elevated liver enzymes, hepatic adenomas, hepatocellular carcinoma, peliosis hepatis, acne and hypertension. Half to three-quarters of patients respond to androgens within three months. Androgens do not prevent progression to AML, which once developed, may increase the risks associated with transplant and when patients are older additional viral infections may have been acquired which may impact transplant outcome.(68)

## SOLID MALIGNANCIES

FA patients are at very high risk of solid tumors. In the NCI's prospective longitudinal cohort study of 130 families including 163 patients, 21/163 (12%) patients were diagnosed with solid tumors: HNSCC n=10, vulva n=3, esophagus n=2, brain n=2, anus n=1, lung n=1, cervix n=1 and breast n=1 and an additional 35/163 (21%) were diagnosed including squamous cell cancer (SCC) n=23 and basal cell cancer (BCC) n=12.(49) The observed to expected ratio, for any malignancy in non-transplanted patients for FA was nineteen. FA patients having undergone hematopoietic stem cell transplant, n=63/163, the frequency of solid tumors was higher at 33 (51%): HNSCC n=5, larynx n=1, vulva n=2, brain n=1, thyroid n=2, BCC n=9 and SCC n=12. The effect of HSCT on increased cancer rates is reflected in the higher observed to expected ratios for transplanted vs. the non-transplanted FA patients 55 versus 19. (49) According to Alter et al. the cumulative incidence of solid tumors reported in FA patients is roughly 20% by the age of 65. As FA patients exhibit extreme sensitivity to alkylating and radiation therapies, there continue to be significant challenges in treating patients and avoiding second malignancies.

## CANCER SCREENING AND RISK REDUCTION

The guidelines for diagnosis management of FA are described in detail at http://fanconi.org/ index.php/publications/guidelines. Fanconi anemia patient require multidisciplinary care and self-directed screening. Proper health maintenance includes coordination of care with ear, nose and throat (ENT) physicians, hematologists, oncologists, gastroenterologists,

orthopedic surgeons, gynecologists (females), endocrinologists, dermatologists, dentists and pediatric developmental physicians.

HNSCC is the most common solid tumor in FA patients. Approximately 1 in 7 (14%) of patients with FA who survive to the age 40 will be diagnosed with HNSCC during their lifetime.(49, 51) Surveillance should begin at age 10 years, which is based on literature reports of the earliest age of diagnosis with head and neck cancer. The sites at risk for the development of HNSCC include all areas of the upper aerodigestive tract. Therefore, all mucosal surfaces of the head and neck should be examined. This exam should also include the proximal and distal oropharynx and should include appropriate equipment such as a transoral mirror and flexible fiberoptic laryngoscope. This exam should occur every six months and suspicious lesions should be appropriately treated and screening should increase to every 2-3 months Patients need to be diligent and perform self-exam bringing any suspicious findings to their physicians' attention. (http://fanconi.org/index.php/publications/guidelines\_for\_diagnosis\_and\_management).

Fanconi patients are at increased risk of skin cancers, thus individuals should perform routine skin checks for suspicious lesions following the ABCD acronym for nevus inspection (asymmetry, bleeding/border changes, color changes and difference in diameter). Patient should also protect their skin with hats, clothing and sunscreen. FA patients require application of sunscreen with an SPF of at least 30 or higher. Individuals should see their dermatologist once a year and strive to maintain adequate vitamin D levels through supplements if necessary.

In regard to hematologic monitoring, most physicians agree that a complete blood count (CBC) and bone marrow aspirate and biopsy are recommended at diagnosis. The bone marrow evaluation should be repeated annually. The CBC should be monitored more frequently for proactive management of cytopenias and myelodysplastic syndrome (Figure 2) (65)

## SUMMARY AND DISCUSSION

Fanconi anemia is a DNA damage syndrome. FA in inherited in an autosomal recessive inheritance disease except for patients with *FANCB* mutations, in which case the disease is X-linked. Multidisciplinary care for Fanconi anemia patients is required.(65) The field's understanding of FA continues to evolve with greater than 20 genes described to play a role in Fanconi biology, a description of the environmental milieu of FA patient's bone marrow, tailored non-myeloablative bone marrow transplant regimens and candidate therapeutics.

At present HCT continues to be the only curative treatment for hematopoietic disease in FA patients.

Chromosomal breakage studies are the gold standard testing assay and there is a continued need for genotypic/phenotypic description and studies. Triple assessment including clinical characteristics, chromosomal breakage studies and molecular testing provide the greatest context for advancing our understanding of FA and better understanding atypical findings.

Studies need to be powered well in order to make valid conclusions from integrated genotype/phenotype analyses. Moreover, uniform annotation of variant calling following rigorous classification guidelines will allow for meaningful clinical translation. At present, identification of FANC complex carriers has broad clinical relevance depending on the specific gene and variant detected. In some instances, such as *FANCD1, PALB2, RAD51C, RAD51, and BRIP1* heterozygous carriers are at risk for adult onset cancers but the extent to which heterozygous carriers of pathogenic variants in other FA genes are at risk is not known and is an area of active study. Carriers should be counseled for reproductive risks as well.(69–71) The continued international effort focused on comprehensive care for Fanconi anemia patients is crucial given the rarity of this disease and for further advancements in caring for patients to be made.

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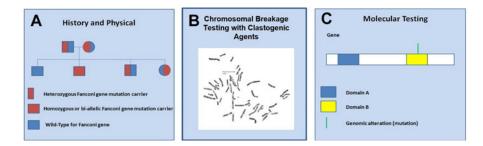
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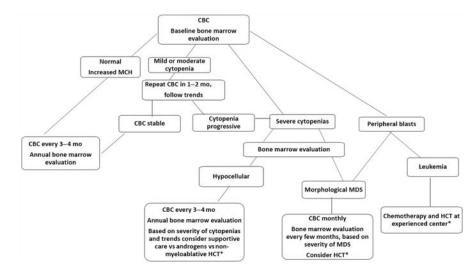
## Key Points (3–5)

- Fanconi anemia is a DNA damage repair syndrome caused by pathogenic variants in key components of the Fanconi DNA repair complex.
- Patients with Fanconi anemia are at very high risk of bone marrow failure, myelodysplastic syndrome, leukemia, head and neck squamous cell carcinoma, and other malignancies.
- Treatment for patients with Fanconi anemia and cancer must be carefully tailored due to exquisite sensitivity to ionizing radiation and alkylating drugs.



## Figure 1.

Fanconi Triple Assessment: Patients being evaluated for Fanconi anemia require a triple assessment including: (A) history/physical, (B) clastogenic testing or chromosomal breakage testing with diepoxybutane +/- mitomycin C, and (C) genetic testing.



#### Figure 2.

Screening for hematologic abnormalities in Fanconi anemia.

^ Persistent drop or rise in blood counts without apparent cause warrants bone marrow evaluation

^^ Specific clonal abnormalities may warrant immediate treatment intervention or closer monitoring

*Adapted from* Alter BP, Hanenberg H, Kinsey S, et al. Hematologic Abnormalities in Patients with Fanconi Anemia. In: Hays L, Frohnmayer D, Frohnmayer L, et al., editors. Fanconi Anemia: Guidelines for Diagnosis and Management, 4<sup>th</sup> edition. Eugene: Fanconi Anemia Research Fund, Inc.: 2014; with permission.

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The Biology of Fanconi Anemia

Fanconi Genes	Inheritance Patterns	Classic Triad (bone marrow failure, chromosomal fragility, malformations)	Number of patients with specific mutation and cancer reported by NCI (without history of transplant) n=163 patients	Adult Onset Cancers for Carriers	Key References
FANCA	AR	Yes	11 / 70		Apostolu et al., Nature Genetics 1996
FANCB	X-Linked Recessive	Yes	1		Meetei et al., Nature Genetics, 1996
FANCC	AR	Yes	3 / 19		Strathdee et al., Nature, 1992
FANCD1/BRCA2	AR / AD	Yes	2/3	Breast, Ovarian, Prostate, Pancreatic, Skin	Timmers et al., Molecular Cell, 2001; Kauff et al. NEJM, 2002, Pritchard et al. NEJM, 2016,
FANCD2	AR	Yes	7		Timmers et al., Molecular Cell, 2001
FANCE	AR	Yes	I		de Winter et al., AJHG, 2000
FANCF	AR	Yes	2		de Winter et al., AJHG, 2000
FANCG	AR	Yes	1 / 7		Garcia-Higuera et al., MCB, 1999
FANCI	AR	Yes	2/3		Dorsman et al., Cell, 2007; Sims et al., Nature Structural Molecular Biology, Smogorzewska et al., Cell, 2007
FANCJ/BRIPI	AR / AD	Yes	I	Ovarian	Cantor et al., Cell, 2001;Tung et al., Nature Reviews, 2016
FANCL	AR	Yes	Ι		Meetei et al., Nature Genetics, 2003
FANCN/PALB2	AR / AD	Yes	I	Breast, Pancreatic, Metastatic Prostate	Xia et al., Molecular Cell, 2006; Tung et al., Nature Reviews, 2016
FA NCP/SLX4	AR	Yes	I		Svendsen et al., Cell, 2009
FANCQ/ERCC4	AR	Yes	I		Bogliolo et al., AJHG, 2013
FANCT/UBE2T	AR	Yes	I		Zhang et al., Genome Research 2000; Hira et al., AJHG 2015
FANCU/XRCC2	AR	Yes	I		Jones et al., Genomics, 1995; Shamselden et al. JMG, 2012
FANCV/REV7	AR	Yes	I		Li and Benezra et al. Science 1996; Bluteau et al. JCI, 2016
FANCW/RFWD3	AR	Yes	1		Knies et al. JCI, 2017
FANCM	AR	No	I		Meetei et al., Nature Genetics, 2005
FACNO/RAD51C	AR / AD	No	I	Ovarian / Metastatic Prostate	Vaz et al. Nature Genetics, 2010 ; Tung et al., Nature Reviews, 2016

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Key References	Ameziane et al. Nature Communications, 2015 ; Tung et al., Nature Reviews, 2016
Adult Onset Cancers for Carriers	Ovarian / Metastatic Prostate
Number of patients with specific mutation and cancer reported by NCI (without history of transplant) n=163 patients	-
Classic Triad (bone marrow failure, chromosomal fragility, malformations)	No
Fanconi Genes Inheritance Patterns Classic Triad (hone marrow failure, chromosomal fragil malformations)	AR / AD
Fanconi Genes	FANCR/RAD51 AR / AD

AR= autosomal recessive, AD=autosomal dominant