



Aldehyde dehydrogenase 2 in aplastic anemia, Fanconi anemia and hematopoietic stem cells



Lauren D. Van Wassenhove^{a,*}, Daria Mochly-Rosen^a, Kenneth I. Weinberg^{b,*}

^a Chemical and Systems Biology, Stanford University School of Medicine, Stanford, CA, USA

^b Division of Stem Cell Biology and Regenerative Medicine, Department of Pediatrics, Stanford University School of Medicine, Stanford, CA, USA

ARTICLE INFO

Article history:

Received 5 May 2016

Received in revised form 13 July 2016

Accepted 13 July 2016

Available online 15 July 2016

Keywords:

Fanconi anemia

Aldehydes

Hematopoiesis

Hematopoietic stem cell

Aplastic anemia

Aldehyde dehydrogenase

ALDH2

ABSTRACT

Maintenance of the hematopoietic stem cell (HSC) compartment depends on the ability to metabolize exogenously and endogenously generated toxins, and to repair cellular damage caused by such toxins. Reactive aldehydes have been demonstrated to cause specific genotoxic injury, namely DNA interstrand cross-links. Aldehyde dehydrogenase 2 (ALDH2) is a member of a 19 isoenzyme ALDH family with different substrate specificities, subcellular localization, and patterns of expression. ALDH2 is localized in mitochondria and is essential for the metabolism of acetaldehyde, thereby placing it directly downstream of ethanol metabolism. Deficiency in ALDH2 expression and function are caused by a single nucleotide substitution and resulting amino acid change, called *ALDH2*2*. This genetic polymorphism affects 35–45% of East Asians (about ~560 million people), and causes the well-known Asian flushing syndrome, which results in disulfiram-like reactions after ethanol consumption. Recently, the *ALDH2*2* genotype has been found to be associated with marrow failure, with both an increased risk of sporadic aplastic anemia and more rapid progression of Fanconi anemia. This review discusses the unexpected interrelationship between aldehydes, ALDH2 and hematopoietic stem cell biology, and in particular its relationship to Fanconi anemia.

© 2016 The Authors. Published by Elsevier Inc. This is an open access article under the CC BY-NC-ND license (<http://creativecommons.org/licenses/by-nc-nd/4.0/>).

Contents

1. Introduction	28
2. Genomic instability and Fanconi anemia.	30
2.1. Aldehyde-mediated DNA damage in Fanconi anemia.	30
3. Acetaldehyde	31
4. The aldehyde dehydrogenase (ALDH) enzyme family	31
4.1. Asian flushing syndrome and ALDH2	31
4.2. Increased risk of other malignancies associated with ALDH2 deficiency	32
4.3. ALDH isoforms and hematopoiesis	32
4.4. <i>ALDH2*2</i> and aplastic anemia	33
4.5. <i>ALDH2*2</i> and Fanconi anemia	33
5. Current therapies for metabolic diseases.	34
6. ALDH activators—a new strategy for treatment of metabolic disease	34
7. Aldehyde sensors to monitor substrate levels	34
8. Conclusions and future directions.	34
Acknowledgements	35
References.	35

1. Introduction

A central problem in hematology is the limited lifespan of mature non-lymphoid cells, which necessitates constant production of new

* Corresponding authors.

E-mail address: ldvanwas@stanford.edu (L.D. Van Wassenhove).

blood cells. In humans, the estimated lifespan of circulating red blood cells (RBCs) is 120 days, polymorphonuclear neutrophils (PMNs) 2 days, and platelets 5–10 days. To maintain the blood system, approximately 2×10^{11} RBCs, 1.6×10^{11} PMNs, and 10^{11} platelets are produced daily [1–4]. The evolutionary resolution of this problem in mammals is based on a hierarchy of self-renewing multipotent hematopoietic stem cells (HSC), which give rise to non-self-renewing, lineage committed hematopoietic progenitor cells (HPC) capable of massive expansion and differentiation into mature blood cells.

The dependence of blood cell production on a limited number of HSC and HPC (hereafter collectively referred to as HSPC) means that protection from potential toxins is essential for maintenance of the hematologic system (see Fig. 1). For example, the generation of blood cells within the intramedullary marrow space of heavily calcified bones

probably protects cells from typical doses of external ionizing radiation. Immature HSPC express a number of proteins which appear to protect them from toxicological injury. For example, the ABC transporter protein MDR1 expressed by HSC increases the export of various xenobiotics, including some chemotherapy drugs [5].

Because genotoxic injury is inevitable, an important protective mechanism is the expression of various proteins involved in DNA repair pathways [6]. These pathways include those for repair of single strand breaks, such as base excision repair (BER), nucleotide excision repair (NER), and mismatch repair (MMR). To repair double strand breaks, cells express pathways for non-homologous end joining (NHEJ), homologous recombination (HR), or microhomology-mediated end joining (MMEJ). Additionally, cells can use tolerance methods to continue to replicate DNA around lesions, called translesion synthesis (TLS).

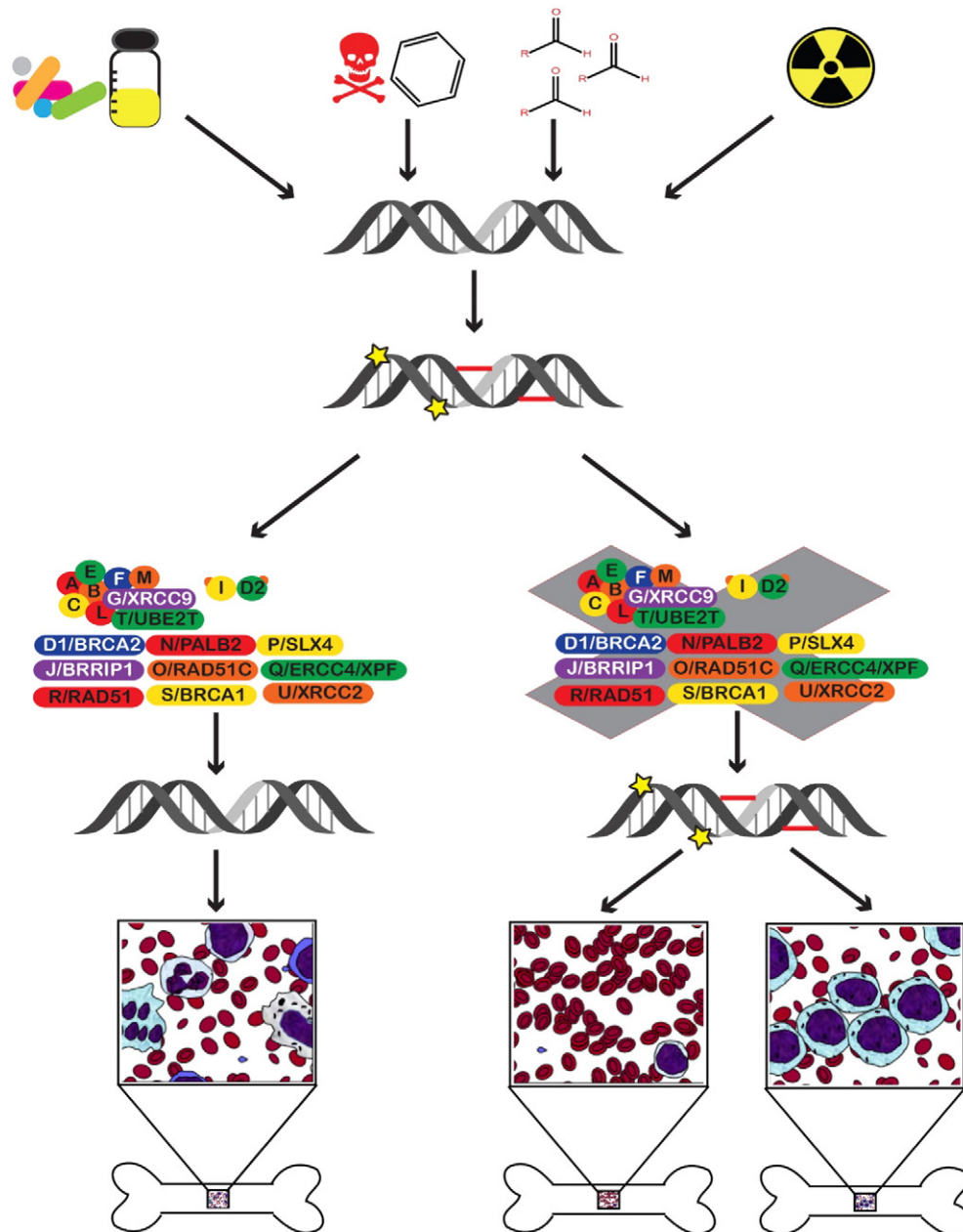


Fig. 1. Genomic instability and bone marrow failure: the genome is subject to a variety of different insults, including chemotherapy drugs, benzene and other toxic substances, radiation, and aldehydes ($RCH = O$). These can induce different types of DNA damage, including adduct formation, single and double strand breaks, and interstrand crosslinks. When the damage is appropriately repaired by DNA repair pathways, one of which is the FA pathway (depicted), there is no damage to HSPC in the bone marrow (depicted by a normal mixture of cells in the left inset). When the damage is not adequately repaired by DNA repair pathways, or by the FA pathway specifically, loss of HSPC can occur, leading to bone marrow failure (aplastic anemia, shown in the middle inset), or acute myelogenous leukemia (AML) shown on the right.

Download English Version:

<https://daneshyari.com/en/article/5513984>

Download Persian Version:

<https://daneshyari.com/article/5513984>

[Daneshyari.com](https://daneshyari.com)