ORIGINAL RESEARCH



Does Genotype–Phenotype Correlation Exist in Vitamin D-Dependent Rickets Type IA: Report of 13 New Cases and Review of the Literature

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Abstract

Vitamin D-dependent rickets type IA (VDDR-IA) is caused by biallelic mutations in CYP27B1. Data regarding genotypephenotype correlation in VDDR-IA are scarce. Here, we aimed to investigate clinical/genotypic features and long-term follow-up of 13 new cases with VDDR-IA and genotype-phenotype correlation of reported cases in the literature. Thirteen patients with VDDR-IA were evaluated. Eight patients had reached their final height at the time of the study and, for whom, long-term outcome data were analyzed. Further, all VDDR-IA patients in the literature (n:183) were analyzed and clinical-genetic features were recorded. The median age of diagnosis was 2.55 ± 1.13 (1.0–12) years. Initial diagnoses before referral to our clinic were nutritional rickets (n:7), hypophosphatemic rickets (n:2), and pseudohypoparathyroidism (n:1). All had biochemical evidence suggestive of VDDR-IA; except one with elevated 1,25(OH)₂D3 and another with hyperphosphatemia, in whom pseudohypoparathyroidism was excluded with molecular tests. Combined analyses of our cohort and other series in the literature demonstrated that three most common CYP27B1 mutations are p.F443Pfs*24, c.195+2T>G, and p.V88Wfs*71. In Turkish population, p.K192E mutation along with the former two is the most common mutations. Comparison of clinical features demonstrated that c.195+2T > G mutation causes the most severe and p.K192E mutation causes the least severe phenotype with respect to age and height at presentation and calcitriol requirement. We found a clear genotype–phenotype correlation in VDDR-IA, notably CYP27B1 intronic c.195+2T>G mutation causes a more severe phenotype with lower height SDS at presentation and, higher calcitriol requirement, while less severe phenotype occurs in p.K192E mutation.

Keywords Rickets · VDDR · CYP27B1 · 1a-hydroxylase · Hypocalcemia · Hyperphosphatemia

Introduction

Vitamin D is essential for bone mineral metabolism, maintenance of calcium, and phosphate homeostasis. The 1 α -hydroxyl group of calcitriol (1,25(OH)₂D3) is required for the binding of vitamin D to its nuclear receptor [1].

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² Department of Medical Genetics, Marmara University School of Medicine, Istanbul, Turkey Therefore, two inactive precursors of vitamin D: ergocalciferol (Vitamin D2) and cholecalciferol (Vitamin D3), need sequential 25- and 1α -hydroxylation to form $1,25(OH)_2$ D3. First, vitamin D is hydroxylated to 25-hydroxy vitamin D (25(OH)D₃) by several hepatic enzymes, mainly microsomal CYP2R1 in the liver, then enters mitochondria where it is further hydroxylated to the biologically active metabolite $1,25(OH)_2D3$ by the 25-OH vitamin D- 1α -hydroxylase in kidneys.

The hereditary defects in vitamin D metabolism and its function cause rare forms of rickets, also known as vitamin D-dependent rickets (VDDR) [2]. VDDR type 1 involves VDDR-IA (1 α -hydroxylase deficiency) and VDDR-IB (25-hydroxylase deficiency). VDDR type 2 involves vitamin D receptor (VDDR-IIA) and post-receptor (VDDR-IIB) defects [2]. More recently, *CYP3A4* mutation as a cause of VDDR type 3 (VDDR-III) by increasing inactivation

of vitamin D has been described [3]. VDDR-IA (OMIM; 264700) is the most common type of VDDR and is caused by biallelic inactivating mutations in *CYP27B1* gene which results in defective 1α-hydroxylation of 25(OH)D3 to 1,25(OH)₂D3 [4]. Despite the paucity of the studies, its prevalence is estimated as around 1-5/10.000 (https://www.orpha .net/consor/cgi-bin/OC_Exp.php?Lng=EN&Expert=28915 7). Patients with VDDR-IA appear normal at birth and usually present during the first two years of life with hypotonia, muscle weakness, growth failure, delay in walking, and bowing in legs [5]. Laboratory characteristics of the patients with VDDR-IA demonstrate typical calciopenic rickets with inappropriately normal or low levels of 1,25(OH)₂D3 and normal or increased concentrations of 25(OH)D3 [6–8].

To the best of our knowledge, over 180 patients with 81 mutations in the *CYP27B1* gene have been reported from different ethnic groups so far. These mutations span all exons of the gene and mostly include missense and nonsense changes (approximately 70% of *CYP27B1* mutations) (http://www.hgmd.cf.ac.uk/ac/index.php) (Fig. 1). However, data regarding genotype–phenotype correlation and long-term follow-up and outcome of patients with VDDR-I-A are scarce.

In this study, we report 13 new patients with VDDR-IA and analyzed the most common genotypes worldwide with extensive literature review. In addition, genotype-phenotype correlation with special emphasis on clinical presentations, response to treatment, and long-term outcome were investigated.

Subjects and Methods

Patients

Patients presenting with clinical and radiological findings of rickets with elevated serum alkaline phosphatase (ALP) and parathyroid hormone (PTH), normal or increased levels of 25(OH)D3, and inappropriately normal or low levels of 1,25(OH)₂D3 if available were included to the study. The diagnosis of VDDR-IA was suggested on clinical ground with unresponsiveness to high-dose cholecalciferol treatment and responsive to standard-dose calcitriol treatment. Eventually, the diagnosis was confirmed by genetic testing in all patients.



Fig. 1 Schematic presentation of *CYP27B1* (NM_000785.4) gene with its exons and introns showing all reported mutations. Arrows show identified variants at respective exons and introns. Mutations identified in Turkish cohorts underlined, and mutations identified in

our patients denoted in bold. Frameshift, splice site and missense/ nonsense mutations showed in upper, middle, and lower boxes, respectively

A comprehensive search of PubMed, using following keywords in the PubMed search query: "CYP27B1," "VDDR," and "Vitamin D-dependent rickets" was conducted to identify all the studies presenting data concerning the genotype-phenotype association and/or long-term follow-up of VDDR-IA patients. We limited our search to human studies. We excluded studies that report cases where the molecular diagnosis of VDDR-IA had not been established. We also checked the bibliographies of included studies for relevant articles. Data were extracted on to a pre-formatted table with details of the study, CYP27B1 variant, and number of patients that carry each individual variant. For the comparison of the clinical features of most common mutations, the clinical and laboratory findings were reviewed and statistical analyses were performed between the mutation groups having more than 20 homozygous patients. Patients with compound heterozygous mutations were excluded from the statistical analyses. The last search was completed on August 24, 2020.

Follow-Up

Serum calcium, phosphate, ALP, PTH concentrations, and urinary calcium excretion were evaluated once per month for the first three months after initiating calcitriol treatment. Thereafter, children were seen every three to four months as long as growth continued. After epiphyseal fusion, followup visits occurred every 6 months. Auxological measurements were obtained at each visit. Renal ultrasound was performed yearly to assess the presence of nephrocalcinosis. The patients were transferred to adult endocrinology clinic after the age of 18 years. The patients with closed epiphysis and no height changes for at least one year were accepted at final height.

Biochemical Analysis

Calcium, phosphate, and creatinine in serum and urine samples were analyzed with automated systems (AU 600, Beckman Coulter, USA). Serum ALP levels were measured by using AU 600 systems using methods based on the recommendations of the IFCC (*International Federation of Clinical Chemistry*) which is a quantitative colorimetric measurement using 2-amino-2-methyl-1-propanol as a buffer. Serum PTH was analyzed by Access Immunoassay Systems (Beckman Coulter, USA).

Plasma 25(OH)D3 measurements were performed by using Thermo Finnigan high-performance liquid chromatography (HPLC) system with ClinRep® HPLC Complete Kit (Recipe Chemicals–Instruments, Germany). Intra-assay and inter-assay coefficients of variability for the levels of 21.5 and 75.5 µg/L were 4.60% and 3.06%, and 3.60% and 4.40%, respectively. The quality evaluation of the method was reported [9].

The measurement of $1,25(OH)_2D3$ levels was performed in different validated commercial laboratories using ELISAbased kits throughout the years when the patients were presented to our clinic before the initiation of calcitriol who were not on calcitriol. For evaluation of the results, reference ranges provided by the laboratory had been used which were given separately for each patient in Table 2.

Radiological Evaluation

Bone mineral density of lumbar vertebrae was measured by dual energy X-ray absorptiometry (Hologic, Bedford, MA, USA).

CYP27B1 Gene Sequencing

Genomic DNA was extracted using a QIAamp DNA Mini Kit (QIAGEN, Germantown, MD USA) from peripheral blood lymphocytes according to standard protocols after informed consent. All coding exons and exon–intron boundaries of the *CYP27B1* gene (NM_000785.4) were sequenced via Sanger sequencing using an ABI 3130XL DNA Sequencer (Thermo Fisher Scientific, Waltham, MA, USA). The primers were designed using Primer3 software (primer3.ut.ee) [10].

Statistical Analysis

Statistical analyses were performed with GraphPad Prism 8.0 (GraphPad Software, Inc., San Diego, CA). The data were presented as mean \pm SEM (median; ranges). The median age of presentation, 1,25(OH)₂D3 levels, height standard deviation score (SDS), and doses of calcitriol were compared using the nonparametric Kruskal–Wallis test, and Mann–Whitney *U* test used for pairwise group comparisons. Spearman rank correlation was used to test associations between prior therapy and 1,25(OH)₂D3 levels at admission. All tests were two-tailed. A *p* value ≤ 0.05 was considered statistically significant.

Results

Clinical Characteristics

Thirteen (7 males) patients from 10 unrelated families (6 consanguineous parents, 4 nonconsanguineous parents but who were originated from nearby villages) were evaluated. The mean age of diagnosis was 4.8 ± 1.13 years (Median: 2.55; min: 1,0; max: 12). All patients had a history of receiving routine prophylactic vitamin D replacement (400 IU/day)

for the first year of life. Ten patients out of 13 came to medical attention within the first 24 months of life. Initial diagnoses of the patients were nutritional rickets (NR) (n=7), hypophosphatemic rickets (HR) (n=2), and pseudohypoparathyroidism (PHP) (n=1) before the referral to our clinic. Seven patients had a history of high-dose vitamin D intake (300,000–1,500,000 IU); two of them (Pt#1.1, Pt#4.1) had toxic levels of 25(OH)D3 (\geq 150 ng/mL) and three (Pt#3.1, Pt#6.1, Pt#10.1) had hypervitaminosis D (\geq 100 ng/mL). The distribution of complaints and available clinical data prior to referral are presented in Table 1. The mean time from the first medical visit to the diagnosis of VDDR-IA was 29.72 ± 10.14 months (Median: 11.10; min: 0.52; max: 96).

Upon referral to our clinic, all patients were diagnosed with VDDR-IA based on the biochemical and radiological findings of active rickets. Hypophosphatemia, elevated ALP and PTH, and normal/elevated 25(OH)D3 levels were detected in all patients except Pt#8.1. She was presented to the clinic with hypocalcemia (Ca: 1.59 mmol/L) and hyperphosphatemia (PO4: 2 mmol/L) and initially diagnosed as PHP. However, VDDR-IA was subsequently considered upon bone pain and the review of the X-rays on initial presentation which was consistent with rickets (Fig. 2). PHP was excluded also after demonstrating normal *GNAS* gene sequencing and Gs α levels (104%, N: 85–115%). All measured 1,25(OH)₂D3 levels were low or inappropriately normal, except mildly elevated in Pt#3.1.

The mean height SDS on admission was -1.71 ± 1.2 (Median: -1.23; min: -4.71; max: -0.25). Six patients had height SDS more than 1 SDS below their mid-parenteral height (MPH). The clinical and biochemical characteristics of the patients are summarized in Table 2.

Treatment and Follow-Up

All patients received calcitriol treatment, which was initiated either at the time of diagnosis (n:6) or after failure of treatment with vitamin D3 (n:7). The initial calcitriol dose

Table 1 Clinical features, working diagnoses, and treatment of the patients before referral to our institution and presenting complaints, renal ultrasound, and urinary calcium excretion at admission to our clinic

	First medical attention (age/initial diagnosis)	Age/presenting complaints (our institu- tion)	Treatment before admission	Renal USG (admission)	U-Ca /Cr (mg/ mg) (admis- sion)
1.1	6 months/NR	16 months/delay in walking, hypocalce- mic seizure	1,500,000 IU ^a Cholecalciferol	NP	NP
1.2	11 months/NR	12 months/delay in walking	1,050,000 IU ^a Cholecalciferol	NP	NP
2.1	17 months/HR	8.6 years/bowed legs	10.5 ng/kg/day Calcitriol 15.75 mg/kg/day ayP04	Microlithiasis (4 mm, unilateral)	0.05
2.2	11 months/HR	3.3 years/history of rickets in sibling	35.71 ng/kg/day Calcitriol 17.85 mg/kg/day P04	Normal	NP
3.1	11 months/NR	14 months/excessive Sweating, elevated serum ALP levels	600,000 IU ^a Cholecalciferol	Microlithiasis (2.5 mm, unilateral)	0.08
4.1	18 months/NR	21 months/excessive Sweating, delay in walking	300,000 IU ^a Cholecalciferol	NP	NP
5.1 ^b	5 years/VDDR	14.9 years/bowed legs, leg pain	None	Normal	NP
6.1	11 months/NR	21 months/delay in walking	600,000 IU ^a Cholecalciferol	NP	0.02
7.1	24 months/NR	12 years/bowed legs	300,000 IU Cholecalciferol	Normal	0.17
7.2	10,6 years/None	11.6 years/bowed legs, gait abnormality	None	Normal	0.02
8.1	36 months/PHP	8.8 years/leg pain	None	Normal	0.007
9.1	14 months/None	14 months/bowed legs, gait abnormality	None	NP	NP
10.1	24 months / NR	5.3 years/bowed legs	675,000 IU ^a Cholecalciferol	Normal	0.03

Ca Calcium, PO4 Phosphate, Cr creatinine, U Urinary, NR Nutritional Rickets, HR Hypophosphatemic rickets, VDDR Vitamin-D-dependent Rickets, PHP Pseudohypoparathyroidism, NP not performed

^aTotal calculated doses of cholecalciferol, which was given either as high-dose stoss therapy (300,000 IU for each time and repeated up to five times) or stoss therapy followed by daily high-dose (2000–3000 IU/day) therapy. Sum of all cholecalciferol received was taken as cumulative dose when presented

^bPatient had been diagnosed previously with VDDR at the age of 5 years and was on calcitriol therapy with inadequate compliance but lost to follow up until the age of 14 years, and at admission to our clinic he was not taking any medication for one year

Fig. 2 Radiologic features of the Pt#8.1 at admission showing active rickets with a loss of provisional calcification zone with irregular metaphysis, fraying and widening of growth plate (a1, b1), and after treatment with normal growth plate with mild osteopenia (a2, b2)

was 41.01 ± 7.59 ng/kg/day (median: 40; min: 14.84; max: 86.2) which was determined by severity of the disease and/ or hypocalcemia. The mean dose for calcitriol to achieve clinical and biochemical recovery was 59.53 ± 13.68 ng/kg/ day (median: 50; min: 12.19; max: 144.23). Clinical, biochemical, and radiological improvement was observed in all patients after initiation of calcitriol therapy. Normocalcemia achieved at 1-3 months of treatment, but normalization of PTH and ALP occurred at a later period (3-12 months). Upon referral to our clinic, one patient with calcitriol and phosphate treatment (Pt#2.1) and one patient who received high-dose vitamin D treatment (Pt#3.1) had microlithiasis on renal ultrasonography, which resolved after initiation of relevant treatment and discontinuation of phosphate therapy. None of the patients developed microlithiasis or nephrocalcinosis during the follow-up.

The onset of walking documented in four children (median age of 14.2 months; range 12–21) who had not started walking before the initiation of treatment. In these children, the median duration until to walk after initiation of calcitriol treatment was 6 months (range: 3–9 months) and the patients could walk at a median age of 20.75 months (range: 16.9–27 months).

The mean duration of calcitriol treatment was 9.45 ± 7.17 years (median: 7.15; min: 1.4; max: 25.16). The mean height SDS change from baseline to last visit was 0.73 ± 0.39 SDS (median:1.04; min: -1.91; max: 2.82). Ten patients had improved height SDS during follow-up, in whom, the mean increase in height SDS from baseline to last follow-up visit was 1.33 ± 0.26 SDS

(median: 1.19; min: 0.25; max: 2.82). In 4 patients with no height SDS improvement during follow-up, two patients (#1.2, #4.1) received treatment in the first two years of life, and they were compliant; however, their parents were short and the final height SDSs were within their MPH SDSs. The other two patients (#7.1 and #7.2) have been diagnosed at older ages (12 and 11.6 years) and noncompliant to treatment.

On the long-term follow-up, eight patients reached adult height. Their mean final height SDS was -1.38 ± 0.46 (Median: -1.39; min: -2.91; max: 0.62), MPH SDS was -1.12 ± 0.39 (Median: -1.50; min: -2.46; max: 0.86). Of these, three patients were fully compliant to treatment and had the highest BMD *z*-scores among patients reaching adult height. Five patients either stopped treatment (n:3) or declared to skip the drug doses (n:2). Lumbar BMD *z*-scores of the patients were -0.14 ± 0.42 (Median: 0.05; min: -1.9; max: 1.7). The clinical characteristics of eight patients who reached final height are summarized in Table 3.

Sequence Analysis of the CYP27B1 Gene

All of the patients had homozygous mutations in *CYP27B1* gene. We revealed a homozygous seven-nucleotide duplication (c.1319_1325dupCCCACCC) causing frameshift mutation (p.F443Pfs*24) in 8 patients from 6 unrelated families. A homozygous missense c.574 A > G (p.K192E) mutation was detected in three patients from two unrelated families. Homozygous c.590G > A (p.G197D) and c.1215 + 2T > A

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no consanguinity ¹ 5.1 ^g Rize / erzurum 0.86 no consanguinity ^f 0.38 6.1 Erzurum 0.38 no consanguinity ^f - 1.61 7.2 - 1.55 7.2 - 0.53 8.1 Bingol consanguineous - 0.36 9.1 Bingol consanguineous - 0.36 10.1 ^j Bingol consanguineous - 1.68		Trabzon	- 2.46	Σ	1.8	- 2.53	2.17	0.74	2495	480	169.0	%	
5.1 ^g Rize / erzurum0.86no consanguinity0.386.1Erzurum0.387.1Bingol0.387.1Bingol- 1.617.2- 1.55- 1.558.1Bingol consanguinity ^f - 1.559.1Bingol consanguineous- 0.539.1Bingol consanguineous- 0.5610.1 ^j Bingol consanguineous- 1.68		no consanguinity [†]			15.9	- 2.08	2.49	1.16	366	12	23	Ι	
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 6.1 Erzurum no consanguinity^f 0.38 7.1 Bingol - 1.61 7.2 - 1.55 7.2 - 1.55 8.1 Bingol consanguineous - 0.36 9.1 Bingol consanguineous - 0.36 10.1^j Bingol consanguineous - 1.68 		no consanguinity			24.2 ^h	- 0.36	2.22	0.58	230	199	31.9	I	
no consanguinity ¹ - 1.61 7.1 Bingol - 1.65 7.2 - 1.55 8.1 Bingol consanguineous - 0.53 9.1 Bingol consanguineous - 0.36 10.1 ^j Bingol consanguineous - 1.68	_	Erzurum	0.38	Σ	1.8	- 1.04	1.97	0.74	9251	877	147.5	× 8	
 7.1 Bingol - 1.61 no consanguinity^f - 1.55 7.2 - 1.55 8.1 Bingol consanguineous - 0.53 9.1 Bingol consanguineous - 0.36 10.1^j Bingol consanguineous - 1.68 		no consanguinity ^f			18.1	0.29	2.42	1.22	271	53	26.1	I	
 7.1 Bingol - 1.61 no consanguinity^f - 1.55 7.2 - 1.55 8.1 Bingol consanguineous - 0.53 9.1 Bingol consanguineous - 0.36 10.1^j Bingol consanguineous - 1.68 					21.2h	0.62	2.14	0.51	982	323	24.8	I	
no consanguinity ¹ – 1.55 7.2 – 1.55 8.1 Bingol consanguineous – 0.53 9.1 Bingol consanguineous – 0.36 10.1 ^j Bingol consanguineous – 1.68	_	Bingol	- 1.61	ц	12	- 1.15	2.14	0.74	2190	237	49.0	21	Exon 3
 7.2 - 1.55 8.1 Bingol consanguineous - 0.53 9.1 Bingol consanguineous - 0.36 10.1^j Bingol consanguineous - 1.68 		no consanguinity ¹			15.6	- 2.6	2.29	0.93	207	108	24.5	I	c.574 A > G (p.K192E)
 7.2 - 1.55 8.1 Bingol consanguineous - 0.53 9.1 Bingol consanguineous - 0.36 10.1^j Bingol consanguineous - 1.68 					18.31	- 2.91	2.02	0.80	154	377	23.6	I	
 8.1 Bingol consanguineous - 0.53 9.1 Bingol consanguineous - 0.36 10.1^j Bingol consanguineous - 1.68 			- 1.55	Σ	11.6	-0.72	2.24	0.96	1048	203	61.1	58.2	
 8.1 Bingol consanguineous - 0.53 9.1 Bingol consanguineous - 0.36 10.1^j Bingol consanguineous - 1.68 					14.6	- 1.36	2.27	1.70	166	45	45.5	I	
 8.1 Bingol consanguineous - 0.53 9.1 Bingol consanguineous - 0.36 10.1^j Bingol consanguineous - 1.68 					17.2'	- 2.37	2.34	1.32	102	193	40.4	I	
9.1 Bingol consanguineous – 0.36 10.1 ^j Bingol consanguineous – 1.68	_	Bingol consanguineous	- 0.53	ц	8.8	- 0.94	1.59	2.00	710	362	33.8	I	
 9.1 Bingol consanguineous - 0.36 10.1^j Bingol consanguineous - 1.68 					11.9	- 0.32	2.37	1.61	347	58	31.0	I	
10.1 ^j Bingol consanguineous – 1.68	_	Bingol consanguineous	- 0.36	Μ	1.2	- 2.46	2.09	0.70	3324	586	43.1	23	Exon 4
10.1 ^j Bingol consanguineous – 1.68					2.6	- 1.2	2.44	1.83	403	73	15.5	I	c.590G>A (p.G197D)
	1 [.]	Bingol consanguineous	- 1.68	ц	5.3	- 4.71	1.99	0.93	3585	1006	138.8	I	Intron 7
					12.4	- 1.64	2.56	1.19	89	26	42.2	I	c.1215 + 2T > A
					16.1	- 1.89	2.24	0.77	144	257	27.1	1	(/ C. SIMACTI)

Table 2 (continued)
MPH mid-parental height, Ca calcium, P phosphorus, ALP alkaline phosphatase, PTH parathyroid hormone
^a Age-specific normal ranges (mmol/L); 0–5 months: 1.7–2.7; 6–12 months: 1.6–2.5; 1–5 years: 1.4–2.0; 6–12 years: 1.2–1.9; >13 years: 0.7–1.4
^b Age- and sex-specific normal ranges (IU/L); 0–14 days: 83–228; 15 days-1 years: 123–377; 1–10 years: 97–362; 10–13 years: 92–418; 13–15 years: 59–275 (female), 88–43 2(male); 15–17 years: 49–154 (female), 68–339 (male); 17–19 years: 45–192 (female), 58–213 (male); > 19 years: 33–98 (female), 43–115 (male)
^c Serum 1,25(OH)2D3 levels were measured in external laboratories; laboratory reference ranges for each individual is as follows; 30–50 pg/mL for Pt#1.1, 16–65 pg/mL for Pt#2.2, Pt#3.1 and Pt#9.1; 15–49 pg/mL for Pt#4.1 and Pt#6.1; 19.6–65 pg/mL for Pt#7.1 and Pt#7.2
^d Diagnosed previously with hypophosphatemic rickets at the age of 17 months at another clinic therefore were already on calcitriol and phosphorus replacement therapy. Biochemical analysis at the time of the initial diagnosis has not been found
^e Admitted to our center at $3^{4/12}$ years of age and height at admission to our center reported. Biochemical analysis of the patient at the time of the initial diagnosis at another center has been found and the clinical and biochemical data at the time of first medical attention (at 11 months of age) have been reported
^f Parents originate from close villages
^g Diagnosed previously with VDDR at the age of 5 years but lost to follow up until age of 14 years; at admission to our clinic patient was not taking any medication for one year
^h Patients 5.1 and 6.1 had not been admitted to regular endocrinology visits and were not taking any medication for one month and two years, respectively
¹ Patients 7.1 and 7.2 had been diagnosed previously with VDDR at the age of 12 and 11.6 years, respectively, but lost to follow up after 3 years until the last admission to our clinic; both patients were not taking any medication for three years

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Pt#10.1, respectively (Table 2). **Genotype-Phenotype Correlation** Combined analyses of the current cohort (n: 13) and other cases in the literature (n: 183) were performed. Three mutations, namely p.F443Pfs*24, c.195+2T>G, and p.V88Wfs* were the most common mutations in the CYP27B1 gene and identified as a homozygous state in 24, 20, and 28 patients, respectively. Comparison of available data regarding clinical features of each mutation group demonstrated that height SDS at presentation was lower and, higher calcitriol treatment doses required during treatment in patients with c.195 + 2T > G mutation, suggesting more severe disease (Table 4). Evaluation of Turkish cohorts separately demonstrated that p.K192E mutation is among the three most common mutation along with p.F443Pfs*24 and c.195+2T>G. Comparison of clinical features of p.K192E mutation with the other two mutations demonstrated that patients with p.K192E had older age and better height SDS at presentation and lower calcitriol replacement dose requirement during treatment, suggesting milder disease (Supplementary Table S1).

(p.L380Afs*57) mutations were detected in Pt#9.1 and

Discussion

Patient also had polymorphism in exon 7 (c.1215T > C; p.Asn405Asn)

In this report, we presented detailed clinical features and long-term follow-up of 13 new patients with VDDR-IA and delineated genotype-phenotype correlation for the most common mutations thorough the literature review. Based on the mutations registered in the Human Gene Mutation Database (HGMD; January 2020) and literature search, a total of 81 different mutations in CYP27B1 gene have been identified as a cause of VDDR-IA (Fig. 1) [11]. Among them, we identified three most commonly observed pathogenic variants which are a seven-nucleotide duplication in exon 8 (p.F443Pfs*24), a small deletion in exon 2 (p.V88Wfs*71), and a splice donor site mutation in intron 1 (c.195+2T>G), all together, observed in 67 [5, 7, 12–22], 63 [5, 12, 13, 20] and 42 [7, 8, 15, 16] alleles, respectively. In Turkish patients cohorts, two most frequently observed pathogenic variants in CYP27B1 are also seven-nucleotide duplication in exon 8 and a splice donor site mutation in intron 1 (c.195+2T>G) as in the rest of the world. However, in Turkish patients and in our cohort reported here, a specific mutation p.K192E appears to be one of the most common mutations.

The seven-nucleotide duplication in exon 8 has been reported in different ethnic groups [5, 13, 14, 17–22] and has been suggested that it has risen by independent de novo events, probably as the result of a slipped strand mispairing

Table 3	Long term	follow-up data o	of the patients wh	o reached adult height
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Pt#	Age (year)	Calcitriol	treatment				Final height SDS MPH SDS Lumb				
		Age of	Duration (year)	Dose (ng/kg/	day)	Compliance			score)		
		onset (year)		At diagnosis	Current						
1.1	18.9	1.16	17.74	86.2	17.2	Noncompliant (last 4 years)	- 0.9	- 1.46	- 1.45		
1.2	17.4	1	16.4	33.7	7.6	Compliant*	- 0.36	- 1.46	0.83		
4.1	27.0	1.75	25.25	58.8	7.7	Compliant*	- 2.63	- 2.46	0.4		
5.1	24.2	5	18	23.6	15.3	Compliant*	- 0.36	0.86	1.7		
6.1	21.2	1.83	17	39.3	0	No treatment (last 2 years)	0.62	0.37	- 1.9		
7.1	18.26	12	3	24.0	0	No treatment (last 3 years)	- 2.91	- 1.61	- 0.8		
7.2	17.19	11.6	3	14.0	0	No treatment (last 3 years)	- 2.63	- 1.55	- 0.3		
10.1	16.13	4.25	11.88	72.9	17.2	Noncompliant (last 5 years)	- 1.89	- 1.68	NA		

MPH mid-parental height, BMD bone mineral density

*Compliant patients were defined as those who took at least 80% of the medication prescribed

 Table 4
 Comparison of phenotypic features of the patients with most common mutations seen worldwide (available data of all patients reported to date, including the patients presented in this paper)

	Exo p.F4	n 8ª 143Pfs*24	Intro c.19	on 1^{b} 95 + 2T > G	Exon 2 ^c p.V88Wfs*71		Р
	n	Median (min; max)	n	Median (min; max)	n	Median (min; max)	
Age at first medical attention (months)	24	16.0 (4; 138)	20	19.2 (9; 36)	21	13 (1.2; 120)	0.0581
Height at first admission (SDS)	10	- 2.22 (- 5.7; - 0.25)	18	- 3.45 (- 5.5; - 2)	7	- 2.7 (- 4.3; 0.8)	0.0391 ^d
Dose of calcitriol treatment (ng/kg/day)	9	17.2 (7.62; 76.92)	12	52.8 (30; 117.6)	0	-	0.0148
Total calcium (mmol/L) (Norm: 2.1-2.7)	19	1.79 (1.27; 2.27)	20	1.84 (1.39; 2.29)	11	1.62 (1.29; 2.49)	0.4129
Phosphate (mmol/L) (Norm*)	20	0.80 (0.51; 1.77)	20	0.87 (0.58; 2.09)	12	1.16 (0.58; 2.48)	0.0874
Alkaline phosphatase (IU/L) (Norm: 46–110)	18	1982 (443; 9251)	20	1862 (496; 3153)	12	1408 (263; 3952)	0.1328
Parathormone (pg/mL) (Norm: 15-65)	18	525.5 (10.8; 1320)	20	617.5 (148.1; 1423)	7	660 (101; 2231)	0.6479
1.25(OH) ₂ D (pg/mL) (Norm: 16–65)	14	5.25 (0; 85.9)	11	12 (0; 38)	4	8.5 (4.9; 55)	0.7282
25(OH)D (ng/mL) (Norm: 20–100)	17	62 (21.1; 304)	20	47.9 (14.2; 238)	8	45.9 (30; 256)	0.7324

Bold P values indicate the statistical significans difference between the groups

P values were calculated using the Kruskal-Wallis and Mann-Whitney test

*Age-specific normal ranges; 0–5 months: 1.7–2.7; 6–12 months: 1.6–2.5; 1–5 years: 1.4–2.0; 6–12 years: 1.2–1.9; > 13 years: 0.7–1.4 mmol/L

^aReferences #: 5. 7, 12-17, 19-21, this study

^bReferences #: 7, 8, 15, 16

^cReferences #: 5, 12, 13, 20

^dThere is no statistically significant difference between exon 2 (p.Val88Trpfs*71) and exon 8 (p.F443Pfs*24) mutation. There is no statistically significant difference between exon 2 (p.Val88Trpfs*71) and intron 1 (c.195+2T>G) mutation

during meiosis [13]. It has also been reported in Turkish cohorts previously [7, 15, 16], which along with our study further confirmed that the seven-nucleotide duplication in exon 8 is a common mutation distributed worldwide.

Splice donor site mutation in intron 1 (c.195 + 2T > G) observed in 42 alleles (20 homozygous, 2 compound heterozygous), which makes it the third most common mutation in *CYP27B1* described in the literature and only found in Turkish population [7, 8, 15, 16]. Since this mutation has not been reported in other ethnic groups, it may delineate a

founder or common ancestor effect [16]. To date, 63 Turkish patients with VDDR-IA (including our patients) have been studied for *CYP27B1* mutations and 11 different mutations have been described. Nine of those described mutations including the common intron 1 (c.195+2T>G) are unique to the Turkish population (Supplementary Table S2) [7, 8, 15, 16, 23].

The second most common mutation worldwide is deletion in exon 2 (p.V88Wfs*71) is a known frameshift mutation that results in a severely truncated protein. It is a founder mutation mostly observed in French Canadian patients, originating from Quebec, where the prevalence of VDDR-IA is reported to be unusually high [12, 20]. This mutation has not been described yet in Turkish patients whereas p.K192 E mutation appears to be the third most common mutation in Turkish patients. p.K192E mutation has not been reported in other ethnic groups, as well (Supplementary Table S2).

Previously, some authors suggested that the severity of the CYP27B1 mutation cannot be predicted from the clinical presentation of the disease [17, 24]. However, analyzing the clinical and molecular characteristics of the patients reported to date, we were able to demonstrate a certain genotype-phenotype correlation in VDDR-IA patients. Although we could not find significant differences in biochemical parameters at presentation between the different genotypes, the age and height SDS at presentation and calcitriol doses required to control the disease differed among the genotypes. It appears that intron 1 c.195 + 2T > G mutation causes a more severe phenotype with lower height SDS at presentation and, higher calcitriol requirement, while a less severe phenotype occurs in patients with p.K192E mutation. The previously studied functional assay of the mutations in CYP27B1 showed that almost all mutations cause either complete loss of enzymatic activity [12, 24] or retained just the 5-12% of normal enzymatic activity [18, 25]. Nevertheless, only a few disease causing mutations showed enzymatic activity of more than 20% of normal [25, 26], in which clinically asymptomatic homozygous mutation carriers in older ages had been described with elevated ALP and PTH levels as the biochemical features of rickets [26]. Although we could not perform a functional assay, we predict that p.K192E mutation leads to less severe loss of enzymatic activity, since the patients with this mutation present in older age and with milder biochemical abnormalities compared to other mutations (Supplementary Table S1).

There are limited data about the long-term outcome of the disease [5, 17]. Edouard et al. [5] observed that treatment with calcitriol resulted in the normalization of biochemical parameters within three months while other studies reported a longer period of time (6–12 months) [16, 17]. The difference in the time for normalization of biochemical parameters could be due to the difference in the initial calcitriol dosage. Edouard et al. [5] started calcitriol treatment at a dose of 1.0 μ g/day and decreased thereafter, while, similar to us, Chi et al. [17] started calcitriol treatment at a median daily dose of 0.5 µg/day (range 0.25–1.5 µg). In our cohort, normocalcemia was achieved at 1-3 months of treatment, but normalization of PTH and ALP occurred later between 3 and 12 months. Thus, our study further supports the relationship between the initial dose of calcitriol therapy and time for normalization of biochemical parameters. Nephrocalcinosis, a complication of treatment, was not detected in any of the patients during follow-up in our cohort or the patients of Chi et al. [17], while 16% of the cases with higher initial calcitriol treatment doses were found to have nephrocalcinosis [5]. Thus, treatment initiation with lower doses and titrating the dose according to biochemical response could be a more appropriate approach to avoid the treatment-related complications.

It has been reported that patients had normal BMD after three months of treatment, even though they received calcitriol only after puberty, suggested that having low levels of $1,25(OH)_2D_3$ during childhood does not affect the achievement of normal peak bone mass [5]. However, our findings dispute that observation. We have demonstrated in our older patients that bad treatment compliance during childhood before the growth cease leads to short adult stature and low BMD in the long term (Table 3). Thus, we suggest that early initiation of treatment and good compliance are essential in achieving normal height and BMD.

A few clinical points in VDDR-IA deserve further discussion. Clinical and laboratory features of VDRR-IA are similar to nutritional rickets (NR) which may cause initial misdiagnosis and treatment delay as in our patients. Normal/high 25(OH)D3 levels, low or inappropriately normal 1,25(OH)₂D3 levels, and unresponsiveness to cholecalciferol treatment are important features of VDDR-IA that differ from NR. Misdiagnosis of HR as in our Pt#3.1, and phosphate treatment may also delay healing of rickets and lead to additional complications such as nephrocalcinosis/ nephrolithiasis [27], since the phosphate treatment further increases phosphate excretion due to stimulatory effect on PTH. Discontinuation of phosphate treatment improved microlithiasis in our case. Mild nephrocalcinosis has been detected in 16% of VDRR-IA [5] whereas about 30-70% of patients with HR have been reported to develop nephrocalcinosis [28]. Another patient (Pt#8.1) in our cohort had hyperphosphatemia and for that reason, initially evaluated for PHP. When we evaluated all reported cases in the literature, although they are not discussed in the relevant papers, there are several normo/hyperphosphotemic cases with proven *CYP27B1* mutation, similar to our Pt#8.1 [5, 12, 13, 25, 26, 29–31]. Falsely elevated serum phosphate concentrations (pseudohyperphosphatemia) have been described in several conditions such as paraproteinemia, hemolysis, and icteric and lipemic serums [32]. However, we could not detect any of these conditions in our patient, and she continued to have upper normal phosphate levels (around 1,6 mmol/L) during follow-up. Although the exact mechanism of normo/hyperphosphatemia in some cases with CYP27B1 mutation is yet unknown, it is important to underline that normo/hyperphosphatemia can be seen rarely in genetically proven VDRRR-1A patients [5, 12, 13, 25, 26, 29-31].

Lastly, in VDRR-IA, low $1,25(OH)_2D3$ levels due to deficient 1 α -hydroxylase enzyme activity is the hallmark of the disease. However, several cases reported in the literature

having normal 1,25(OH)₂D3 levels [6, 14-16, 25]. In our study, Pt#3.1 had mildly elevated 1,25(OH)₂D3 level as well. A small increase of 1,25(OH)₂D3 level has been observed in CYP27B1 knockout mice after given high dietary vitamin D and, normalization of 1,25(OH)₂D3 level after vitamin D replacement in a case with VDDR-IA has been also reported [33]. However, we could not detect any correlation between prior cholecalciferol intake (r: 0.487; p: 0.40) or 25(OH)D3 levels (r: -0.154; p:0.83) and 1,25(OH)₂D3 levels at admission in our cohort. Furthermore, it seems that variation in 1,25(OH)₂D3 levels could not be explained with genotype, since, in our cohort, the patients with the same mutation (p.F443Pfs*24) having low to high 1,25(OH)₂D3 levels independently from serum 25(OH)D3 levels. These findings suggest that 25(OH)D3 could be converted to 1,25(OH)₂D3 by a non-*CYP27B1* enzyme [34] or other enzymatic pathways involving in vitamin D metabolism or degradation might have a role in 1,25(OH)₂D3 levels. Recently, novel CYP11A1-derived hydroxy derivatives of D3 have been described and, 20(OH)D3 is the main metabolite of this reaction, and acts as biased agonists on VDR [35, 36]. It is also possible that 20(OH)D3 might cross reacts with 1,25(OH)₂D3 in the assays. It should also be considered that reference ranges for 1,25(OH)₂D3 levels are determined in normal subjects with normal PTH and phosphate levels. In situations with elevated PTH and low phosphate levels, due to their stimulatory effect on 1α -hydroxylase, 1,25(OH)₂D3 levels should be much higher. Thus, normal or mildly elevated 1,25(OH)₂D3 levels in VDDR-IA can be considered as inappropriately normal in that setting. Thus, clinical and biochemical findings of cholecalciferol treatment-resistant calciopenic rickets with normal or increased 25(OH)D3 levels, together with good treatment response to the standard-dose calcitriol treatment, will exclude NR, VDDR-IB,-II, and-III and make the most probable diagnosis of VDRR-IA, even without measurement of 1,25(OH)₂D3 levels. We made the clinical diagnosis of VDDR-IA in all of our patients and Pt#3.1 as well with these principles and eventually confirmed with genetic testing. Genetic analysis is required in cases with unusual features to identify the exact type of rickets and/or hypocalcemia as in our case with hyperphosphatemia and elevated 1,25(OH)₂D3 level.

In conclusion, early and correct diagnosis of rare genetic forms of rickets is challenging, especially in developing countries where NR is the most common form of the disease. VDDR-IA should be considered when a patient develops signs of rickets while under vitamin D prophylaxis and has a normal 25(OH)D3 level or is unresponsive to cholecalciferol treatment for NR. Genetic analysis is important for making an accurate diagnosis since there are rare cases with VDDR-IA who have normo/hyperphosphatemia or normal 1,25(OH)₂ D3 levels. Proper and timely treatment is important for better long-term outcomes as we have demonstrated poor final height and osteopenia in noncompliant patients.

Finally, our cohort together with the reviewed cases in the literature suggests a genotype–phenotype correlation in patients with VDDR-IA notably that intron 1 c.195+2T > Gmutation causes more severe phenotype with lower height SDS at presentation and, higher calcitriol requirement, while less severe phenotype occurs in patients with p.K192E mutation.

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Compliance with Ethical Standards

Conflicts of interest Sare Betul Kaygusuz, Ceren Alavanda, Tarik Kirkgoz, Mehmet Eltan, Zehra Yavas Abali, Didem Helvacioglu, Tulay Guran, Pinar Ata, Abdullah Bereket, and Serap Turan declare that they have no conflicts of interest.

Ethical Approval All procedures performed in studies involving human participants were in accordance with the ethical standards of the Marmara University School of Medicine Ethical Committee and with the 1964 Helsinki declaration and its later amendments.

Consent to Participate Informed consent was obtained from parents of all patients for genetic testing.

Consent for Publication Informed consent was obtained from parents of all patients for publication of data.

References

- 1. Chiellini G, DeLuca F (2011) The importance of stereochemistry on the actions of vitamin D. Curr Top Med Chem 11:840–859
- Acar S, Demir K, Shi Y (2017) Genetic causes of rickets. J Clin Res Pediatr Endocrinol 9:88–105. https://doi.org/10.4274/jcrpe .2017.S008
- Roizen JD, Li D, O'Lear L et al (2018) CYP3A4 mutation causes vitamin D-dependent rickets type 3. J Clin Invest 128:1913–1918
- Fraser D, Kooh SW, Kind HP et al (1973) Pathogenesis of hereditary vitamin-D-dependent rickets: an inborn error of vitamin D metabolism involving defective conversion of 25-hydroxyvitamin D to 1α, 25-dihydroxyvitamin D. N Engl J Med 289:817–822
- Edouard T, Alos N, Chabot G et al (2011) Short- and long-term outcome of patients with pseudo-vitamin D deficiency rickets treated with calcitriol. J Clin Endocrinol Metab 96:82–89. https ://doi.org/10.1210/jc.2010-1340
- Giannakopoulos A, Efthymiadou A, Chrysis D (2017) A case of vitamin-D-dependent rickets type 1A with normal 1,25-dihydroxyvitamin D caused by two novel mutations of the CYP27B1 gene. Horm Res Paediatr 87:58–63. https://doi.org/10.1159/000446774
- 7. Tahir S, Demirbilek H, Ozbek MN et al (2016) Genotype and phenotype characteristics in 22 patients with vitamin

D-dependent rickets type I. Horm Res Paediatr 85:309–317. https://doi.org/10.1159/000444483

- 8. Demir K, Kattan WE, Zou M et al (2015) Novel CYP27B1 gene mutations in patients with vitamin D-dependent rickets type 1A. PLoS ONE 10:1–14. https://doi.org/10.1371/journ al.pone.0131376
- Baykan O, Yaman A, Arpa M et al (2014) The comparison of serum vitamin D3 measurement with HPLC, HPLC coupled tandem mass spectrometry using atmospheric pressure chemical ionization, and immunoassay methods. Clin Chem Lab Med 52:3. https://doi.org/10.1515/cclm-2014-4042
- Untergasser A, Cutcutache I, Koressaar T et al (2012) Primer3—new capabilities and interfaces. Nucleic Acids Res 40:e115-e115
- Stenson PD, Ball EV, Mort M et al (2003) Human gene mutation database (HGMD®): 2003 update. Hum Mutat 21:577–581
- Wang JT, Lin CJ, Burridge SM et al (1998) Genetics of vitamin D 1α-hydroxylase deficiency in 17 families. Am J Hum Genet 63:1694–1702. https://doi.org/10.1086/302156
- Chan JK, Kaplan LE, Perwad F et al (2007) Vitamin D 1α-hydroxylase gene mutations in patients with 1α-hydroxylase deficiency. J Clin Endocrinol Metab 92:3177–3182. https://doi. org/10.1210/jc.2006-2664
- Ito N, Peña AS, Perano S et al (2014) First Australian report of vitamin D-dependent rickets type I. Med J Aust 201:420–421. https://doi.org/10.5694/mja13.00220
- Durmaz E, Zou M, Al-Rijjal RA et al (2012) Clinical and genetic analysis of patients with vitamin D-dependent rickets type 1A. Clin Endocrinol (Oxf) 77:363–369. https://doi.org/10.111 1/j.1365-2265.2012.04394.x
- Fatma D, Gamze O, Heves K et al (2019) Genetic and clinical characteristics of patients with vitamin D dependent rickets type 1A. J Clin Res Pediatr Endocrinol 11:34–40. https://doi. org/10.4274/jcrpe.galenos.2018.2018.0121
- Chi Y, Sun J, Pang L et al (2018) Mutation update and longterm outcome after treatment with active vitamin D3 in Chinese patients with pseudovitamin D-deficiency rickets (PDDR). Osteoporos Int 30:481–489. https://doi.org/10.1007/s00198-018-4607-5
- Cui N, Xia W, Su H et al (2012) Novel mutations of CYP27B1 gene lead to reduced activity of 1α-hydroxylase in Chinese patients. Bone 51:563–569. https://doi.org/10.1016/j. bone.2012.05.006
- Smith SJ, Rucka AK, Berry JL et al (1999) Three families with pseudovitamin D-deficiency rickets resulting in loss of functional enzyme activity. J bone Miner Res 14:730–739
- Yoshida T, Monkawa T, Tenenhouse HS et al (1998) Two novel 1α-hydroxylase mutations in French-Canadians with vitamin D dependency rickets type I. Kidney Int 54:1437–1443. https://doi. org/10.1046/j.1523-1755.1998.00133.x
- Beck-Nielsen SS, Hertel NT, Brock-Jacobsen B (2006) Vitamin D 1 alpha-hydroxylase deficiency as the cause of severe rickets in a 1-year-old-old boy. Ugeskr Laeger 168:700–702
- Porcu L, Meloni A, Casula L et al (2002) A novel splicing defect (IVS6+ 1G> T) in a patient with pseudovitamin D deficiency rickets. J Endocrinol Invest 25:557–560

- 23. Orbak Z (2017) A novel mutation of CYP27B1 in two siblings with vitamin D-dependent rickets type 1A. In: 8th International Conference on Children
- 24. Kitanaka S, Murayama A, MToshioyuki S et al (1999) No enzyme activity of 25-hydroxyvitamin D3 1-a hydroxylase gene product in pseudovitamin D deficiency rickets, including that with mild clinical manifestation. J Clin Endocrinol Metab 84:4111–4117
- 25. Wang X, Zhang MYH, Miller WL, Portale AA (2002) Novel gene mutations in patients with 1α-hydroxylase deficiency that confer partial enzyme activity in vitro. J Clin Endocrinol Metab 87:2424–2430. https://doi.org/10.1210/jc.87.6.2424
- Alzahrani AS, Zou M, Baitei EY et al (2010) Alzahrani2010 a novel G102E mutation of CYP27B1 in a large family with vitamin D-dependent rickets type 1.pdf. J Clin Endocrinol Metab 95:4176–4183
- Füchtbauer L, Brusgaard K, Ledaal P et al (2015) Case report: vitamin D-dependent rickets type 1 caused by a novel CYP27B1 mutation. Clin Case Reports 3:1012–1016. https://doi. org/10.1002/ccr3.406
- Haffner D, Emma F, Eastwood DM et al (2019) Clinical practice recommendations for the diagnosis and management of X-linked hypophosphataemia. Nat Rev Nephrol 15:435–455. https://doi. org/10.1038/s41581-019-0152-5
- Hu WW, Ke YH, He JW et al (2014) A novel compound mutation of CYP27B1 in a Chinese family with vitamin D-dependent rickets type 1A. J Pediatr Endocrinol Metab 27:335–341. https:// doi.org/10.1515/jpem-2013-0183
- Kim CJ (2011) Vitamin D dependent rickets type I. Korean J Pediatr 54:51–54. https://doi.org/10.3345/kjp.2011.54.2.51
- Kitanaka S, Takeyama K, Murayama A et al (1998) Inactivating mutations in the 25-hydroxyvitamin D3 1α-hydroxylase gene in patients with pseudovitamin D-deficiency rickets. N Engl J Med 338:653–662
- Larner AJ (1995) Pseudohyperphosphatemia. Clin Biochem 28:391–393
- Koek WNH, Zillikens MC, van der Eerden BCJ, van Leeuwen JPTM (2016) Novel compound heterozygous mutations in the CYP27B1 gene lead to pseudovitamin D-deficient rickets. Calcif Tissue Int 99:326–331. https://doi.org/10.1007/s0022 3-016-0165-z
- Nishikawa M, Yasuda K, Takamatsu M et al (2019) Generation of 1, 25-dihydroxyvitamin D3 in Cyp27b1 knockout mice by treatment with 25-hydroxyvitamin D3 rescued their rachitic phenotypes. J Steroid Biochem Mol Biol 185:71–79
- Slominski AT, Manna PR, Tuckey RC (2015) On the role of skin in the regulation of local and systemic steroidogenic activities. Steroids. https://doi.org/10.1016/j.steroids.2015.04.006.On
- Wang J, Slominski A, Tuckey RC, Janjetovic Z (2012) 20-Hydroxyvitamin D3 inhibits proliferation of cancer cells with high efficacy while being non-toxic. Anticancer Res 32:739–746

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