

## REVIEW ARTICLE

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## Genotype and phenotype in patients with dihydropyrimidine dehydrogenase deficiency

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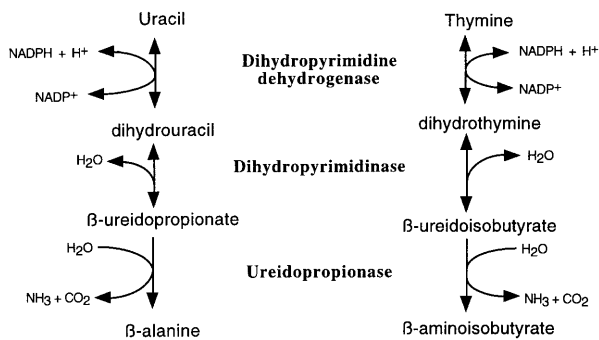
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**Abstract** Dihydropyrimidine dehydrogenase (DPD) deficiency is an autosomal recessive disease characterised by thymine-uraciluria in homozygous deficient patients and has been associated with a variable clinical phenotype. In order to understand the genetic and phenotypic basis for DPD deficiency, we have reviewed 17 families presenting 22 patients with complete deficiency of DPD. In this group of patients, 7 different mutations have been identified, including 2 deletions [295–298delTCAT, 1897delC], 1 splice-site mutation [IVS14+1G>A] and 4 missense mutations (85T>C, 703C>T, 2658G>A, 2983G>T). Analysis of the prevalence of the various mutations among DPD patients has shown that the G→A point mutation in the invariant splice donor site is by far the most common (52%), whereas the other six mutations are less frequently observed. A large phenotypic variability has been observed, with convulsive disorders, motor retardation and mental retardation being the most abundant manifestations. A clear correlation between the genotype and phenotype has not been established. An altered β-alanine, uracil and thymine homeostasis might underlie the various clinical abnormalities encountered in patients with DPD deficiency.

### Introduction

In mammalian liver, the pathway for the catabolism of uracil and thymine consists of three consecutive steps (Fig. 1). Dihydropyrimidine dehydrogenase (DPD, EC 1.3.1.2) is the initial and rate-limiting enzyme in the catabolism of the pyrimidine bases; it catalyzes the reduction of uracil and thymine to 5,6-dihydrouracil and 5,6-dihydrothymine, respectively. The second step consists of a hydrolytic ring-opening, which is catalyzed by dihydropyrimidinase (EC 3.5.2.2). Finally, β-ureidopropionic acid (N-carbamyl-β-alanine) or β-ureidoisobutyric acid (N-carbamyl-β-amin-



**Fig. 1** Catabolic pathway of the pyrimidines uracil and thymine

oisobutyric acid) is converted to  $\beta$ -alanine or  $\beta$ -aminoisobutyric acid, ammonia and  $\text{CO}_2$  by  $\beta$ -ureidopropionase (EC 3.5.1.6).

In children, a deficiency of DPD is often accompanied by a neurological disorder but a considerable variation in the clinical presentation among these patients has been reported (Van Gennip et al. 1994, 1997). In these patients, a large accumulation of uracil and thymine has been detected in urine, blood and cerebrospinal fluid, whereas no activity of DPD has been detected in fibroblasts and mononuclear cells (Bakkeren et al. 1984; Van Gennip et al. 1993, 1994, 1997).

DPD is also responsible for the breakdown of the widely used anti-neoplastic agent 5-fluorouracil (5FU). The catabolic route plays a significant role, since more than 80% of the administered 5-FU is catabolized by DPD (Heggie et al. 1987). In this light, a pharmacogenetic disorder has been described concerning cancer patients with a complete or partial deficiency of DPD and suffering from a severe or even life-threatening toxicity after the administration of 5FU. Recently, it has been shown that two such patients are genotypically heterozygous for a mutant allele of the gene encoding DPD (Wei et al. 1996; Van Kuilenburg et al. 1997a, 1998a, b). The detection of more than 30 patients of various nationalities with a (partial) DPD deficiency within 15 years in The Netherlands alone suggests that this type of inborn error is less rare than previously assumed (Van Gennip et al. 1997).

The recent cloning of the cDNA coding for human DPD and the sequence of the entire human DPD gene (*DPYD*) has allowed the detection of the defects at the molecular level. In this manuscript, we review the results of the analysis of the genotype and phenotype of 17 families presenting 22 patients who have complete deficiency of DPD (i.e. no detectable DPD activity in fibroblasts, <0.2% of controls) and who have been diagnosed in our laboratories. Clinical histories have been updated and three newly identified patients have been included. In addition, the possible pathological mechanisms underlying the various clinical abnormalities are discussed.

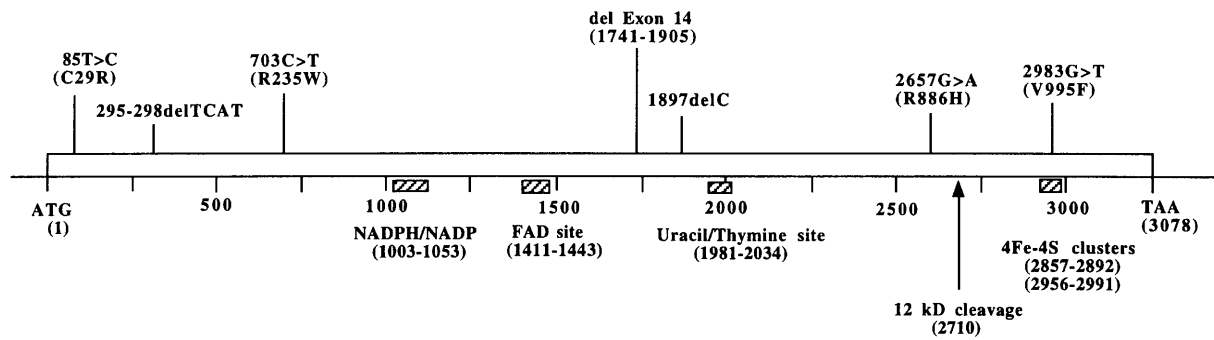
## Structural and kinetic properties of dihydropyrimidine dehydrogenase

DPD catalyzes the NADPH-dependent reduction of uracil and thymine to 5,6-dihydrouracil and 5,6-dihydrothymine, respectively. The activity of DPD is exclusively present in the cytosol (Van Kuilenburg et al. 1997b) and can be detected in a variety of tissues with the highest activity being found in the liver (Naguib et al. 1985; Ho et al. 1986) and peripheral blood monocytes (Van Kuilenburg et al. 1997c, 1998c, d). The enzyme has been purified and extensively characterised from liver tissues of rat (Shiotani and Weber 1981), pig (Podschun et al. 1989), bovine (Porter et al. 1991) and man (Lu et al. 1992, 1993). These studies have demonstrated that the native mammalian enzyme has a molecular mass of approximately 210 kDa and is composed of two identical subunits. The C-terminal region of DPD is sensitive to proteolysis resulting in the cleavage of a 12-kDa peptide from the native enzyme. DPD contains various tightly associated prosthetic groups including two FMN, two FAD and at least two [4Fe-4 S] clusters (Podschun et al. 1989; Lu et al. 1992). With respect to the kinetic properties of the enzyme, a non-classical two-site ping-pong mechanism has been proposed for the pig liver enzyme, whereas a rapid equilibrium random kinetic mechanism has been put forward for the bovine liver enzyme (Podschun et al. 1990; Porter and Spector 1993). However, both kinetic mechanisms necessitate that two separate binding sites are available for uracil (or thymine) and NADPH, respectively.

## Structural organisation of the human DPD gene

The cDNAs coding for human DPD, pig DPD and bovine DPD have been isolated and sequenced (Yokota et al. 1994; Albin et al. 1996). Mammalian DPD appears to be relatively conserved throughout evolution, since a comparison of the deduced amino acid sequences of bovine DPD with that of pig and human DPD has shown a homology of 93% and 92%, respectively. The human cDNA encodes a protein containing 1025 amino acids with a calculated molecular weight of 111 kDa (Fig. 2). The conserved domains corresponding to a possible NADPH-binding site and FAD-binding site have been found in the N-terminal and middle region of the enzyme. In the C-terminal region, typical motifs for [4Fe-4 S] clusters have been found between residues 953 and 964 and residues 986 and 997. On the basis of chemical modification studies, the putative uracil-binding site of DPD has been located between Gly-661 and Arg-678. Thus, the functional domains of DPD can be arranged, from the N-terminus, in the order of NADPH-FAD-uracil-[4Fe-4 S].

Recently, the entire human DPD gene has been isolated and shown to consist of 23 exons, with exon 15 (69 bp) being the smallest and exon 23 (961 bp) the largest (Johnson et al. 1997; Wei et al. 1998). A physical map indicates that the human DPD gene is a least 950 kb in length with 3 kb of



**Fig. 2** Localisation of mutations in human DPD cDNA. The nucleotides involved in the three frameshift mutations are given. In addition, four missense mutations and their corresponding amino acid substitutions (given *in parenthesis*) are shown. The *hatched boxes* represent the localisation of conserved motifs corresponding to putative binding sites of the prosthetic groups. The nucleotide sequences involved are given *in parenthesis*. The proteolytic-sensitive site (*vertical arrow*) is close to the carboxy terminus and encompasses the iron-sulphur clusters

coding sequence and a minimal average intron size of about 43 kb (Wei et al. 1998). In addition, the human DPD gene has been mapped to chromosome 1p22 and is present as a single-copy gene (Takai et al. 1994; Wei et al. 1998).

### DPD mutations in patients with DPD deficiency

In order to understand the genetic basis of DPD deficiency, we have analyzed 17 families presenting 22 patients with complete deficiency of DPD. In this large group of patients, we have identified 7 different mutations, including 1 splice-site mutation, 2 deletions and 4 missense mutations (Table 1). The G→A point mutation changes an invariant GT splice donor site to AT; this leads to the skipping of exon 14 immediately upstream of the mutated splice donor site, during the splicing of DPD pre-mRNA. As a consequence, a 165-bp fragment encoding amino acid residues 581–635 of the primary sequence of the DPD protein is lacking in the mature DPD mRNA (Meinsma et al. 1995; Vreken et al.

1996). Both the four-base deletion 295–298delTCAT and the 1897delC mutation cause a frameshift leading to a premature stop codon shortly thereafter (Vreken et al. 1997a, b). The TCAT(295–298) deletion is located in a TCAT tandem-repeat sequence and most probably results from slipped mispairing or unequal chromosome crossing-over. Expression of the missense mutations C29R, R235W, R886H and V995F in *Escherichia coli* has demonstrated that C29R, R235W and V995F result in mutant DPD proteins with no significant residual enzyme activity (Vreken et al. 1997c, 1998). However, the DPD protein with the R886H mutation still possesses a residual activity of 25% and it is therefore unlikely that this mutation is responsible for the observed complete deficiency in one patient who has proved to be homozygous for both the C29R and R886H mutations (Vreken et al. 1997c). The mutations identified so far are randomly distributed along the cDNA and there are no apparent hot spots present (Fig. 2). All mutations are located outside those regions known to be involved in the binding of the various substrates and prosthetic groups, with the exception of the V995F mutation, which is located in the C-terminal region of DPD, a region thought to be involved in the binding of a [4Fe-4 S] cluster.

The majority of the patients (68%) have proved to be homozygous for one of the identified mutations, whereas the remaining patients are compound heterozygotes. Analysis of family members of the index patients for the presence of mutant alleles has shown that the mutations segregate following a recessive pattern of inheritance (Meinsma et al. 1995; Vreken et al. 1997c), in accordance with the pattern

**Table 1** Mutations in patients with DPD deficiency

	Name <sup>a</sup>	Genotype <sup>b</sup>	Effect <sup>c</sup>	Location <sup>d</sup>	Allele frequency
	Splicing				
	<i>DPYD</i> *2A	IVS14+1G>A	Del (exon 14)	IVS14	23/44 (52%)
	Frameshift				
	<i>DPYD</i> *7	295–298delTCAT	Frameshift	EX4	7/44 (16%)
	<i>DPYD</i> *3	1897delC	Frameshift	EX14	3/44 (7%)
	Missense				
	<i>DPYD</i> *9A	85T>C	C29R	EX2	7/44 (16%)
	<i>DPYD</i> *8	703C>T	R235W	EX7	1/44 (2%)
		2657G>A	R886H	EX21	2/44 (4%)
	<i>DPYD</i> *10	2983G>T	V995F	EX23	2/44 (4%)
		Unknown	-	-	1/44 (2%)

<sup>a</sup> Nomenclature according to McLeod et al. (1998)

<sup>b</sup> Nomenclature according to Antonarakis (1998)

<sup>c</sup> Effect of the mutation on DPD protein or mRNA

<sup>d</sup> According to Wei et al. (1998)

**Table 2** Genotype and phenotype of patients with DPD deficiency at diagnosis (– none, + mild, ++ severe)

Patient <sup>a</sup>	Genotype	Convulsions	Motor retardation	Mental retardation	Growth retardation	Microcephaly	Dysmorphism	Autism	Ocular abnormalities	Others <sup>b</sup>
1 (NL)	ΔEX14/ΔEX14	–	+	++	–	–	–	–	+	1
2 (NL)	ΔEX14/ΔEX14	+	+	+	–	–	–	+	–	2
3 (DK)	ΔEX14/ΔEX14	+	+	+	–	–	–	–	–	3
4 (DK)	ΔEX14/ΔEX14	–	–	–	–	–	–	–	–	4
5 (S)	ΔEX14/ΔEX14	–	+	–	+	–	–	–	+	5
6 (DK)	ΔEX14/ΔEX14	++	++	++	–	–	–	–	–	6
7 (NL)	ΔEX14/ΔEX14	–	–	–	–	–	–	–	–	6
8.1 (SF)	ΔEX14/ΔEX14	+	–	–	–	–	–	–	–	7
8.2 (SF)	ΔEX14/ΔEX14	–	–	–	–	–	–	–	–	8
9.1 (NL)	ΔTCAT/ΔTCAT	+	+	–	–	+	–	–	–	9
9.2 (NL)	ΔTCAT/ΔTCAT	–	–	–	–	–	–	–	–	9
9.3 (NL)	ΔTCAT/ΔTCAT	+	–	+	–	–	–	–	–	9
10 (IT)	C29R/C29R	–	+	++	–	–	+	++	–	10
11 (NL)	V995F/V995F	+	–	+	–	–	–	+	–	11
12.1 (TUR)	ΔC1897/ΔC1897	–	–	–	–	–	–	–	–	12
12.2 (TUR)	ΔC1897/R235 W	+	++	++	++	++	+	–	+	13
13 (TUR)	C29R, R886H/ C29R, R886H	–	–	–	+	–	–	–	–	14
14.1 (NL)	C29R/ΔEX14	+	–	–	–	–	–	–	–	15
14.2 (NL)	C29R/ΔEX14	–	–	–	–	–	–	–	–	15
15 (NL)	ΔEX14/ΔTCAT	+	++	++	–	+	–	++	+	16
16 (NL)	ΔEX14/C29R	–	–	–	–	–	–	–	–	17
17 (NL)	ΔEX14/?	–	+	+	+	–	+	–	+	18
Total		10/22 (45%)	10/22 (45%)	10/22 (45%)	4/22 (18%)	3/22 (14%)	3/22 (14%)	4/22 (18%)	5/22 (23%)	

<sup>a</sup> The nationality of the patient is given in parenthesis

<sup>b</sup> (1) Ocular abnormalities (bilateral microphthalmia, iris and choroida coloboma) and nystagmus (Bakker et al. 1994). (2) CT scan showed strong contrast between white and grey matter. (3) Delayed development of speech (Christensen et al. 1998). (4) Lethargy (Christensen et al. 1998). (5) Bilateral ptosis, progressive external ophthalmoplegia, retinitis pigmentosa and muscle weakness caused by Kearns-Sayre syndrome with a verified mtDNA deletion and haemolytic anaemia resulting from hereditary spherocytosis. (6) Initially suffering from dizziness. (7) Status epilepticus, dizziness, minor difficulties in learning and in mathematics at school (Holopainen et al. 1997). (8) Minor difficulties in learning speech and language (Holopainen et al. 1997). (9) Generalised tonic clonic seizures; EEG showed generalised epileptic activity (Bakkeren et al. 1984; Braakhekke et al. 1987). (10) Slight white matter hyperintensity (Van Kullenburg et al. 1999). (11) Delayed development of speech and hyperactivity (Van Gennip et al. 1981; Berger et al. 1984). (12) Suspected of having monoplegia. (13) Mild dysmorphic features (low set and posteriorly rotated ears, high arched palate) were noted. The child was

in a state of unconsciousness with no response to verbal, sensory or physical stimuli; only massive motor reaction to pain was noted. There was no reaction to light with preserved corneal reflexes. Bilateral optic atrophy was present. Superficial abdominal and anal reflexes were absent. Tetraparesis and flexion contractures were present. EEG revealed severe dysrhythmia and multifocal sharp/ spike waves. Generalised loss of white and grey matter and diffuse cerebral atrophy was observed on cranial MRI. Died at the age of 8 years. (14) Upper airway infection, Bartter's syndrome (hypokalaemia, 2.5 mM), enuresis nocturna. (15) Paroxysmal vertigo with attacks of 30 min to 2 h, hemiparaesthesia, diplopia, hemiparesis, headache. (16) Severe behavioural disorder and delayed development of speech, episodic tempers, chronic hypernatremia, spastic diplegia, partial agenesis of corpus callosum, delayed myelination, hamartoid cerebral lesion, epileptic discharges, megalocorneae, hypopigmentation of the fundus and pallor of optic discs (Brockstedt et al. 1990). (17) ALTE with hypothermia (29°C) and shock. (18) Pseudostrabismus, ugly formed, coarse, notched and fawn-coloured teeth (Van Gennip et al. 1987)

observed for the DPD activity in mononuclear cells and fibroblasts (Meinsma et al. 1995; Van Gennip et al. 1995; Vreken et al. 1997b). Analysis of the prevalence of the various mutations among DPD patients has shown that the G→A point mutation in the invariant splice donor site leading to the skipping of exon 14 is by far the most common, whereas the other six mutations are less frequently observed (Table 1). In addition, there appears to be some homogeneity for the G→A point mutation in Northern Europe, since homozygosity for this mutation has been observed in 9 individuals from Denmark, Sweden, Finland

and The Netherlands (Table 2). Furthermore, the majority of the DPD patients are of Dutch origin (55%); this probably reflects the fact that, in The Netherlands, screening for inborn errors of pyrimidine degradation is part of an intensive screening program for inborn errors in general.

So far, the frequency of these mutations in a normal population is not known. Based on the analysis of the DPD activity in various populations, it has been estimated that the frequency of heterozygotes might be as high as 3% (Gonzalez and Fernandez-Salguero 1995). Fortunately, the G→A point mutation destroys a unique *MaeII* restriction



**Table 3** Onset phenotype of patients with DPD deficiency (AS asymptomatic, – not available for analysis)

Patient	Genotype	Age of onset of symptoms	Age at diagnosis	Epilepsy in family	Consanguinity	Treatment <sup>a</sup>
1	$\Delta$ EX14/ $\Delta$ EX14	At birth	2 years, 1 months	No	No	
2	$\Delta$ EX14/ $\Delta$ EX14	3 years	7 years	Yes	No	1
3	$\Delta$ EX14/ $\Delta$ EX14	1–2 years	2 years	No	No	
4	$\Delta$ EX14/ $\Delta$ EX14	At birth	At birth	–	No	
5	$\Delta$ EX14/ $\Delta$ EX14	10 years 2 months	14 years, 6 months	No	No	
6	$\Delta$ EX14/ $\Delta$ EX14	6 months	6 years, 2 months	No	No	2
7	$\Delta$ EX14/ $\Delta$ EX14	Childhood	–	–	–	
8.1	$\Delta$ EX14/ $\Delta$ EX14	7 years, 5 months	8 years, 1 month	Yes	No	3
8.2	$\Delta$ EX14/ $\Delta$ EX14	Childhood	4 years, 5 months	Yes	No	
9.1	$\Delta$ TCAT/ $\Delta$ TCAT	3 years	6 years	Yes	Yes	4
9.2	$\Delta$ TCAT/ $\Delta$ TCAT	AS	4 years	Yes	Yes	
9.3	$\Delta$ TCAT/ $\Delta$ TCAT	12 years	28 years	Yes	Yes	5
10	C29R/C29R	1 years	3 years	No	No	
11	V995F/V995F	1 years, 6 months	3 years, 1 months	No	No	
12.1	$\Delta$ C1897/ $\Delta$ C1897	–	30 years	Yes	No	
12.2	$\Delta$ C1897/R235 W	4 months	6 years, 2 months	Yes	Yes	
13	C29R,R886H/ C29R, R886H	8 years	8 years	–	–	
14.1	C29R/ $\Delta$ EX14	17 years, 6 months	18 years, 2 months	Yes	No	
14.2	C29R/ $\Delta$ EX14	AS	26 years, 4 months	Yes	No	
15	$\Delta$ EX14/ $\Delta$ TCAT	6 months	1 years, 3 months	No	No	6
16	$\Delta$ EX14/C29R	5 weeks	5 weeks	No	No	
17	$\Delta$ EX14/?	At birth	1 years, 10 months	–	No	

<sup>a</sup> (1) Responded to valproate. (2) No response to treatment with valproate. Good response to carbamazepine and lamotrigine. After interruption of medication, symptoms of complex partial epilepsy recurred. After reintroduction of medication, the patient is again seizure free

(Christensen et al. 1998). (3) Good response to oxcarbazepine (Holopainen et al. 1997). (4) Phenobarbital and ethosuximide (Bakkeren et al. 1984; Braakhekke et al. 1987). (5) Phenytoin and phenobarbitone (Bakkeren et al. 1984). (6) Responded to pipamperon, clonazepam.

site present in an amplified genomic DNA fragment encompassing the skipped exon and its flanking sequences, allowing the rapid screening of this mutation in patients (Vreken et al. 1996; Wei et al. 1996). Screening for the presence of the G→A splice site mutation in a limited number of individuals of various nationalities has revealed heterozygosity for this mutation in 1% of the Finnish population (180 alleles analysed) and none in British (60 alleles), Japanese (100 alleles), African-American (210 alleles) or Dutch (100 alleles) populations (Wei et al. 1996, 1998; Vreken et al. 1996). Initially, an allele frequency of 5% was found for the G→A splice site mutation in Taiwanese subjects (72 alleles analysed; Wei et al. 1996). However, in subsequent studies, the G→A splice site mutation has not been detected in a larger group of Taiwanese subjects (262 alleles analysed; Wei et al. 1998). In addition, neither the splice-site nor the 1897delC has been detected in 60 Caucasian subjects (Ridge et al. 1998).

### Clinical phenotype of patients with DPD deficiency

A thorough investigation of clinical symptoms in the patients with complete DPD deficiency has shown a considerable variation in the clinical presentation among these pa-

tients (Table 2). Convulsive disorders (seizures and epileptic insults), motor retardation and mental retardation have been observed in approximately half of the patients, whereas growth retardation, microcephaly, autism and dysmorphism are less frequently observed. In this respect, it is worthwhile to note that, in 5 out of the 17 families, a history of convulsions is present (Table 3). A conspicuous finding has been the observation that five patients presented with ocular abnormalities. In one of the patients, the ocular symptoms are part of Kearns-Sayre syndrome with a verified mtDNA deletion but, even if this case is omitted, the four cases indicate a possible association with DPD deficiency. To our knowledge, a possible association of ocular abnormalities with DPD deficiency has not previously been recognised. Surprisingly, one patient suffers from a combined deficiency of DPD and the relatively rare syndrome of Kearns-Sayre, whereas one other patient suffers from a combined deficiency of DPD and Bartter's syndrome. The phenotypic variability of DPD deficiency is demonstrated by the finding that two asymptomatic patients have been identified and that seven other patients do not show the previously mentioned neurological and developmental abnormalities. However, 6 out of these 7 patients presented with other (neurological) abnormalities, such as lethargy, dizziness, monoplegia, an acute life-threatening event (ALTE) with hypothermia, and minor difficulties in learning speech and

language. This latter neurological abnormality has also recently been described in two other families with otherwise healthy DPD-deficient siblings (Henderson et al. 1995; Fernandez-Salguero et al. 1997). As discussed below, a number of the aforementioned symptoms might be explained by the altered  $\beta$ -alanine, uracil and thymine homeostasis in patients with DPD deficiency.

In all patients, the onset of the clinical phenotype occurred during childhood with the majority of the patients showing clinical abnormalities during the first years of life (Table 3). In general, a good response was noted when patients with convulsions/epileptic attacks were treated with anti-epileptic medication.

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### Phenotype-genotype relationship

An investigation into whether a correlation exists between the genotype and the observed clinical phenotype has revealed that all patients homozygous for the G→A splice site mutation presented with clinical abnormalities ranging from very mild (patients 5 and 8.2) to quite severe (Table 2). However, in a family presenting three subjects homozygous for the four-base deletion (295–298delTCAT), clinical abnormalities were clearly present in the index patient (no. 9.1) and his mother (no. 9.3), whereas the same genotype did not lead to a clinical phenotype in the brother of the patient (no. 9.2; Braakhekke et al. 1987; Vreken et al. 1997a). In addition, neurological abnormalities were seen in a patient who proved to be compound heterozygous for C29R/ $\Delta$ EX14 (patient 14.1) but not in her sister (patient 14.2). Thus, the absence of a characteristic phenotype in some patients with complete DPD deficiency indicates that other factors play an important role in the clinical manifestation of this disorder. In this respect, it has been shown that the involvement of a second gene closely linked to the DPD gene on chromosome 1 in the expression of the clinical symptoms is not very likely (Fernandez-Salguero et al. 1997).

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### $\beta$ -Alanine homeostasis

$\beta$ -Alanine is a structural analogue of  $\gamma$ -aminobutyric acid (GABA) and glycine, which are the major inhibitory neurotransmitters in the central nervous system. It has been suggested that  $\beta$ -alanine itself is involved in synaptic transmission, since  $\beta$ -alanine is present in central nervous tissue (Martin Del Rio et al. 1977), is released upon depolarisation by high potassium (Sandberg and Jacobson 1981) or electrical stimulation (Kihara et al. 1989), inhibits neuronal excitability (Choquet and Korn 1988; Mathers et al. 1990) and is removed from the extracellular fluid by a high affinity uptake system shown to operate in brain slices (Kontro 1983), cerebellar granule cells (Saransaari and Oja 1993), synaptosomal preparations (Zafra et al. 1984) and glial cells (Schon and Kelly 1975; Holopainen and Kontro 1986;

Mabjeesh et al. 1992). At least in chick spinal cord neurons and in mouse brain, it has been demonstrated that  $\beta$ -alanine activates both glycine and GABA<sub>A</sub> receptors with an efficacy similar to that for glycine and GABA, respectively (Choquet and Korn 1988; Horikoshi et al. 1988; Wu et al. 1993). Whether  $\beta$ -alanine is also able to bind to a unique receptor is still a matter of debate. Following synaptic release, GABA is transported into presynaptic endings and into glial cells where some of the neurotransmitter is metabolized by GABA transaminase. Most importantly,  $\beta$ -alanine has been shown to be a potent blocker of the uptake of GABA in glial cells (Mabjeesh et al. 1992). Reduced levels of  $\beta$ -alanine in patients with DPD deficiency might therefore have a profound effect on the degree of activation of the glycine and GABA<sub>A</sub> receptors and on GABA transport into glial cells. Since convulsions are often noted in patients with DPD deficiency, it is worthwhile to note that many anti-convulsant drugs act by potentiating the GABA-mediated inhibition in the nervous system and that GABA uptake blockers, such as  $\beta$ -alanine, possess profound anti-convulsant effects (Pfeiffer et al. 1996).

A conspicuous finding has been the presence of ocular abnormalities in 5 patients with a DPD deficiency. It has been suggested that, in addition to GABA,  $\beta$ -alanine is a neurotransmitter in the visual system (Sandberg and Jacobson 1981). Recently, a novel GABA receptor (GABA<sub>p1</sub>) has been detected in the retina; this receptor responds to glycine and  $\beta$ -alanine (Calvo and Miledi 1995). Therefore, an altered regulation of this GABA<sub>p1</sub> receptor by decreased  $\beta$ -alanine concentrations may also play a role in the observed ocular abnormalities.

The regulation of body temperature takes place in the central nervous system in the hypothalamus and is affected by neurotransmitters such as serotonin and GABA (Dhumal et al. 1974). In addition, after the administration of  $\beta$ -alanine to mammals, both hypothermia and hyperthermia have been reported (Gomez et al. 1978; Peters et al. 1987). Whether the occurrence of severe hypothermia in one of the DPD deficient patients might have been related to reduced levels of  $\beta$ -alanine is not yet known.

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### Pyrimidine bases

There is an increasing awareness that pyrimidines play an important role in the regulation of the central nervous system and that metabolic changes affecting the levels of pyrimidines may lead to abnormal neurological activity (Connolly et al. 1996). Indeed, the anti-convulsant effects of uridine in animals with experimentally induced seizures indicate that pyrimidine compounds may play an important role in regulating the activity of the nervous system (Roberts 1973). Moreover, disturbances of pyrimidine metabolism by anti-metabolites in cancer treatment are thought to be responsible for neurotoxicity presenting as seizures (Wiley et al. 1982).

In patients with complete DPD deficiency, a large accumulation of uracil and thymine and a moderate amount of

5-hydroxymethyluracil is present in urine, blood and cerebrospinal fluid (Bakkeren et al. 1984; Van Gennip et al. 1993, 1994, 1997). Altered uracil and thymine homeostasis in addition to altered pools of their downstream products might conceivably underlie some of the clinical abnormalities of patients with DPD deficiency. Parenteral administration of low doses of uracil and thymine to mice has been shown to increase their spontaneous activity, whereas their activity is decreased at higher doses of these pyrimidine bases (Krooth et al. 1978). Very high concentrations of both uracil and thymine have even proved to be lethal. Most interestingly, the intraperitoneal administration of the degradation products has no effect on spontaneous activity, although the subcutaneous administration of high-doses of  $\beta$ -alanine leads to a moderate depression of activity (Krooth et al. 1978). The phenomena observed after the administration of pyrimidine bases to mice are in line with those observed in tumour patients treated with high doses of thymidine, which in vivo is rapidly catabolized into thymine. The central nervous system toxicities encountered in these patients include somnolence, headache, visual illusions and memory impairment (Chiuten et al. 1980). Thus, the elevated levels of pyrimidine bases in patients with complete DPD deficiency might underlie both the occurrence of lethargy and hyperactivity, as observed in two of the DPD-deficient patients.

### Concluding remarks

In this paper, we have shown that a wide spectrum of clinical abnormalities, ranging from very mild to quite severe, are encountered in patients with complete DPD deficiency. The finding that some patients with complete deficiency of DPD do not present with any clinical abnormalities suggests that additional factors are involved determining the clinical outcome. Reasoning along these lines, we have speculated that an altered  $\beta$ -alanine homeostasis might be compatible with the various neurological symptoms seen in these patients. Since  $\beta$ -alanine can also be derived from dietary sources, such as carnosine, anserine and balanine (Van Gennip et al. 1993), the availability of these substances and of carnosinase activity in the relevant compartments may also affect the concentration of  $\beta$ -alanine in these compartments. In addition, we feel that not only a decreased level of  $\beta$ -alanine itself, but also the relative concentration of  $\beta$ -alanine compared with other neurotransmitters, such as GABA, might determine whether a clinical phenotype will emerge. Therefore, DPD deficiency is probably a necessary, but not a sole, prerequisite for the onset of a clinical phenotype. Nevertheless, the diagnosis of DPD deficiency is of paramount importance, not only in order to avoid severe toxicity during the 5FU treatment of tumour patients, but also in order to gain further insight into the relationship between the biochemical abnormalities and the onset of a clinical phenotype.

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