

# Loss of function *CYP24A1* mutations in patients with hypercalcemia and low PTH level: an autosomal dominant or recessive trait



Arnaud Molin<sup>a</sup>, Roseline Baudouin<sup>b</sup>, Nadia Coudray<sup>a</sup>, Marie-Lucille Figueres<sup>c</sup>, Glennville Jones<sup>d</sup> & Marie-Laure Kottler<sup>a</sup>

<sup>a</sup>CHU, Caen, France, <sup>b</sup>CHU, Bordeaux, France, <sup>c</sup>CHU, Nantes, France, <sup>d</sup>Queen's University, Kingston, Canada



## Abstract

Homozygous or compound heterozygous mutations of the gene *CYP24A1* coding vitamin D 24-hydroxylase have recently been reported to cause Idiopathic Infantile Hypercalcemia (IIH) due to increased intestinal absorption of calcium [1]. However, an autosomal dominant transmission with partial penetrance of the trait was also suggested [2]. So far, only case-reports have been published. Frequency of *CYP24A1* mutations in hypercalcemic patients remains unknown.

Here we describe a cohort of patients presenting with hypercalcemia and low PTH rate, to better define the phenotype of patients who should benefit of *CYP24A1* genetic screening and to evaluate the frequency of the disease.

In addition, we also show that simultaneous measurement of vitamin D metabolites by liquid chromatography-tandem mass spectrometry (LC-MS/MS) is a valuable screening tool for these patients.

## Objectives

- 1) To evaluate the frequency of *CYP24A1* mutation in hypercalcemic patients with low PTH rate
- 2) To study the impact of *CYP24A1* heterozygous mutation on calcium metabolism
- 3) To highlight the usefulness of LC/MSMS for vitamin D metabolites measurement

## References

- 1-Schlingmann *et al.* 2011 *N Engl J Med* 365:410-21
- 2-Tebben P *et al.* 2011, *J Clin Endocrinol Metab* 97:E423-E7
- 3-Kaufmann M *et al.*, 2014. *J Clin Endocrinol Metab* 99:2567-2574.

## Methods

### Patients

We studied 72 index cases presenting with hypercalcemia (>2.6 mmol/L) and low PTH levels (<20pg/mL) and 22 heterozygous relatives.

### Biochemical parameters

Data on clinical symptoms, renal ultrasound examination and biological explorations were collected at the time of the diagnosis, or retrospectively using records from hospitals or primary care physicians.

### Simultaneously assay of vitamin D metabolites

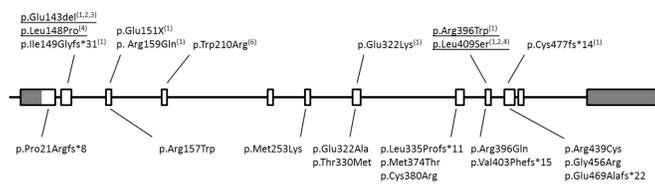
LC-MS/MS analysis was performed at the time of molecular diagnosis as previously described [3] including 25-OH D<sub>3</sub> and 24,25-(OH)<sub>2</sub>D<sub>3</sub> using 100µl of serum; results are expressed as a ratio of 25-OH D<sub>3</sub>:24,25-(OH)<sub>2</sub>D<sub>3</sub>. Values under 25 indicated no defect in 24-hydroxylase activity and were considered as normal.

### Molecular analysis

11 coding exons of *CYP24A1* and their intron-exon junctions were sequenced as previously described (Castanet *et al.* 2013). New variations of sequence interpreted according to pathogenicity prediction programs (PolyPhen-2, Align-GVGD, MutationTaster, SIFT).

## Results 1

We identified 25 patients (35%) harboring mutations in coding sequence of *CYP24A1*: 20 patients (28%) with bi-allelic mutations (10 homozygous, and 10 compound heterozygous) and 5 children with heterozygous mutation (7%). All were neonates, under 2 weeks (range 1 to 13 days). In these patients, hypercalcemia was found during routine exams performed for another pathology: prematurity, growth retardation, infection or apnea. None presented with renal pathology

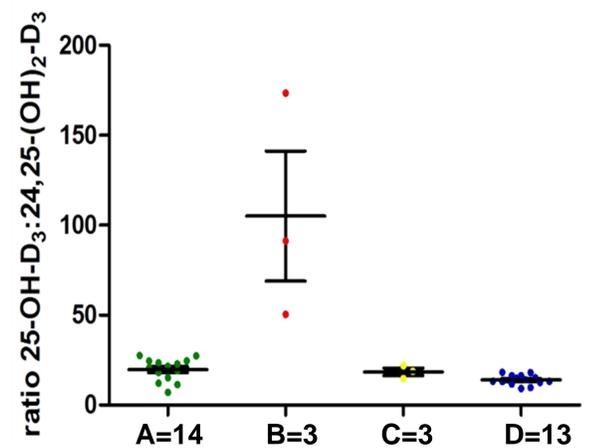


(1) Schlingmann *et al.* (2) Dauber *et al.* (3) Dinour *et al.* (4) Nesterova *et al.* (5) Colussi *et al.* (6) Meusburger *et al.*

Positions of observed mutations in the human *CYP24A1* gene.

## Results 2

Patients with bi-allelic *CYP24A1* mutations (group B) exhibit a dramatic increase in 25-OH D<sub>3</sub>:24,25-(OH)<sub>2</sub>D<sub>3</sub> ratio (105 [48.8-173.4]) providing evidence "in vivo" for the loss of *CYP24A1* enzyme activity. By contrast, 25-OH D<sub>3</sub>:24,25-(OH)<sub>2</sub>D<sub>3</sub> ratio remains within the normal range (R= 19.7[7-27.5]) in probands without *CYP24A1* mutation (group A).



A : no mutation      C : heterozygous index  
B : bi-allelic mutation      D : heterozygous relatives

In patients heterozygous for *CYP24A1* mutations, probands (group C) as well as relatives (group D), simultaneous assay of both 25-OH D<sub>3</sub> and 24,25-(OH)<sub>2</sub>D<sub>3</sub> provides evidence for the presence of normal *CYP24A1* activity with a 25-OH D<sub>3</sub>:24,25-(OH)<sub>2</sub>D<sub>3</sub> ratio within the normal range (R=13.7 [9.3-18]).

## Conclusions

We identified mutation in *CYP24A1* as a major cause of hypercalcemia associated to low PTH level.

We confirm the accuracy and effectiveness of a novel blood test estimating the ratio between relevant vitamin D metabolites 25-OH D<sub>3</sub> and 24,25-(OH)<sub>2</sub>D<sub>3</sub>. This test constitutes a useful screening tool.

We suggest that in patients with *CYP24A1* haplo-insufficiency, vitamin D supplementation associated with a low renal function could trigger hypercalcemia and hypercalciuria.