

Severe Fluoropyrimidines Toxicities: a simple and effective way to avoid them.

Screen Effectively for DPD Deficiencies

Boisdron Celle M¹, Capitain O¹, Metges JP², Guerin-Meyer V¹, Faroux R³, Stampfli CI⁴, Matysiak-Budnik T⁵, Leguellec Ch⁶, Gamelin E¹, Morel A¹

¹ Institut de Cancérologie de l'Ouest, Angers, France, ² CHU Hôpital Morvan, Brest, France, ³CHU, La Roche/Yon, France, ⁴CH, Laval, France, ⁵CHU Hôtel Dieu, Nantes, France, ⁶CHU Bretonneau, Tours, France

BACKGROUND

Severe, even fatal toxicities which occur during the first course of chemotherapy treatments using fluoropyrimidines pose a serious public health problem. This family of chemotherapy molecules is used in over 60% of protocols, in adjuvant and metastatic settings. Even protocols using orally-administered molecules cause the same toxicities.

PATIENTS

11,351 patients were screened for the risk of severe toxic reaction (April 2001-December 2011).

11,104 before treatment, 247 after toxicities (grade III-IV or death).

Chemotherapy protocols: FEC, LV5FU2, FOLFOX, FOLFIRI, Capecitabine, UFT.

METHODS

Toxicity risk evaluation combines **genotyping** ⁽¹⁾ (24 mutations) by pyrosequencing (Qiagen, Paris, France) and **phenotyping (UH₂/U ratio)** ⁽²⁾. **Multiparametric risk calculations** (genotyping + phenotyping + physiological + physiopathological parameters) were done via **ODPM Tox™** (ODPM, Angers, France). Patients deemed at risk before treatment received an individually-adapted dose via **ODPM Protocol™** (ODPM, Angers, France).

RESULTS

► **Results for the total population (11,351 patients)** are presented in Figure 1 (A).

- **346 patients (3%)** had **one or more mutations**.
- The most frequent mutation is the **2846 A>T (1.6%)**

► **Results for the population screened before treatment (11,104 patients)** are shown in Figure 1 (B).

- % of patients with one or more mutations among the 24 sought is the **same** as in the general population (**2.4%**).
- The mutation **most frequently** found is the **2846 A>T (1.3%)**.
- **1 patient** was a **homozygote DPYD2A** and one was **doubly mutated**.

Pre-treatment screening saved their lives.

► **Results for the patients screened after toxicities (247 patients)**

- **27 patients (11%) died** (5FU, Xeloda, UFT,...)
- **220 patients (89%)** showed **grade III-IV toxicities** (diarrhea, febrile neutropenia, coma, multi-organ toxicity).

The distribution of mutations in this population is shown in Figure 2: Total population (A), Patients who **died** (27 patients) (B).

- **60%** of patients who **died** had **one or more mutations**

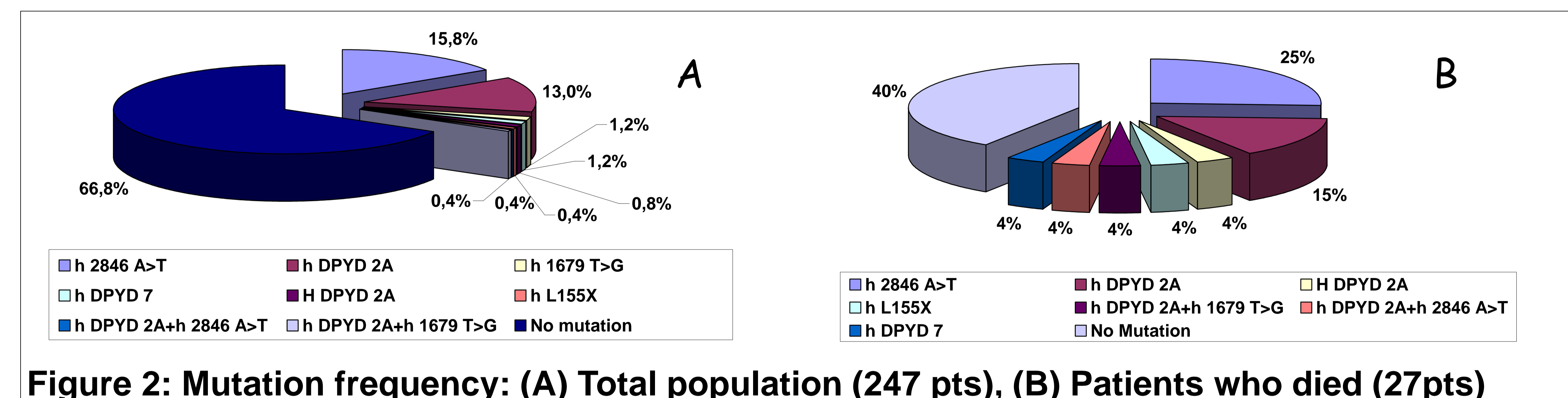


Figure 2: Mutation frequency: (A) Total population (247 pts), (B) Patients who died (27pts)

EFFECTIVE SCREENING FOR DPD DEFICIENCY

Patients	One or plus mutations	Deficient phenotypic status*	Multiparametric approach (ODPM Tox™)*
N	N (%)	N (%)	(%)
247	82 (33%)	211 (85%)	242 (98%)
27	16 (59%)	24 (89%)	27 (100%)

*Deficient phenotypic status = UH₂/U

*Multiparametric approach (ODPMTox™) = genotyping + phenotyping + physiological + physiopathological characteristics

CONCLUSION

Multiparametric approach (ODPM Tox™) is the key to **avoiding 100% of mortalities related to toxicities and 98% of severe toxicities**. Neither genotyping nor phenotyping alone are sufficient predictors of early onset toxicity.

Currently **2,000 patients screened annually using this approach** (ICO, France).

Clinical practice: 1 blood sample 10 days before starting treatment. Results in 8 days. Even DPD deficient patients can be treated using pharmacokinetic monitoring and appropriate dose adjustment (ODPM Protocol™).

References

Morel et al., Mol Cancer Ther. 2006; Morel et al. Clin Biochem 2007
Boisdron Celle M et al. Cancer Lett. 2007; Patents : 0503616, O6290592.2

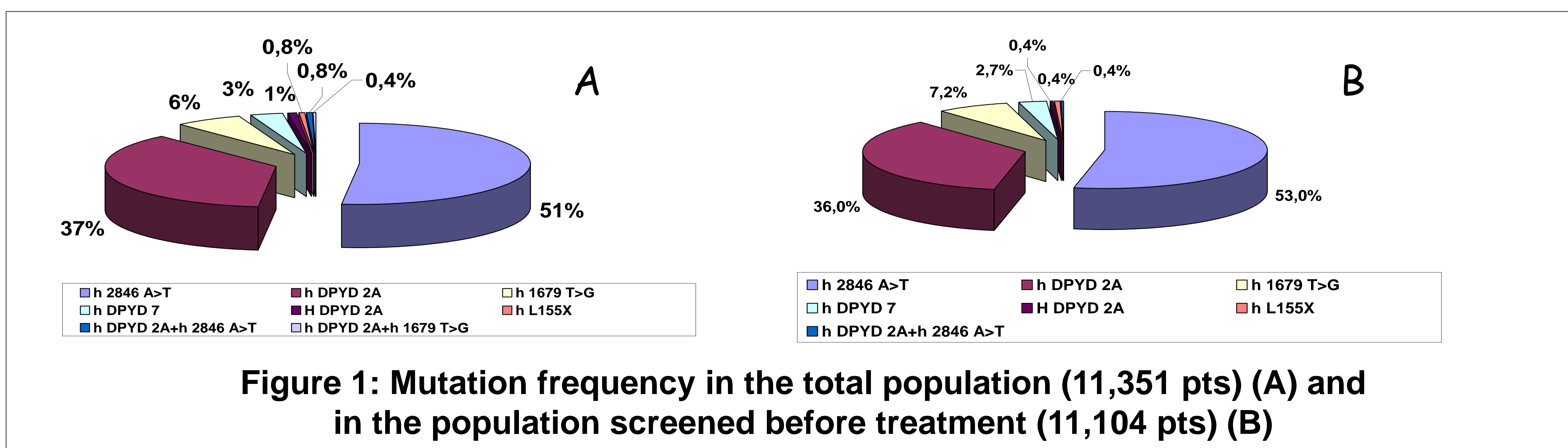


Figure 1: Mutation frequency in the total population (11,351 pts) (A) and in the population screened before treatment (11,104 pts) (B)