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Screening of DPD Deficiency before Fluoropyrimidine Chemotherapy

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Editorial Note:

Screening of dihydropyrimidine dehydrogenase (DPD) deficiency before systemic fluoropyrimidine chemotherapy can improve safety and prevent the occurrence of the associated toxicity. Both genotyping and phenotyping approaches have been advocated. In this review, Dr Felix Wong compares and contrasts both approaches and explains the upcoming situation in Hong Kong in the near future. We welcome any feedback or suggestions. Please direct them to <u>Dr Esther Hung</u> of Education Committee, the Hong Kong College of Pathologists. Opinions expressed are those of the authors or named individuals, and are not necessarily those of the Hong Kong College of Pathologists.

Introduction:

Pre-emptive testing for dihydropyrimidine dehydrogenase (DPD) deficiency is currently recommended for the prevention of fluoropyrimidine toxicity. Dihydropyrimidine dehydrogenase (DPD), which is encoded by the DPYD gene on chromosome 1, is the major catabolic enzyme of fluoropyrimidines responsible for 80% of their metabolism. Fluoropyrimidines include the chemotherapeutic agents 5-fluorouracil (5-FU) and its prodrugs (capecitabine and tegafur). DPD converts 5-FU to dihydrofluorouracil, which is non-cytotoxic. Genetic variants of the DPYD gene are associated with decreased DPD activity and increased fluoropyrimidine-related toxicity (bone-marrow and gastrointestinal toxicity, even death due to severe toxicity) due to an accumulation of 5-FU and its downstream active cytotoxic metabolites. In recent years, multiple guidelines recommending preemptive *DPYD* testing before fluoropyrimidine chemotherapy from different countries or regions have been published¹⁻⁶. These guidelines are focused on targeted genotyping of the following variants based on studies done in the Caucasian population: c.1905+1G>A (rs3918290, also known as *DPYD*2A*, *DPYD*:IVS14 + 1G>A), c.1679T>G (rs55886062, *DPYD*13*, p.1560S), c.2846A>T (rs67376798, p.D949V), and c.1129–5923C>G (rs75017182, HapB3 – corresponds to a combination of five genetically linked polymorphisms. It includes four intronic variants and one exon variant, two of which are in complete linkage disequilibrium: the c.1129-5923C>G variant (intronic variant) and the c.1236G>A variant (exon variant). The c.1129-5923C>G variant introduces a cryptic splice site and the partial production of a non-functional transcript.^{5,7}). Nevertheless, other rarer decreased- or no-function DPYD variants exist. Notably, based on

multiethnic population databases, e.g. Genome Aggregation Database (gnomAD, https://gnomad.broadinstitute.org/) and multiple studies performed in the East Asian population, it is known that these 4 variants are rare or absent in the East Asian population. For Hong Kong, an international city with a predominantly Chinese population, DPYD genotyping has been mostly performed in a retrospective manner to ascertain the cause in patients who had suffered from severe fluoropyrimidine toxicity⁸. Obviously, this approach does not prevent fluoropyrimidine toxicity or improve patient outcomes. It is uncertain how preemptive *DPYD* testing should be conducted in this locality. Whole DPYD gene sequencing with investigation of DPD enzyme activity for patients with novel variants has been proposed by some local experts⁹.

An alternative approach to DPYD genotyping is DPD phenotyping. 5-FU is an synthetic analog of uracil, which is the native substrate of the DPD enzyme located in the liver as part of the pyrimidine catabolic pathway. DPD deficiency is associated with an increase in uracil and decrease in dihydrouracil, the product of the DPD enzymatic reaction. Therefore, a high plasma uracil (U) concentration or a low dihydrouracil (UH2) to uracil ratio (UH2/U ratio) indicates DPD deficiency. DPD phenotyping by plasma uracil measurement is recommended by the European Medicines Agency¹⁰ for the detection of DPD deficiency and has been adopted in France¹¹ since 2018 using a cutoff of >=16 ng/mL, above which DPD deficiency is diagnosed. France is the first country in the world adopting DPD phenotyping by plasma uracil measurement for the screening of DPD deficiency and there has been an increase in the number of tests and European countries offering the test from 2019 to 2021¹².

Genotyping

The Activity Score system, proposed by Henricks et al13 and endorsed by the Clinical Pharmacogenetics Implementation Consortium (CPIC)⁵, translates DPYD genotype to a predicted DPD phenotype. An Activity Score of 2 represents normal DPD activity and is a sum of a score of 1 for each normal allele that an individual carries. A decreased-function allele is assigned a score of 0.5 and a no-function allele is assigned a score of 0. DPYD*2A and *13 alleles are assigned a score of 0 (no-function variant) while c.2846A>T and c.1129-5923C>G alleles are assigned a score of 0.5 (decreased-function variant). Based on this system, an Activity Score of 2 translates into DPYD normal metabolizer, while an Activity Score of 1 or 1.5 translates into DPYD intermediate metabolizer and an Activity Score of 0 or 0.5 translates into DPYD poor metabolizer. For example, homozygous DPYD c.1129-5923C>G has an Activity Score of 0.5 + 0.5 = 1, which is equal to the Activity Score of heterozygous DPYD*2A. Dosage recommendations based on the metabolizer status are available, ranging from a reduced starting dose of 50% in intermediate metabolizers to a complete avoidance of fluoropyrimidines for poor metabolizers with an Activity Score of 0. Furthermore, therapeutic drug monitoring of 5-FU is recommended for guiding dosage adjustment following initial dosing. Prospective clinical trials involving upfront testing for these variants with genotype-based dosage adjustments have demonstrated improved patient safety^{14,15} and cost-effectiveness^{15,16}, with no adverse impact on treatment response rates for the reduced fluoropyrimidine dose given to DPYD variant carriers¹⁷. Nevertheless, it is known that this approach has limited sensitivity (17%) for severe fluoropyrimidine-related toxicity [>=grade 3 according to the Common Terminology Criteria for Adverse Events (CTCAE)] because only 39 - 61% of patients having severe fluoropyrimidine-related toxicity are due to DPD deficiency, and even if DPD deficiency is present, it cannot be always traced back to a genetic alteration in the DPYD gene despite full gene sequencing¹⁸.

On the other hand, multiple studies performed in the East Asian population (Chinese^{19–21}, Korean^{21,22}, Japanese^{21,23}) have

demonstrated the scarcity of the *2A, I560S, D949V and c.1129-5923C>G in the studied populations. Indeed, fluoropyrimidine-induced toxicity appears to be less common in the East Asian population than other populations^{24,25}. From the DPYD allele frequencies based on population databases together with a comprehensive evaluation of variants by functional data/in silico predictions, it has been deduced that the East Asian population may have a lower incidence of DPD deficiency than the European population (3.4% and 0.02% of intermediate metabolizer and poor metabolizer, respectively, in East Asians, in contrast to 7.6% and 0.09%, respectively, in non-Finnish Europeans). 26 Table 1 shows a comparison of minor allele frequencies (MAF) in Genome Aggregation Database (gnomAD) of the 4 variants in the European and East Asian populations. It can be deduced that targeted genotyping of these 4 variants in the local Chinese population will result in poor sensitivity of detecting DPD deficiency, and whole DPYD gene sequencing is required for the detection of other deleterious DPYD alleles. Based on the literature review performed by Zhou et al, 49 DPYD variants (*2A, I560S and D949V included) are considered to be pathogenic²⁶, and by in silico predictions of all observed DPYD variants in the global population in the same study, it was estimated that a large number of DPYD variants (174 variations, which corresponds to only 34.3% of deleterious DPYD variants) have to be tested in order to explain 95% of genetically encoded functional DPD variability in the population, underscoring the presence of significant allelic heterogeneity. DPYD variants with unknown functional impact may be analyzed by in silico prediction tools, which are computational algorithms developed to predict the functional impact of DPYD variants, e.g. DPYD-Varifier²⁷, Absorption, Distribution, Metabolism and Excretion (ADME) Prediction Framework^{26,28}, or evaluated using in vitro functional assays of DPD enzyme activity, which involves expression of DPYD variants in mammalian cells and incubation of radiolabelled 5-FU coupled with the measurement of rate of formation of dihydrofluorouracil from 5-FU^{29–32}. It has been suggested that computation algorithms specifically developed for pharmacogenes (e.g. ADME Prediction Framework) or in a gene-specific manner (e.g. DPYD-Varifier) may be preferable to conventional prediction methods for non-pharmacogenes because conventional methods are primarily based on evolutionary conservation while pharmacogenes are poorly conserved³³. Given the rarity of each individual DPYD variant, in vivo data from clinical studies is mostly unavailable or limited to isolated case reports. Nevertheless, the approach of including additional DPYD deficiency variants with evidence of a deleterious impact on protein function equivalent to the CPIC high level evidence variants has been shown to improve the sensitivity of the prediction of grade 3/4 haematological toxicity while retaining excellent specificity.34

DPYD Variant	MAF (European, non-Finnish)(%)	MAF (East Asian)(%)	
c.1905+1G>A (rs3918290, *2A)	2.385	0	
c.1679T>G (rs55886062, I560S)	0.06220	0	
c.2846A>T (rs67376798, D949V)	0.5163	0.005013	
c.1129-5923C>G (rs75017182)	2.102	0.1926	

Table 1. Minor allele frequencies (MAF) for *2A (c.1905+1G>A), I560S (c.1679T>G), D949V(c.2846A>T) and c.1129–5923C>G. Source: Genome Aggregation Database, accessed 20 December 2023.

A prospective study was performed from November 2020 to December 2021 in the author's hospital recruiting patients requiring fluoropyrimidine-containing chemotherapy. Polymerase chain reaction followed by Sanger sequencing of the 23 coding exons, their flanking intronic regions and HapB3 (c.1129-5923C>G included) of the *DPYD* gene was performed. 103 patients (97 Chinese, 94%) requiring chemotherapy for colorectal, stomach and other cancers were recruited. 4

heterozygous decreased-function/no-function *DPYD* variants were detected in 4 Chinese patients (heterozygous c.2210C>T, p.Thr737lle: 2 patients; heterozygous c.220C>T, p.Arg74Ter and heterozygous c.1314T>G, p.Phe438Leu: 1 patient each). In addition, two heterozygous VUS were detected in three patients (heterozygous c.2303C>A, p.Thr768Lys: two Chinese patients; heterozygous c.2528T>C, p.lle843Thr in a Filipino patient). None of the well-established alleles affecting *DPYD* function were detected in the study population.

Phenotyping

The DPD enzyme activity is mainly located in the liver in vivo. It is also present in peripheral blood mononuclear cells (PBMCs) which can be measured ex vivo and acts as a surrogate marker of the hepatic enzymatic activity³⁵. Nevertheless, the assay is not widely available because it is time-consuming, requires a large volume of blood and the use of radiolabeled materials³⁶. Of note, only 25% of patients with a decreased DPD enzyme activity carries one of the four DPYD variants³⁷. Alternatively, plasma uracil and dihydrouracil measurements by High Performance Liquid Chromatography - Ultraviolet detection (HPLC-UV) or Liquid Chromatography Tandem Mass Spectrometry (LC-MS/MS) have been studied as surrogate markers of DPD enzyme activity. The theoretical advantage of DPD phenotyping resides in the possibility of identifying more patients with DPD deficiency beyond which can be explained by DPYD genetic variants alone. In a study of 550 patients requiring fluoropyrimidine chemotherapy with pretreatment serum uracil measured, a concentration of >=16 ng/mL strongly associated with global severe toxicity (odds ratio 5.3, p = 0.009). Serum uracil was shown to correlate better with DPD activity in PBMCs than UH2/U ratio and be superior to UH2/U ratio as a predictor of severe toxicity³⁸. The same cutoff in another study demonstrated a sensitivity of 67% and specificity of 92% in the prediction of grade 4 capecitabine toxicity with a relative risk of 20.6 (p = 0.021)³⁹. The distribution of plasma uracil in the pre-treatment population across studies is shown in Table 2. One of these studies was performed in the Chinese population⁴⁰. Plasma uracil >= 16 ng/mL was observed in approximately 10 - 15% of the population 39,41 . Plasma uracil measurement has been endorsed by the European Medicines Agency (EMA) in 2020 for the detection of DPD deficiency 10. Detailed guidelines on the detection of DPD deficiency by plasma uracil measurement are available in France (>=16 ng/mL to 150 ng/mL: partial DPD deficiency; >= 150 ng/mL: complete DPD deficiency)⁴² and Belgium (>= 14 ng/mL to 100 ng/mL: partial DPD deficiency; >=100 ng/mL: complete DPD deficiency)⁴³. Dosage adjustment based on plasma uracil leads to a decreased 5FU toxicity in patients with partial DPD deficiency, while the resulting lower dose of 5FU received by these patients suggests the need for therapeutic drug monitoring of 5FU to uphold treatment efficacy⁴⁴. Plasma uracil is unstable in whole blood samples and strict preanalytical requirements are required (maximum delay between sample and centrifugation of 1h30 mins if the sample is stored at ambient temperature, and 4h if it is stored at + 4°C, centrifugation preferably at + 4°C then immediate freezing of the resulting plasma, transport respecting the cold chain)⁴², otherwise, artefactual elevations may occur⁴⁵. There are known effects of food and circadian rhythm affecting plasma uracil concentration 45,46. Plasma uracil concentration may be elevated in renal impairment^{47,48}, abnormal liver function⁴⁸ and tumor lysis syndrome⁴⁹. Patients with end stage renal disease on haemodialysis were shown to have a higher uracil concentration pre-analysis (mean = 14 ng/mL) than post-dialysis (mean = 8 ng/mL)⁵⁰. Plasma uracil concentration is also elevated if fluoropyrimidine has already been administered because of the competition between uracil and 5-FU for metabolism by DPD⁵¹.

In the author's laboratory, residual plasma from 74 specimens delivered on ice for plasma renin analysis was used for plasma uracil analysis. The average, median, 5th and 95th percentile of plasma uracil were: 9.8, 9.6, 5.9 and 16.2 ug/L. Therefore,

around 5% of samples is expected to show a plasma uracil >=16 ug/L ("Flagging rate"), indicating a positive result for DPD screen. Uracil >= 150 ug/L is expected to be rare (occurrence rate 0.08% in France)¹¹ and there were no such cases in our 74 specimens.

Study	Median (ng/mL)	Range (ng/mL)	No. of subjects (n)	Methodology	Diagnosis	Ethnicity
Coudoré et al, 2012 ⁵²	8.4	0.6 – 43.3	26	LC-MS/MS	advanced or recurrent cancer	French
Etienne- Grimaldi et al, 2017 ³⁹	9.6	3.9 – 75.3	205	HPLC-UV	Advanced breast cancer patients	French
Pan et al, 2017 ⁴⁰	11.2	5.16 – 120.6	68	HPLC-UV	Colorectal cancer patients	Chinese
Dolat et al, 2020 ⁵³	10.8	3 – 37.6	169	LC-MS/MS	Colorectal, pancreas and stomach cancer	French
Tafzi et al, 2020 ⁴¹	10.6	3.9 – 81.6	526	LC-MS/MS	Patients before 5FU treatment	French

Table 2. Distribution of plasma uracil concentrations across different studies. LC-MS/MS: liquid chromatography tandem mass spectrometry; HPLC-UV: high performance liquid chromatography – ultraviolet spectroscopy.

Genotyping and Phenotyping

Plasma uracil concentration correlates with the presence of known deleterious DPYD variants *2A, I560S and D949V – such variants were associated with a higher plasma uracil concentration, though not necessarily higher than the cutoff of 16 ng/mL, while this association appears to be weaker for HapB3^{38,39,54}. The presence of any of the 4 variants has a sensitivity and specificity of 11% and 95%, respectively, for the detection of DPD deficiency defined as a uracil concentration of > 16 ng/mL⁵⁴. A retrospective study of 472 patients with DPYD targeted genotyping (the 4 DPYD variants recommended by CPIC, plus DPYD*7, a frameshift variant classified to be a no-function allele by CPIC) and phenotyping (plasma uracil and UH2/U ratio) done without fluoropyrimidine dosage being adjusted according to such results (either because the patient has already suffered from toxicity of fluoropyrimidine or because such testing was done in parallel with treatment initiation) demonstrated sensitivities and specificities of targeted genotyping and plasma uracil to be 33%/59% and 95%/84%, respectively, for the prediction of grade 3 or higher fluoropyrimidine toxicity⁵⁵.

For the local population who is predominantly Chinese, it was demonstrated that none of the 4 variants recommended for testing by CPIC (c.1905+1G>A, c.1679T>G, c.2846A>T and c.1129–5923C>G) were detected in the author's laboratory, while

sequencing of whole *DPYD* gene enabled the detection of other *DPYD* variants enriched in the East Asian population which may lead to DPD deficiency and confer susceptibility to fluoropyrimidine toxicity. Nevertheless, these variants are less well characterized and their interpretation in the local population as decreased or no function variants relies on in vitro or in silico data, i.e. human data is lacking. According to the CPIC guideline, when two different decreased-/no -function variants are detected in a patient, it is assumed that the two variants are in trans, i.e. located on different alleles, while this may not be true if one of the two detected variants is novel. In the absence of haplotyping techniques, phenotyping may be helpful to differentiate partial from complete DPD deficiency.⁵ This approach has been highlighted in a review by Knikman et al. which outlined the possibility of combining phenotyping with genotyping in the same patient. They proposed that while dose modification may be based on genotyping result if one known variant is detected, dose modification should be based on phenotyping results in the event of two variants detected or no variants detected. In fact, even if two deleterious alleles are proven to be in trans by haplotyping techniques, the Dutch Pharmacogenetics Working Group (DPWG) guideline recommends the use of phenotyping to determine the residual DPD activity for guiding the fluoropyrimidine use at significantly reduced doses².

Screening of DPD deficiency in Hong Kong

While screening of DPD deficiency by plasma uracil has not been formally studied in the Chinese population, it is assumed to be applicable to all populations because the phenotype of uracilemia is reasoned to a biomarker of DPD deficiency independent of ethnicity⁵⁷. A decision was made to go for the phenotyping rather than genotyping approach and Queen Mary Hospital Chemical Pathology will launch preemptive screening of DPD deficiency by plasma uracil measurement to all patients requiring systemic fluoropyrimidine chemotherapy in the Hospital Authority in 2024 Q1. Plasma uracil is measured by liquid chromatography tandem mass spectrometry, which requires in-house method development and evaluation. We have subscribed to an external quality assurance program based in France (ASQUALAB). Uracil is an unstable analyte in plasma and an unbroken cold chain is required from specimen collection to centrifugation and storage. A turnaround time of less than 10 calendar days, which fulfils the French standard⁴², will be implemented to minimize the delay before treatment initiation. Under this service, genotyping is not required and will not be performed, even for patients screened positive for partial or complete DPD deficiency by plasma uracil. While fluoropyrimidines are contraindicated in patients with complete DPD deficiency, patients with partial DPD deficiency may be started on a reduced dose and the option of therapeutic drug monitoring of intravenous infusion of 5-fluorouracil (the topic of which is beyond the scope of this article) will be available as a part of the service to improve the drug safety of fluoropyrimidines. The service of measuring plasma 5FU level will be launched simultaneously with that of plasma uracil. According to an article published in 2023, no preventable toxicity-related death has been declared to the French Pharmacovigilance Network in patients with complete or partial DPD deficiency since 2020⁵⁷. We hope that the same benefits will soon become a reality in Hong Kong.

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