1 2	Review paper FANCONI ANEMIA GENES AND REACTIVE OXYGEN SPECIES IN CANCER
3	DEVELOPMENT
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5	ABSTRACT
6 7	Fanconi Anemia (FA) is an autosomal recessive disease of childhood. However, the FA pathway is
8	responsible for the development of leukemia and the other cancers. It has been also demonstrated
9	that FA, an only human genomic instability syndrome is very sensitive to oxidative stress and
10	reactive oxygen species (ROS) overproduction. In the present work, we consider major mechanisms
11	of antioxidant protection in Fanconi anemia cells. We showed that there are two types of such
12	mechanisms: the suppression of reactive oxygen species overproduction by Fanconi anemia genes
13	through the activation of basic Fanconi anemia proteins under the conditions of oxidative stress and
14	the application of free radical scavengers able to react with iron-dependent reactive oxygen species
15	such as flavonoids rutin and quercetin. The last nontoxic compounds of vitamin P group might be
16	recommended for the treatment of Fanconi anemia patients. Then, we discussed the role of Fanconi
17	anemia proteins in cancer development.
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19	Key words: Fanconi anemia, ROS, antioxidants, rutin
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Fanconi anemia (FA) is an autosomal recessive disease of childhood characterized by progressive pancytopenia, developmental abnormalities, bone marrow failure, and high disposition to leukemia and other cancers. It has been demonstrated that the FA pathway is an important way in the development of such deadly adult pathologies as breast and cervical cancer [1,2]. These findings sharply increase an interest to studying FA molecular mechanisms. The special interest is attracted

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to Fanconi anemia because FA is the only human genomic instability syndrome uniquely sensitive
to oxidative stress. Earlier, an importance of reactive oxygen species (ROS) overproduction i.e.
oxidative stress has been demonstrated. Joenje *et al.* has shown that erythrocyte superoxide
dismutase (SOD) decreased in FA [3]. These authors also proposed that the formation of
chromosomal aberrations in FA anemia might be explained by the genetic toxicity of oxygen [4]
underlining an important role of oxidative stress in FA development.

32 EARLIER STUDIES

33 In 1960, medicine doctors from Russian Institute of Hematology for Children (Moscow) asked us as scientists working in the field of free radical-associated diseases to study the possible application of 34 35 antioxidants for the treatment of FA patients. We hoped that antioxidants and free radical scavengers 36 could act positively on behalf of FA patients. To study the effects of antioxidants on FA cells, ROS production especially superoxide in these cells was measured. Major results of this in vitro study 37 have been summarized as follows [5,6]: ROS formation was measured by lucigenin- and luminol-38 39 dependent chemiluminescence (CL) in non-stimulated and stimulated blood and bone marrow 40 leukocytes of FA patients. It was found that the FA blood leukocytes produced the enhanced level of 41 luminol CL in comparison with leukocytes from normal donors. Lucigenin CL was also higher in 42 FA blood leukocytes, although its effect was not very significant. Similar but smaller effects were observed for bone morrow FA leukocytes. Earlier, we have already demonstrated that flavonoids 43 44 rutin and its aglycone quercetin possess both free radical scavenging and chelating properties. We have shown that both flavonoids were the most effective inhibitors of iron-dependent microsomal 45 lipid peroxidation comparing to lipid peroxidation initiated by carbon tetrachloride [7]. (This work 46 47 had a good response; at present, it has been cited more than 800 times).

We proposed that ROS formation in FA leukocytes was the iron-catalyzed process because it was characterized by the enhanced luminol CL typical for the formation of iron-dependent ROS.

This proposal was supported by studying the effects of various free radical inhibitors on luminol CL 50 of FA leukocytes [5]: Superoxide dismutase (SOD) inhibited only slightly luminol CL, while 51 mannitol (the typical hydroxyl radical scavenger) and rutin were the strongest inhibitors. (Later on, 52 it was also demonstrated that rutin was a strong inhibitor of iron-dependent lipid peroxidation of rat 53 brain homogenates [8] and the free radical formation in iron-overloaded rats [9]. Moreover, it was 54 found that rutin efficiently inhibited free radical formation and oxyhemoglobin oxidation in β-55 thalassemic red blood cells [10]). These findings reassured us to apply the non-toxic flavonoid rutin 56 57 (vitamin P) for the treatment of FA patients.

The results of rutin therapy for a small group of FA patients were encouraging. Rutin (vitamin P) was permitted for application to patients (FA patients were under the supervision of medical doctors). No toxic side effects were observed in patients. ROS production by FA leukocytes sharply decreased and patients' health was essentially improved [6]. Unfortunately, there was no possibility to continue our study despite the numerous requests of medical doctors.

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64 THE NATURE OF REACTIVE OXYGEN SPECIES IN FA CELLS

65 Our findings demonstrated that ROS formation in FA cells was connected with the iron species. In 66 1992, in accord with the views of that time we proposed that the major damage produced by ROS in 67 cells is the direct interaction of ROS with biomolecules. As the superoxide, a major precursor of 68 ROS in cells is unreactive in free radical processes [11,12], we proposed that the reactive ironcontained species could be responsible for free radical damage in FA cells. It was suggested that the 69 70 iron-catalyzed superoxide conversion into reactive hydroxyl radicals (the Fenton reaction) was 71 responsible for their formation in FA cells. Indeed, there are some evidences of hypersensitivity of 72 FA cells in the presence of oxygen and iron [13]. Although reactive hydroxyl radicals are probably unable to achieve the target biomolecules, they can be formed during the contacts of superoxidewith DNA "iron fingers" [14].

A major conclusion from our previous work was that FA characterized by the enhanced 75 76 production of reactive iron-dependent species, which might be the source of fatal disorders in this hereditary disease. At present, many authors agreed that FA cells exist under the conditions of ROS 77 overloading. Thus, Hadjur et al. concluded that the abnormality of FA cells depended on ROS 78 overproduction [15]. Du et al. pointed out that FA is only human genomic instability syndrome, 79 80 which was uniquely sensitive to oxidative stress [16]. Therefore, there is no doubt about importance of ROS in Fanconi anemia. However, the damaging mechanisms of ROS activity remain 81 to be investigated. 82

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84 THE FA GENETIC PATHWAY

The most important discovery in FA molecular mechanisms was the identification of Fanconi 85 86 anemia genes responsible for synthesis of special FA proteins FANC (among them FANCA, 87 FANCE, FANCE, FANCE, FANCE, FANCG, FANCL and FANCM). It was found that eight major FA proteins FANCA, B, C, E, F, G, L, and M) formed a nuclear complex [16]. In response to DNA 88 89 damage or DNA replication stress FA complex monoubiquitinate into the two FA proteins FANCD2 90 and FANCI, which then recruite the other downstream FA proteins including FANCD1 (which is 91 also named the breast cancer protein BRCA2), FANCJ, and FANCN to enter nuclear loci containing 92 the damaged DNA.

It has been shown that FA proteins play the critical role in the regulation of oxidative stress. Thus, deficiency in FA genes apparently affect mitochondrial ROS [17]. The redox-sensitive proteins FANCA and FANCF exist as monomers under non-oxidizing conditions but form a new nuclear complex through the intermolecular disulfide bonds in response to oxidative damage [18].

97 FANCA, FANCC, and FANCG participate in redox processes in mice with combined deficiencies of
98 the genes encoding FANCC and Cu/Zn superoxide dismutase [15].

FA proteins function through the interaction with some enzymes. Saadatzadeh et al. showed that 99 100 Fancca-/-cells were highly sensitive to oxidants (hydrogen peroxide) and underwent enhanced 101 apoptosis [19]. Antioxidative compounds enhanced the survival of these cells. Thus, the redoxdependent ASK1 kinase was hyperactive in hydrogen peroxide-treated Fancca-/-cells. Another FA 102 103 protein FANCG interacted with mitochondrial antioxidant enzyme peroxiredoxin-3 and cytochrome 104 P450 2E1 (CYP2E1) [20]. This member of P450 superfamily responsible for ROS production and the activation of carcinogens. These findings suggested that the interaction of FANCG with 105 106 CYP2E1 might increase DNA oxidation.

107 It is known that the tumor protein p53 plays an important role in the prevention of cancer. 108 Furthermore, some findings demonstrate that p53 deficiency might enhance cancer development in 109 FA patients and FA mice. Therefore, it was suggested that FA proteins could interact with p53 110 under the conditions of oxidative stress. Freie *et al.* showed that ionizing radiation (IR) induced p53 111 elevated levels in cells from Fance mutant mice and that the inactivation of p53 enhanced TNFinduced apoptosis in myeloid cells from Fance-/-mice [21]. Rani et al. demonstrated that FA 112 113 proteins protected cells from the stress-induced proliferative arrest and tumor evolution 114 through the modulation of signaling pathways which connected FA proteins to p53 [22]. Du et 115 al. proposed that two major FA proteins FANCA and FANCC might coordinate with p53 in the regulation of oxidative stress response [16]. 116

117 It has been shown that the *Foxo3a* gene might be involved in ROS formation [23]. For 118 example, Tothova *et al.* showed that ROS levels increased in Foxo-deficient hematopoietic stem 119 cells that correlated with the changes in expression of ROS-regulated genes [24]. Correspondingly, 120 *Foxo3a* plays an important role in ROS regulation in FA cells. Thus, Li *et al.* showed that the treatment of FA cells with hydrogen peroxide stimulated the formation of a complex between FANCD2 and FOXO3a with subsequent monoubiquitination of FANCD2 [25]. It was suggested that the overexpression of *Foxo3a* reduced abnormal accumulation of ROS, enhanced cellular resistance to oxidative stress, and increased antioxidant gene expression only in cells corrected by a FANCD2 protein capable of interacting with FOXO3a.

It has been shown that ROS accelerated the development of hydrocephalus (abnormal 126 accumulation of cerebrospinal fluid in the brain) in mouse model of FA [26]. The deletion of 127 128 Foxo3a in FA mice increased the ROS accumulation and subsequently deregulated mitosis and ultimately apoptosis in the neural stem and progenitor cells NSPCs, leading to hydrocephalus 129 development. The antioxidants NAC and quercetin reduced ROS formation in both neural stem 130 131 cell NSCs and in the brain of double knockout (DKO) offspring mice. Importantly, quercetin greatly diminished the synthetic lethality imposed by DKO and completely eliminated 132 hydrocephalus in DKO mice. 133

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135 MECHANISMS OF ANTIOXIDANT PROTECTION IN FANCONI ANEMIA

136 ANTIOXIDANTS AND FREE RADICAL SCAVENGERS

As it was aforementioned, it has been previously shown that oxidative stress (ROS overproduction) in FA cells might be diminished by the application of flavonoid rutin, an antioxidant and free radical scavenger [5,6]. It was suggested that rutin might be able to scavenge reactive iron-depend ROS. Rutin is a nontoxic compound (vitamin P) permitted for the treatment of patients. Therefore, medical doctors were allowed to use rutin for the treatment of FA patients. They observed certain positive effects in patients. (It should be mentioned that several FA families asked us to prolong the treatment of FA children but we were unable to continue our work).

Now we were encouraged to find out that in subsequent studies flavonoids were used for the 144 suppression of oxidative stress in FA. It has been shown that quercetin (the aglycone of rutin) turned 145 out to be an effective antioxidant in FA cells. Antioxidant effect of quercetin was also shown in FA 146 147 animals. Li et al. found that mice deficient for the Fanca or Fance genes were diabetes-prone 148 when fed with a high-fat diet [27]. Treatment of FA mice with quercetin diminished diabetes and obesity. It was already demonstrated that quercetin reduced ROS formation and eliminated 149 hydrocephalus in double knockout mice [26]. Ponte et al. proposed that the cocktail of antioxidants 150 151 lipoic acid and NAC might be applied as a prophylactic approach to delay progressive clinical 152 symptoms in FA patients [28].

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156 ANTIOXIDANT EFFECTS OF FA GENES

157 In 2001, Hadjur *et al.* showed that encoding *Fancc* and Cu/Zn superoxide dismutase genes might be useful for the treatment of defective hematopoiesis and hepatic steatosis in mice [13]. These findings 158 159 suggested an important role of FA genes in protection of FA cells from oxidative stress. In 160 subsequent work, it was confirmed that FA genes were able to suppress ROS overproduction. Du et 161 al. showed that major antioxidant defense genes were downregulated in FA patients due to the 162 increased oxidative DNA damage in the promoters of antioxidant genes [29]. They showed that FA 163 proteins together with the chromatin-remodeling factor BRG1 protected the promoters of 164 antioxidant defense genes. Oxidative stress activated FA pathway through monoubiquitination of 165 FANCD2. After this, FANCA or FANCD2 proteins formed the ternary complex with BRG1 at the promoters of antioxidant genes. It has been suggested that this complex played essential role in the 166 protection of promoters from ROS damage. As it has been mentioned, ROS influenced the 167

development of hydrocephalus in mouse model of FA [26]. Combined deficiency of two FA genes *Fancc* and *Fancd2* led to the inactivation of *Foxo3a* gene, the enhancement of ROS level, and
apoptosis of neural stem and progenitor cells. Antioxidants quercetin and NAC reduced ROS
formation in the brain of mice, while quercetin completely suppressed hydrocephalus.
Mukhopadhyay *et al.* identified the FA group G (FANCG) protein in mitochondria, which interacted
with the mitochondrial peroxidase peroxiredoxin-3 (PRDX3) [30]. The formation of this complex
prevented the destruction of PRDX3 peroxidase and diminished ROS formation.

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176 **DISCUSSION**

177 Fanconi anemia is a rare chromosome instability syndrome, which is characterized by aplastic 178 anemia in childhood. It is known that FA possesses high disposition to leukemia and other cancers. 179 Now, there are new findings, which make us reconsider more carefully the connection between FA and cancer. Some new contemporary data demonstrate that the FA pathway is an important route for 180 181 the development of such deadly adult pathologies as breast and cervical cancer. Thus, D'Andrea 182 pointed out that abnormalities in the FA pathway are found not only in childhood Fanconi 183 anemia but also in sporadic cancers in adults [1]. Narayan et al. investigated the molecular genetic basis and the role of (FA)-BRCA pathway in cervical cancer (CC) [2]. These authors showed 184 that the *fancf* gene was disrupted by either the promoter hypermethylation or the deregulated gene 185 186 expression in cervical cancer. It was also found that gene inactivation in the FA-BRCA pathway by epigenetic alterations showed its major role in the development of cervical cancer. Chen et al. 187 demonstrated that the regulation of FANCD2 by m-TOR pathway led to the resistance of cancer 188 189 cells to DNA double-strand breaks [31]. It is not surprising that among thirteen FA genes one gene is the well-known breast cancer susceptibility gene brca2. 190

191 In normal cells, the FA pathway is not constitutively active but it activates by DNA damage. As

DNA damage depends on oxidative stress, ROS overproduction should be an important factor of FA 192 pathway activation. Mechanism of response of FA genes to DNA damage is not fully established, 193 but it has been shown that FA genes exist as nuclear complex before activation. In response to the 194 195 DNA damage or DNA replication stress FA complex monoubiquitinate into two FA proteins 196 FANCD2 and FANCI, which then recruite the other downstream FA proteins including the breast cancer protein BRCA2, FANCJ, and FANCN to enter nuclear loci containing the damaged DNA. 197 Then FANCA and FANCF form a new complex through the intermolecular disulfide bonds after 198 199 exposure to oxidative stress [16]. Thus, FA genes accomplish important defense function, which is confirmed by high sensitivity of *Fanc-/-* cells to ROS. 200

We now can visualize the complete defense scheme of ROS inhibition in FA cells. It has been shown that the oxidative stress (ROS overproduction) initiates the activation of FA genes, which remain inactive under unstressed conditions. After activation, FANCA and FANCF proteins form a complex through the intermolecular disulfide bonds, and we might propose that the disulfide complex formed by the reaction of superoxide with these proteins:

206 FANCA-SH + FANCF-SH + O_2 \rightarrow FANCA-SS-FANCF + H_2O_2

207 This proposal supported by the effect of SOD on the redox processes of FANCA, FANCC, and208 FANCG in mice [13].

Another mode of antioxidant activity of FA proteins is the protection of antioxidant gene promoters from ROS overproduction. Du *et al.* pointed out that oxidative stress is an important pathogenic factor in leukemia-prone marrow diseases such as Fanconi anemia [29]. It has been proposed that the FA pathway plays a crucial role in protecting major antioxidant defense genes from oxidative damage. This protection probably accomplished in response to oxidative stress by the interaction with the chromatin-remodeling machinery. Indeed, it has been shown that the oxidative stress-induced activation of FA pathway (FANCD2 ubiquitination) required for the

- 216 formation of the FA-BRG1-promoter complex. This complex is essential for the protection of the
- 217 antioxidant gene promoters from oxidative damage.

218 Thus, the activation of FA pathway under the conditions of oxidative stress led to cellular

219 protection through various ways. Taking into account the above-mentioned consideration, these

- 220 ways might be presented as follows:
- 221

222 **Table 1**

223 FA PATHWAYS OF CELLULAR PROTECTION

224	
225	1. Protection of antioxidant gene promoters
226 227	FANCD2 ubiquitination
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229	FA-BRG1-promoter complex
230	\downarrow
231	Protection of antioxidant gene promoters
252	
233	2. Formation of the disulfide complex
234	2. Formation of the disulfide complex
235	
236	$FANCA + FANCF + O_2^{T} \rightarrow FANCA - SS - FANCF$
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239	3. Suppression of DNA double-strand breaks
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241	FANCD2 \rightarrow m-TOR pathway \rightarrow DNA double-strand breaks
242	
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244	4. Suppression of ROS overproduction
245	
246	Antioxidants (quercetin and NAC) \rightarrow ROS \downarrow (diabetes and hydrocephalus in FA patients)
247	
248	5. Enhancement of gene antioxidant activity
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250	The interaction of genes with enzymes (Foxo3a, p50, p450)
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253 CONCLUSIONS

254 (1) Direct interaction of FANCA and FANCF with ROS (superoxide) forms a complex through the

255 intermolecular disulfide bonds.

- 256 (2) The regulation of FANCD2 by the m-TOR pathway might lead to the resistance of cells to DNA
- 257 double-strand breaks as in cervical cancer.
- 258 (3) FANCD2 ubiquitination following the formation of FA-BRG1-promoter complex capable of
- 259 protection of antioxidant gene promoters.
- 260 (4) Antioxidant activity of FA genes enhanced by the interaction with various enzymes such as
- 261 Foxo3a, p50, p450.
- 262 (5) Application of antioxidants and free radical scavengers (quercetin and NAC) to decrease ROS
- 263 overproduction in such diseases as diabetes or hydrocephalus in FA patients.
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