

TWO FRENCH FAMILIES WITH VITAMIN D DEPENDENCY RICKETS TYPE 1B HARBOR HOMOZYGOUS RECESSIVE EXPRESSION OF CYP2R1 MUTATIONS L99P and G42_L46delinsR.

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Introduction: Mutations of *CYP2R1* (11p15.2) encoding the main vitamin D 25-hydroxylase have been associated with a rare recessive autosomal form of rickets, also called vitamin D dependency rickets type 1B (VDDR-1B) (Cheng et al. 2004).

We describe rickets & loss-of-function *CYP2R1* mutations in 6/10 individuals tested from two unrelated families: five patients in family 1 (F1) with homozygous L99P mutations and one boy in family 2 (F2) with a novel homozygous mutation G42_L46delinsR.

Calcifediol (25-OH D₃) therapy resulted in complete normalization of biochemical and bone defects.

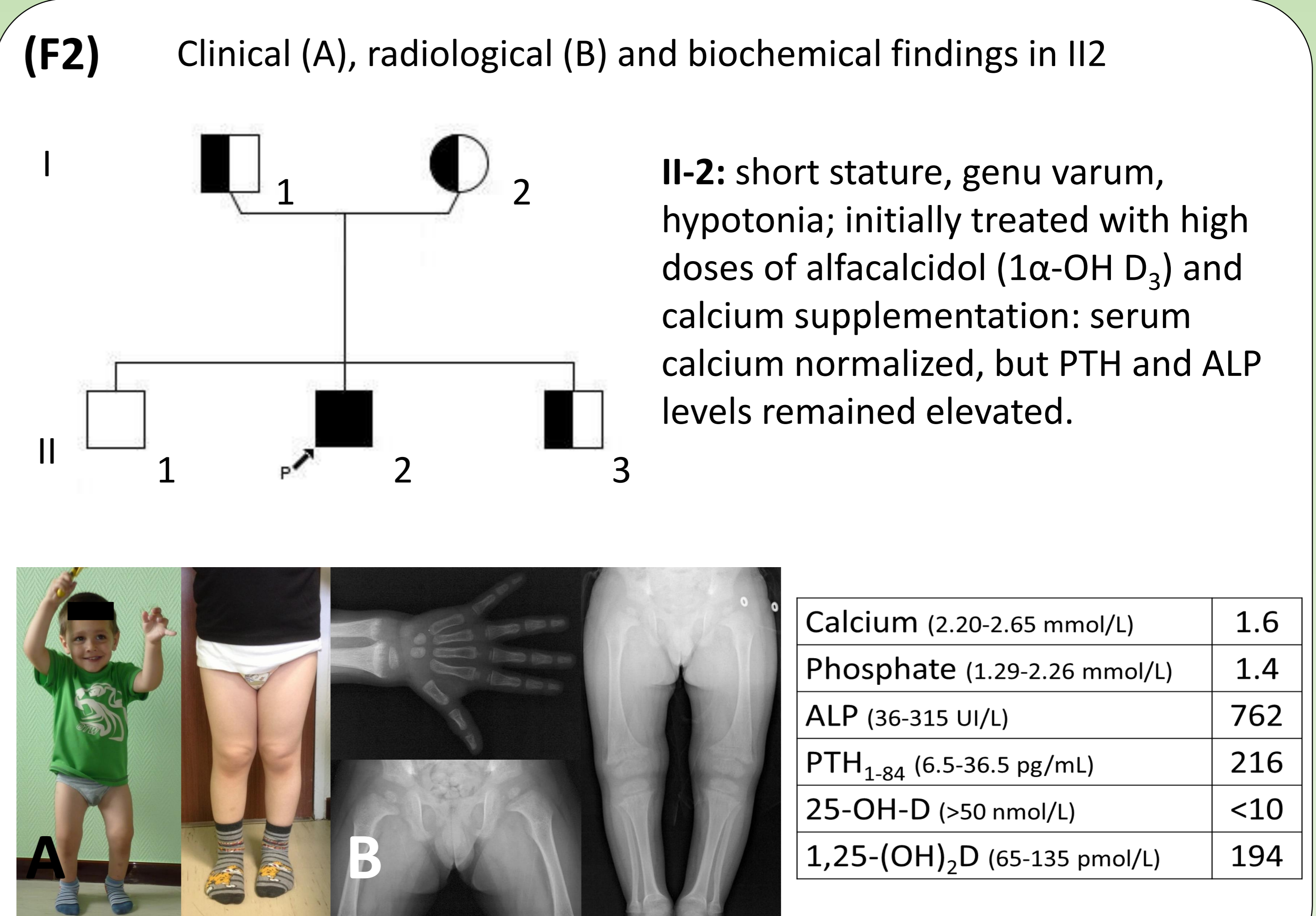
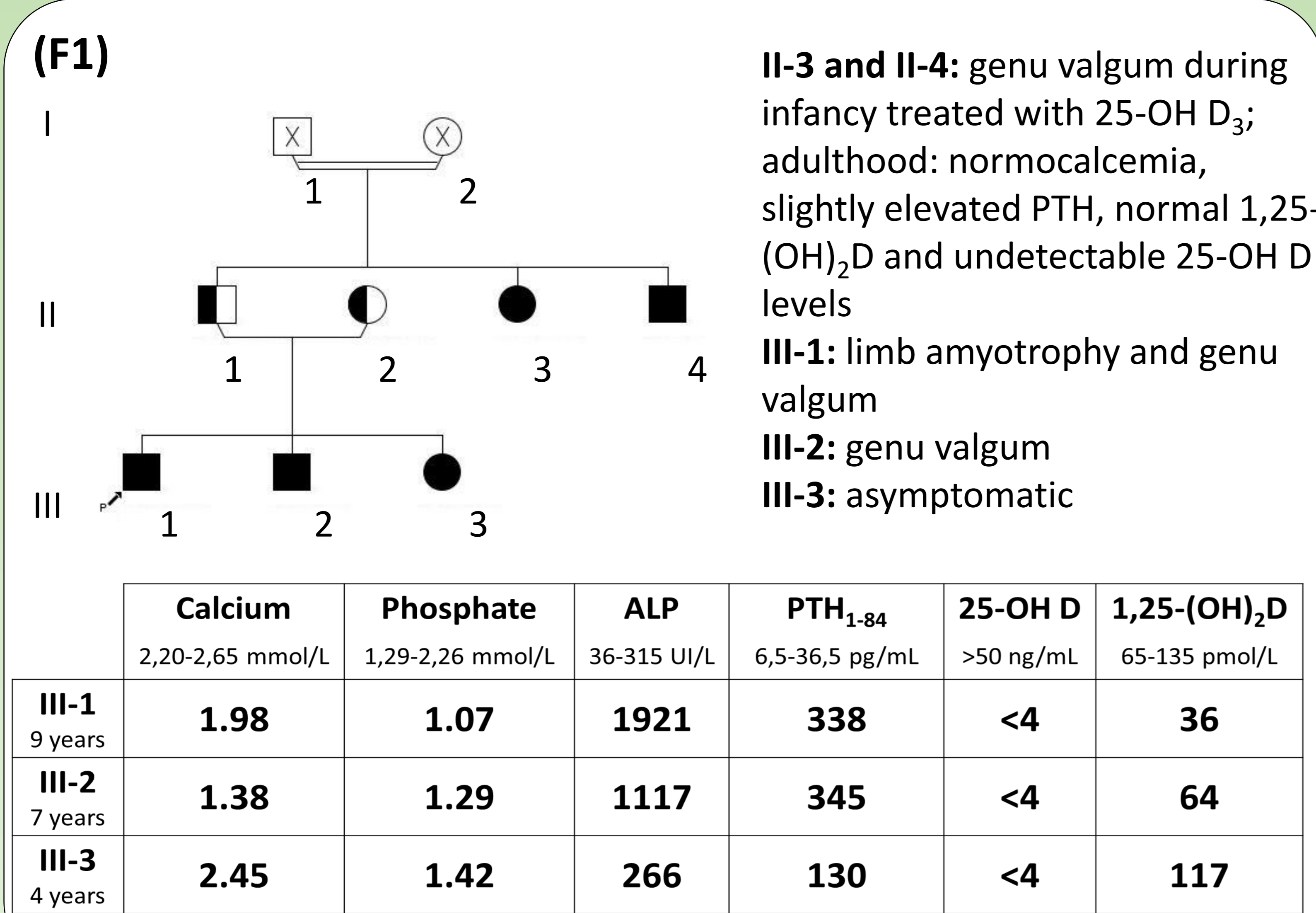
Methods:

Clinical and routine laboratory explorations: (Ca and P, PTH, alkaline phosphatase (ALP) and vitamin D metabolites) Data were collected at the time of the diagnosis, retrospectively and prospectively, using records from hospitals.

Molecular analysis: *CYP2R1* exons 1-5 and their intron-exon junctions were sequenced using standard procedures on a Beckman Coulter DNA Sequencer.

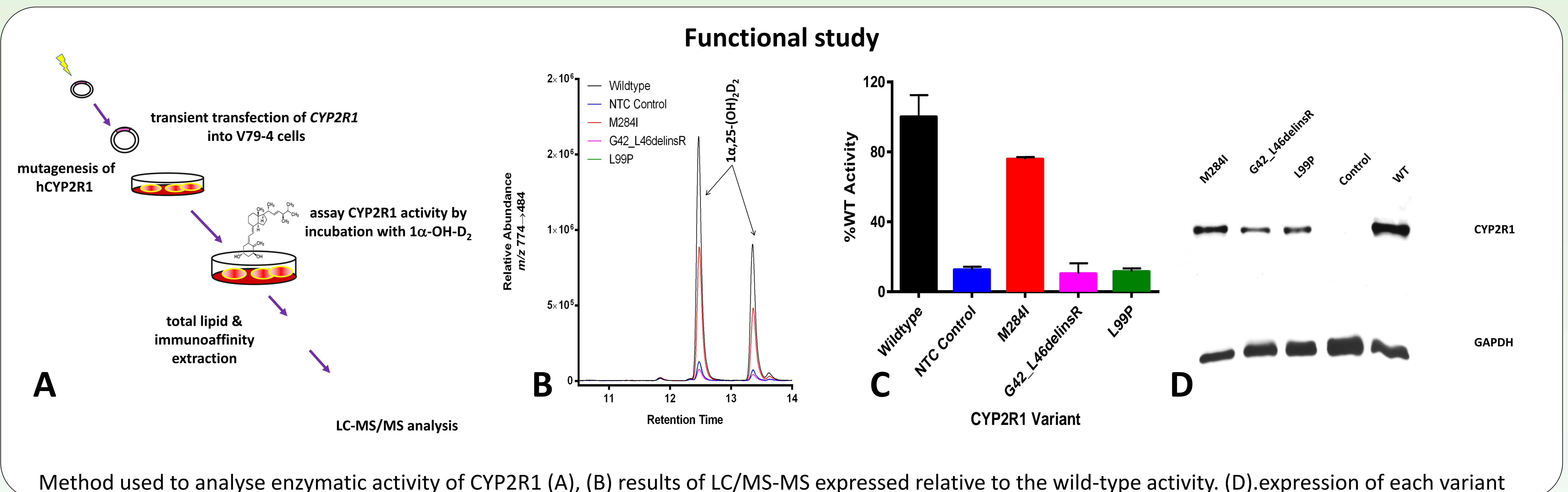
Functional study: The mutations, as well as another variant M248I found in the French population, were recreated and tested using an in vitro mammalian expression system described previously (*JBiolChem* 286:28729).

Results : radiological findings, were typical of rickets. All affected children presented with similar biochemical findings : hypocalcemia, hypo/normo phosphatemia, high PTH and ALP levels. While serum 1,25-(OH)₂D levels were within the normal range (F1 : III3) or even high (F2 : II2), 25(OH)D was undetectable:



Molecular analysis: Sequence analysis of *CYP2R1* in the probands revealed the previously described c.296T>C (L99P) mutation in the exon 2 in F1, and a novel c.124_138delinsCGG (p.G42_L46delinsR) in the exon 1 in F2. Mutations were in a homozygous state in both probands, and in a heterozygous state in their parents. The brother and the sister of the probands in F1 also carried the L99P mutation in a homozygous state.

Functional study: L99P and G42_L46delinsR showed <5% of wild type *CYP2R1* enzyme activity and are presumed to be loss-of-function mutations, while the M284I variant had 75% activity and is thus likely a polymorphism.



CYP2R1 deficiency should be investigated in patients presenting rickets with low/undetectable 25-OH-D serum concentration even so 1,25(OH)₂D concentration is normal. The precise identification of the genetic defect allowed an appropriate therapy resulting in complete normalization of bone defect and biochemical parameters.