### **Dienst Oncologie**



# **DPD Deficiency**

#### General Overview

- Dihydropyrimidine dehydrogenase (DPD), the first of three enzymes in the fluoropyrimidine metabolic pathway is the rate-limiting enzyme in the FU catabolism
- Patients who are partially or totally deficient in DPD activity cannot adequately degrade fluoropyrimidine, leading to increased toxicity, sometimes fatal.
- Molecular analysis of patients with DPD deficiency has identified over 128 mutations and polymorphisms in the *DPYD* gene that may result in partial or total loss of DPD activity.
- Only 4 have been consistently associated with enhanced 5-FU toxicity
  - o DPYD\*2A SNP
  - DPYD\*13 SNP (c.1679T>G)
  - DPYD\*9B SNP (c.2846A>T)
  - HapB3 (3 intronic variants and one synonymous variant)
- Complete DPD deficiency is rare. Partial deficiency occurs in 3.5% of women and 1.9% of men
- Inheritance of one of these high risk alleles does not account for all cases of DPD deficiency (sometimes epigenetic mechanisms).
- Other potential genotypes that cause toxicity: DPYD\*4 and DPYD\*5
- Preemptive testing for DPYD variants is recommended by the EMA (but not the FDA)
- Phenotypic testing (enzymatic function) may be an alternative
- More information: <u>All You Need to Know About DPYD Genetic Testing for Patients Treated With</u> <u>Fluorouracil and Capecitabine: A Practitioner-Friendly Guide | JCO Oncology Practice</u> (ascopubs.org)
- Implementation of dihydropyrimidine dehydrogenase deficiency testing in Europe (nih.gov)
- <u>Dihydropyrimidine Dehydrogenase Testing prior to Treatment with 5-Fluorouracil, Capecitabine,</u> and Tegafur: A Consensus Paper (karger.com)

#### Guidelines for treatment

Genotype	Activity score
Not carrier of decreased function/no function <i>DPYD</i> variant (*1/*1)	2.0
Heterozygous carrier of decreased function DPYD variant (*1/c.1236G>A or *1/c.2846A>T)	1.5
Heterozygous carrier of no function DPYD variant (*1/*2A or *1/*13)	1
Carrier of two decreased function <i>DPYD</i> variants (e.g., * <i>1/c.1236G&gt;A</i> and * <i>1/c.2846A&gt;T</i> ) or Carrier of one reduced function <i>DYPD</i> variant and one no function variant (combination of <i>c.1236G&gt;A</i> or * <i>1/c.2846A&gt;T</i> with *2 <i>A</i> or * <i>13</i> , such as <i>c.2846A&gt;T</i> )	0.5*
Homozygous carrier of no function <i>DPYD</i> variant (*2 <i>A</i> /*2 <i>A</i> ; *13/*13) or Heterozygous carrier of two no function <i>DPYD</i> variants (*2 <i>A</i> /*13)	0

The activity score of 0.5 is not always reliable and requires additional phenotyping [24].

# UZA

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