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1 **Vitamin D Receptor FokI polymorphism is a determinant of both maternal and**  
2 **neonatal Vitamin D concentrations at birth**

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38

39 **Abstract**

40 Maternal vitamin D deficiency is considered to be the key determinant of the  
41 development of neonatal vitamin D deficiency at birth and during early infancy.  
42 Specific vitamin D receptor (*VDR*) gene polymorphisms have been associated with  
43 adverse pregnancy and offspring outcomes. The aim of this study was to evaluate the  
44 effect of maternal and neonatal *VDR* polymorphisms (ApaI, TaqI, BsmI, FokI, Tru9I)  
45 on maternal and neonatal vitamin D status. *VDR* polymorphisms were genotyped in  
46 70 mother-neonate pairs of Greek origin, and classified according to international  
47 thresholds for Vitamin D status. Mean neonatal and maternal 25-hydroxy-vitamin D  
48 [25(OH)D] concentrations were  $35 \pm 20$  and  $47 \pm 26$  nmol/l, respectively. Neonatal  
49 *VDR* polymorphisms were not associated with neonatal 25(OH)D concentrations. In  
50 contrast, mothers with the FokI FF polymorphism had a 70% lower risk of vitamin D  
51 deficiency [25(OH)D <30 nmol/l] compared with ff ones, after adjustment for several  
52 confounders. They were also in 73% and 88% lower risk of giving birth to vitamin D  
53 deficient [25(OH)D <30 nmol/l] neonates compared with Ff and ff mothers,  
54 respectively. These results suggest a protective role of maternal FokI FF genotype  
55 against both maternal and neonatal vitamin D deficiency. Further studies are needed  
56 to clarify the complex gene-gene and gene-environment interactions that determine  
57 vitamin D status at birth.

58

59 **Keywords:** Vitamin D; pregnancy; neonatal health; calcium; rickets.

60

## 61 1. Introduction

62 The teleological purpose of an ongoing pregnancy is to fulfil its fundamental role in a  
63 successful, uncomplicated delivery, along with an optimal intrauterine environment  
64 for the developing fetus [1]. Vitamin D homeostasis during pregnancy is adapted to  
65 meet both those demands, first, by stimulation of calcium (Ca) absorption for  
66 adequate intrauterine bone mineral accrual of the fetus and second, by enhancing  
67 systemic and local maternal tolerance to paternal and fetal alloantigens [1]. On that  
68 basis, data from observational studies during the last decade have suggested a  
69 potential adverse effect of maternal hypovitaminosis D during pregnancy on maternal  
70 and offspring health outcomes. Randomized trials of moderate quality indicate that  
71 vitamin D supplementation during pregnancy might reduce the risk of pre-eclampsia,  
72 gestational diabetes, low birth weight and severe postpartum hemorrhage [1].  
73 Maternal vitamin D deficiency is also considered to be the key determinant of the  
74 development of neonatal vitamin D deficiency at birth and during early infancy.  
75 Maternal 25-hydroxy-vitamin D [25(OH)D] crosses the placental barrier and  
76 represents the main pool of vitamin D for the fetus [2]. Serum fetal (cord blood)  
77 25(OH)D concentrations correlate strongly with maternal 25(OH)D concentrations,  
78 being on average 25% lower compared with the latter [3,4].  
79 Guidelines suggest a maternal vitamin D intake of >600 IU/day, to prevent elevated  
80 cord blood alkaline phosphatase, increased fontanelle size, neonatal hypocalcemia and  
81 congenital rickets [2] and to ensure the adequacy of maternal vitamin D status,  
82 especially in women at risk of deficiency [5]. However, there is an ongoing  
83 controversy among experts worldwide about the definition of maternal vitamin D  
84 deficiency during pregnancy, especially about the optimal thresholds of maternal  
85 25(OH)D concentrations ( $\geq 50$  nmol/l vs.  $\geq 75$  nmol/l) [6,7]. On the other hand,

86 different criteria are used to define optimal neonatal vitamin D status (sufficiency >50  
87 nmol/l, insufficiency 30-50 nmol/l, deficiency <30 nmol/l) [5].

88 Well-designed clinical trials indicate that maternal 25(OH)D concentrations >100  
89 nmol/l (40 ng/ml) during pregnancy are associated with a 60 % reduction in preterm  
90 birth risk in the daily clinical obstetric care [8]. In this setting,  
91 prenatal screening programs for optimizing 25(OH)D concentrations have been  
92 demonstrated as an effective approach to detect deficient women and prevent  
93 pregnancy complications [9]. Previously published randomized controlled trials  
94 demonstrated that a daily vitamin D dose of 4000 IU safely elevated circulating  
95 25(OH)D concentrations and normalized vitamin D metabolism and Ca homeostasis  
96 in pregnant women, regardless of race [10,11]. More specifically, a circulating  
97 25(OH)D level of about 100 nmol/l was found to be the required concentration to  
98 optimize production of 1,25(OH)<sub>2</sub>D during pregnancy through renal and/or placental  
99 production of the hormone [12].

100 Based on these findings, a target of maternal 25(OH)D >100 nmol/l seems to be  
101 biologically and scientifically sound [13]. The fact that the aforementioned data was  
102 not incorporated into previous systematic reviews [1], might lie in the fact that most  
103 of those systematic analyses did not include trials in which any amount of the  
104 investigated agent was given to the control group, including the study by Hollis et al.  
105 [12], in which the authors considered unethical not to supplement with minimal dose  
106 the control group.

107 Despite the differences in definitions of maternal and neonatal vitamin D status and  
108 the lack of uniform results on the association between maternal thresholds and  
109 neonatal outcomes, the appliance of criteria for vitamin D status resulted in an  
110 improvement of the management of maternal hypovitaminosis D in the daily clinical

111 setting [5-7]. On the other hand, the clinical aspects of this controversy are largely  
112 reflected in the conflicting results between observational and supplementation studies  
113 [14], also affected by country-specific dietary patterns, public health policies and  
114 variation in ultraviolet B (UVB) exposure, due to cultural and life-style reasons [15].  
115 In this context, the effects of genetic variations of vitamin D receptor (*VDR*) gene on  
116 maternal and neonatal vitamin D status are gaining increasing interest. Specific *VDR*  
117 polymorphisms have been associated with adverse pregnancy and offspring outcomes  
118 [16-18]. However, robust evidence of such an association is currently unavailable,  
119 given that various studies present significant heterogeneity in terms of maternal and  
120 neonatal criteria for vitamin D status, study design, sample size and racial descent of  
121 the included subjects.

122 The aim of this study was to evaluate the effect of maternal and neonatal *VDR*  
123 polymorphisms (ApaI, TaqI, BsmI, FokI, Tru9I) on maternal and neonatal vitamin D  
124 status, by applying internationally-adopted criteria for maternal and neonatal vitamin  
125 D deficiency.

126

## 127 **2. Methods**

### 128 **2.1. Inclusion and exclusion criteria**

129 This study included data and samples from a cohort of mother-child pairs at birth that  
130 has been previously described [3]. Pregnant women on regular follow-up were  
131 recruited from the Maternity Unit of the 1<sup>st</sup> Department of Obstetrics and Gynecology,  
132 Aristotle University, Thessaloniki, Greece. The inclusion criterion was full-term  
133 pregnancy (gestational week 37-42). Maternal exclusion criteria were primary  
134 hyperparathyroidism, secondary osteoporosis, heavy alcohol use ( $\geq 7$  alcohol units per  
135 week or  $\geq 6$  units at any time during pregnancy), hyperthyroidism, nephritic syndrome,

136 inflammatory bowel disease, rheumatoid arthritis, osteomalacia, obesity [body mass  
137 index (BMI) >30 kg/m<sup>2</sup>], gestational diabetes and use of medications affecting Ca or  
138 vitamin D status (e.g. corticosteroids), except for Ca and vitamin D supplements.  
139 Neonatal exclusion criteria were being small-for-gestational age (SGA) and presence  
140 of severe congenital anomalies. Informed consent was obtained from all mothers. **The**  
141 **study was conducted from January 2018 to September 2018.** The protocol received  
142 approval from the Bioethics Committee of the Aristotle University of Thessaloniki,  
143 Greece (approval number 1/19-12-2011).

144

## 145 **2.2. Demographics and dietary assessment**

146 At enrolment, maternal demographic and social characteristics, as well as dietary  
147 habits, were recorded. Ca and vitamin D dietary intake during the last month of  
148 pregnancy were assessed through a validated, semi-quantitative, food frequency  
149 questionnaire that includes 150 foods and beverages [19-21]. For each dietary item,  
150 participants were asked to report their frequency of dairy products consumption and  
151 portion size. From these data, calculations were made for estimations of consumed  
152 quantities (in g per day) based on a food composition database, modified to  
153 accommodate the particularities of the Greek diet [22] for estimating daily dietary  
154 calcium and vitamin D intake. Maternal education was classified as elementary  
155 (primary), standard (secondary) and higher (tertiary and holding of academic  
156 degrees). Maternal alcohol use during pregnancy was treated as a dichotomous  
157 variable, defined either as none (subdivided in never drinking alcohol or drinking  
158 alcohol but not during pregnancy) or light (1-2 units per week or at any one time  
159 during pregnancy) / moderate (3-6 units per week or at any one time during  
160 pregnancy) [23].



161

162 **2.3. Biochemical and hormonal assays**

163 Blood samples were obtained from mothers by antecubital venipuncture 30-60 min  
164 before delivery. Umbilical cord blood was collected immediately after clamping, from  
165 the umbilical vein. Serum and umbilical cord specimens were stored at -20°C prior to  
166 analysis for the following parameters: Ca, phosphorus (P), parathyroid hormone  
167 (PTH), 25-hydroxyvitamin D<sub>2</sub> [25(OH)D<sub>2</sub>] and 25(OH)D. Serum Ca and P  
168 determinations were performed using the Cobas INTEGRA clinical chemistry system  
169 (D-68298; Roche Diagnostics, Mannheim, Germany). The inter- and intra-assay  
170 coefficients of variation (CVs) were 1.0% and 3.5% for Ca, and 1.3% and 2.5% for P,  
171 respectively. PTH determinations were performed using the electro-  
172 chemiluminescence immunoassay ECLIA (Roche Diagnostics GmbH, Mannheim,  
173 Germany). Reference range for PTH was 15-65 pg/ml, functional sensitivity 6.0  
174 pg/ml, within-run precision 0.6-2.8% and total precision 1.6-3.4%. Concentrations of  
175 25(OH)D<sub>2</sub> and 25(OH)D were determined using novel assay, liquid chromatography-  
176 tandem mass spectrometry (LC-MS/MS), with lower limits of quantification (LLOQ):  
177 25(OH)D<sub>2</sub>(0.5 ng/ml), 25(OH)D (0.5 ng/ml). Briefly, the assay involves analyte  
178 purification using liquid-liquid extraction followed by chromatographical separation  
179 using a chiral column in tandem with a rapid resolution microbore column. Full  
180 method validation parameters have been previously reported [24,25].

181

182 **2.4. Neonatal and maternal vitamin D status**

183 Differences in the frequency of neonatal *VDR* polymorphisms were determined  
184 between three groups of neonates, according to their vitamin D status at birth:  
185 25(OH)D <30 nmol/l (deficiency), 30 ≤ 25(OH)D ≤ 50 nmol/l (insufficiency) and

186 25(OH)D >50 nmol/l (sufficiency) [2]. Differences in the genotype distribution of  
187 maternal *VDR* polymorphisms were evaluated between different groups of Vitamin D  
188 status, defined by using two thresholds for maternal 25(OH)D concentrations:  
189 [25(OH)D <30 nmol/l] vs. [25(OH)D ≥30 nmol/l] and [25(OH)D <50 nmol/l] vs.  
190 [25(OH)D ≥50 nmol/l] [26].

191

## 192 **2.5. *VDR* analysis**

193 DNA was isolated from peripheral blood samples by QIAamp DNA Blood Mini Kit  
194 (Cat. No. 51304, QIAGEN, **Hilden, Germany**) according to manufacturer's protocol.

195 In order to determine the genotypes of rs7975232 (ApaI), rs7731236 (TaqI), rs757343  
196 (Tru9I) and rs1544410 (BsmI) SNPs within *VDR* gene, Polymerase Chain Reaction  
197 (PCR) and Restriction Fragment Length Polymorphism (RFLP) methods were  
198 performed as previously described [27]. Real-Time PCR (RT-PCR) method was used  
199 for determining genotypes of rs2228570 (FokI) SNP by using Simple Probe  
200 (LightSNiP, TibMolBiol, **Berlin, Germany**) and LightCycler Fast Start DNA Master  
201 HybProbe Kit (Cat. No. 12239272001, Roche Diagnostics, **Mannheim, Germany**)  
202 with LightCycler 480 Instrument II (Roche Diagnostics, **Mannheim, Germany**).

203 Melting curve analysis were performed for genotyping as previously described [28].

204 Each SNP allele named after as follows: for rs7731236 (TaqI), "t" represents C, "T"  
205 represents T nucleotide; for rs7975232 (ApaI), "a" represents C, "A" represents A  
206 nucleotide, for rs757343 (Tru9I), "u" represents A, "U" represents G nucleotide, for  
207 rs1544410 (BsmI), "b" represents G, "B" represents A nucleotide, and for rs2228570  
208 (FokI), "f" represents T, "F" represents C nucleotide.

209

## 210 **2.6. Statistical analysis**

211 Kolmogorov-Smirnov and Shapiro-Wilk analyses were carried out on the data to test  
212 for normality. Group differences were tested using the chi-square test, whereas  
213 Fisher's exact test was applied, in case the expected values were less than five.  
214 Categorical data were presented as absolute numbers and frequencies (percentages).  
215 Univariate logistic regression was performed to identify independent associations of  
216 maternal and neonatal polymorphisms with both neonatal and maternal vitamin D  
217 status, after adjusting for confounders, such as age, smoking status, education level,  
218 alcohol consumption, Ca supplementation, dietary daily Ca and vitamin D intake  
219 during the third trimester, pre-pregnancy BMI and delivery BMI. The genotype  
220 frequencies were tested for the Hardy-Weinberg equilibrium using the  
221 [https://ihg.gsf.de/ihg/index\\_engl.html](https://ihg.gsf.de/ihg/index_engl.html) for cases and controls through the Pearson chi-  
222 square ( $\chi^2$ ) test. ORs and *p*-values, were all adjusted for confounders and tested for  
223 co-dominant, dominant and additive genetic models.

224

### 225 **3. Results**

226 Seventy mother-neonate pairs were included in the study. Demographic and  
227 laboratory data of mothers and neonates are presented in **Table 1**. Mean neonatal  
228 25(OH)D concentrations were  $35 \pm 20$  nmol/l. Overall, 52% (n=36) of the neonates  
229 were vitamin D deficient, 27% (n=19) insufficient and 21% (n=15) sufficient. Mean  
230 maternal age and 25(OH)D concentrations were  $33 \pm 6$  years and  $47 \pm 26$  nmol/l,  
231 respectively. Overall, 34% (n=24) of the mothers were vitamin D deficient, 30%  
232 (n=21) insufficient and 36% (n=25) sufficient. No deviations from HWE were  
233 observed.

234

#### 235 **3.1. Association between neonatal polymorphisms and neonatal Vitamin D status**

236 No differences in the genotype distribution of the *VDR* gene polymorphisms (ApaI,  
237 TaqI, BsmI, FokI, Tru9I) were detected among sufficient, insufficient and deficient  
238 neonates (**Table 2**). In the case of the FokI polymorphism, the difference in the  
239 frequency of the FF genotype between sufficient and deficient neonates approached,  
240 but did not reach, significance (67 vs. 36%, p=0.05).

241

### 242 **3.2. Association between maternal polymorphisms and maternal Vitamin D** 243 **status**

244 FokI FF genotype was more frequent among mothers with 25(OH)D concentrations  
245  $\geq 30$  nmol/l compared with those with  $< 30$  nmol/l (57 vs. 25%, p=0.02) (**Table 3**).  
246 Similar results were yielded when the threshold of 50 nmol/l was used to define  
247 groups of maternal Vitamin D status [64% in mothers with 25(OH)D  $\geq 50$  nmol/l vs.  
248 36% in those  $< 50$  nmol/l, p=0.02] (**Table 4**). No differences in the genotype  
249 distribution of the other polymorphisms (ApaI, TaqI, BsmI, FokI, Tru9I) were  
250 detected between maternal groups of Vitamin D status, irrespectively of the Vitamin  
251 D threshold applied. The probability of maternal deficiency [25(OH)D  $< 30$  nmol/l]  
252 was 70% lower in FokI FF mothers compared with Ff ones [odds ratio (OR) 0.3, 95%  
253 confidence interval (CI) 0.09-0.92, p=0.03] and 88% lower in carriers of the FF  
254 genotype than those of the ff genotype (OR 0.12, 95% CI 0.02-0.78, p=0.03).

255

### 256 **3.3. Association between maternal polymorphisms and neonatal Vitamin D** 257 **status**

258 Maternal FokI FF genotype was more frequent among mothers of non-deficient  
259 [25(OH)D  $\geq 30$  nmol/l] neonates compared with those of deficient ones [25(OH)D  $< 30$   
260 nmol/l] (62 vs. 31%, p<0.01) (**Table 5**). When the same analysis was performed

261 using a neonatal 25(OH)D threshold of 50 nmol/l [25(OH)D <50 nmol/l vs. 25(OH)D  
262  $\geq$ 50 nmol/l], no differences in the distribution of maternal *VDR* genotypes were  
263 observed between groups. Mothers with the FokI FF genotype presented a 73% lower  
264 risk of giving birth to vitamin D deficient neonates (logistic regression - OR 0.27,  
265 95% CI 0.1-0.77, p=0.01) compared with carriers of the Ff genotype (**Table 6**).

266

#### 267 **4. Discussion**

268 This study aimed to evaluate the effects of maternal and neonatal *VDR*  
269 polymorphisms on maternal and neonatal vitamin D status at birth, including a  
270 population from a sunny Mediterranean area in Northern Greece. Results from this  
271 maternal-neonatal pair cohort indicate that specific maternal genotypes might be  
272 protective against neonatal vitamin D deficiency, defined according to internationally  
273 applied criteria for vitamin D status [5], irrespective of neonatal *VDR* genetic  
274 variation. These findings are the first to be reported on the association between  
275 neonatal *VDR* polymorphisms and vitamin D status from this region.

276 A protective effect of maternal FokI FF genotype against the development of neonatal  
277 vitamin D deficiency [25(OH)D <30 nmol/l] was demonstrated. This effect was  
278 rationally mediated through the attainment of sufficient maternal vitamin status  
279 [25(OH)D  $\geq$ 30 nmol/l and  $\geq$ 50 nmol/l] [6,7], in mothers with the FF genotype.  
280 Neonatal 25(OH)D concentrations at birth roughly follow the maternal pattern in the  
281 deficient and insufficient mother groups, while follow the normal distribution in the  
282 group of mothers with sufficient vitamin D status [3,4]. The attainment of maternal  
283 vitamin D sufficiency during pregnancy is associated with a decreased prevalence of  
284 maternal and neonatal complications [29]. Still, there is a lack of consensus regarding  
285 specific maternal 25(OH)D thresholds that affect neonatal outcomes. Taking into

286 account that neonatal vitamin D status at birth is decreased by approximately 25%  
287 compared with the respective maternal vitamin D concentrations, a maternal  
288 25(OH)D threshold of  $\geq 50$  nmol/l, could theoretically be able to prevent the  
289 development of neonatal vitamin D deficiency. However, the extent to which this  
290 phenomenon is affected by genetic variants and ethnic differences has not been  
291 elucidated.

292 The present study was conducted in an area with a high prevalence of maternal  
293 vitamin D deficiency during pregnancy, albeit abundant sunshine [30], mainly due to  
294 sartorial habits, lack of food fortification [31] and reduced sunshine exposure,  
295 especially during the hot summer months [32]. The identification of maternal FokI FF  
296 carriers could contribute to the overall improvement of prediction scores for  
297 management of maternal and neonatal vitamin D deficiency, yet to be developed in  
298 this region. In addition, such an approach could provide individualized management  
299 of vitamin D supplementation to the future mother.

300 The FokI FF genotype was associated with optimal maternal 25(OH)D concentrations  
301 ( $\geq 30$  nmol/l and  $\geq 50$  nmol/l), still not with an increased probability of neonatal  
302 vitamin D sufficiency [25(OH)D  $\geq 50$  nmol/l]. Possible reasons for this observation  
303 might be related to the small study sample or other parameters implicated in the  
304 regulation of maternal-neonatal vitamin D equilibrium. Apart from low dietary  
305 vitamin D intake during pregnancy, additional factors such as sunlight / UVB  
306 exposure, dark skin pigmentation and maternal anthropometry may constitute gene-  
307 environment interactions that affect neonatal Vitamin D status.

308 Although an association between *VDR* polymorphisms and adverse pregnancy  
309 outcomes, such as preterm birth and SGA neonates [18,33,34] has been suggested,  
310 relative evidence is still inconclusive. In a case-control study, maternal but not

311 placental, *VDR* FokI Ff genotype was found to be lower in preeclamptic women  
312 compared with controls [35]. On the other hand, FokI *VDR* variant was associated  
313 with a higher risk for preterm birth and recurrent pregnancy loss [36,37]. The  
314 significant heterogeneity among the studies might be explained by the lack of  
315 standardized thresholds for vitamin D status, which would enable a universal  
316 stratification of mothers and neonates. The racial diversity of included populations  
317 might also contribute to the inconsistency of the results, underlying the importance of  
318 regionally-derived data in the implementation of national health policies for the  
319 prevention of vitamin D-associated adverse maternal and neonatal outcomes [38, 39].  
320 The main limitation of the present study is the relatively small sample size, which  
321 attenuates its power to reveal gene-outcome associations. Consequently, the  
322 probability that potentially significant associations may have been missed should be  
323 considered. Furthermore, neonatal vitamin D status and adverse pregnancy outcomes  
324 are dependent on complex gene-gene and gene-environment interactions, that  
325 interplay at both maternal and neonatal levels. As a result, the variation in a single  
326 gene cannot sufficiently explain the entire spectrum of the pathophysiology of vitamin  
327 D deficiency at birth. On the other hand, the strengths of the present study are the  
328 inclusion of an ethnically homogenous population of mothers and neonates and the  
329 use of multiple thresholds to determine vitamin D status. Further studies with larger  
330 sample sizes that will involve subjects of different ethnic origins are needed to  
331 replicate the findings of the present study and clarify the complex underlying  
332 mechanisms. To conclude, this study highlights the value of population-specific,  
333 genetic profiling in understanding vitamin D deficiency among neonates and their  
334 mothers.

335

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337

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339

340 **Informed consent:** Informed consent was obtained from all participants.

341

342 **Ethical approval:** The study protocol was approved by the Bioethics Committee of  
343 the Aristotle University of Thessaloniki.

344



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- 488

489 **Table 1.** Demographic and laboratory features of mothers and neonates.

<b>Variable</b>	<b>Value</b>
Mothers' age (years)	33 ± 6
BMI pre-pregnancy (kg/m <sup>2</sup> )	25 ± 5
BMI term (kg/m <sup>2</sup> )	30 ± 6
Weeks of gestation	39 ± 2
Smoking (%)	11 (15)
Alcohol consumption (%)	None: 51 (73) / Light: 12 (17) / Moderate: 7 (10)
Higher education (%)	Primary: 13 (19) / Secondary: 42 (60) / Higher: 15 (21)
Calcium supplementation (%)	20 (29)
Maternal 25(OH)D (nmol/l)	47 ± 26
Neonatal 25(OH)D (nmol/l)	35 ± 20
Calcium (mg/dl)	9 ± 1
PTH (pg/ml)	27 ± 13

490

491 Data are presented as mean ± standard deviation or absolute value (percentage).

492 Abbreviations: BMI: body mass index; PTH: parathyroid hormone; 25(OH)D: 25-

493 hydroxy-vitamin D.

494

495 **Table 2.** Genotype distribution of neonatal VDR polymorphisms according to  
 496 neonatal vitamin D status.

Polymorphism	Genotype	Deficient	Insufficient	Sufficient	p-value
		n=36 (52%)	n=19 (27%)	n=15 (21%)	
APAI	AA	12 (33)	8 (42)	3 (20)	0.57
	Aa	20 (56)	10 (53)	9 (60)	
	aa	4 (11)	1 (5)	3 (20)	
TAQI	TT	15 (42)	4 (21)	8 (53)	0.39
	Tt	16 (44)	11 (58)	5 (33)	
	tt	5 (14)	4(21)	2 (13)	
BSMI	BB	9 (25)	7 (37)	3 (20)	0.84
	Bb	14 (39)	7 (37)	6 (40)	
	bb	13 (36)	5 (26)	6 (40)	
FOKI	FF	13 (36)	10 (53)	10 (67)	0.05
	Ff	19 (53)	8 (42)	5 (33)	
	ff	4 (11)	1 (5)	0 (0)	
TRU9I	UU	24 (66)	11 (58)	11 (73)	0.70
	Uu	10 (28)	8 (42)	4 (27)	
	uu	2 (6)	0 (0)	0 (0)	

497

498 Deficient: 25(OH)D <30 nmol/l, Insufficient: 30 ≤ 25(OH)D ≤ 50 nmol/l, Sufficient:  
 499 25(OH)D >50nmol/l. Data are presented as absolute value (percentage).

500 Abbreviations: VDR: Vitamin D receptor; 25(OH)D: 25-hydroxy-vitamin D.

501



502 **Table 3.** Genotype distribution of maternal VDR polymorphisms according to  
 503 maternal vitamin D status [25(OH)D <30 nmol/l vs. 25(OH)D ≥30 nmol/l].

Polymorphism	Genotype	Maternal vitamin D status		p-value
		25(OH)D <30 nmol/l n=24 (34%)	25(OH)D ≥30 nmol/l n=46 (66%)	
APAI	AA	10 (42)	19 (43)	0.98
	Aa	11 (46)	22 (46)	
	aa	3 (12)	5 (11)	
TAQI	TT	7 (29)	18 (39)	0.60
	Tt	12 (50)	21 (46)	
	tt	5 (21)	7 (15)	
BSMI	BB	10 (42)	16 (35)	0.63
	Bb	8 (33)	13 (28)	
	Bb	6 (25)	17 (37)	
FOKI	FF	<b>6 (25)</b>	<b>26 (57)</b>	<b>0.02</b>
	Ff	14 (58)	18 (39)	
	ff	4 (17)	2 (4)	
TRU9I	UU	16 (67)	25 (54)	0.30
	Uu	8 (33)	18 (39)	
	Uu	0 (0)	3 (7)	

504

505 Data are presented as absolute value (percentage). Significant differences are  
 506 presented in bold.

507 Abbreviations: VDR: Vitamin D receptor; 25(OH)D: 25-hydroxy-vitamin D

508

509 **Table 4.** Genotype distribution of maternal VDR polymorphisms according to  
 510 maternal Vitamin D status [25(OH)D <50 nmol/l vs. 25(OH)D ≥50 nmol/l].

Polymorphism	Genotype	Maternal vitamin D status		p-value
		25(OH)D <50 nmol/l n=45 (64%)	25(OH)D ≥50 nmol/l n=25 (36%)	
APAI	AA	19 (42)	10 (40)	0.68
	Aa	22 (49)	11 (44)	
	aa	4 (9)	4 (16)	
TAQI	TT	14 (31)	11 (44)	0.60
	Tt	23 (51)	10 (40)	
	tt	8 (18)	4 (16)	
BSMI	BB	17 (38)	9 (36)	0.57
	Bb	15 (33)	6 (24)	
	bb	13 (29)	10 (40)	
FOKI	FF	<b>16 (36)</b>	<b>16 (64)</b>	<b>0.02</b>
	Ff	23 (51)	9 (36)	
	ff	6 (13)	0 (0)	
TRU9I	UU	27 (60)	14 (56)	0.91
	Uu	16 (36)	10 (40)	
	uu	2 (4)	1 (4)	

511

512 Data are presented as absolute value (percentage). Significant differences are  
 513 presented in bold.

514 Abbreviations: VDR: Vitamin D receptor; 25(OH)D: 25-hydroxy-vitamin D.

515

516 **Table 5.** Genotype distribution of maternal VDR polymorphisms according to  
 517 neonatal Vitamin D status [Vitamin D deficient: 25(OH)D <30 nmol/l, Vitamin D  
 518 non-deficient: 25(OH)D ≥30 nmol/l].

Polymorphism	Genotype	Neonatal vitamin D status		p-value
		<b>25(OH)D &lt;30 nmol/l</b> <b>n=36 (51%)</b>	<b>25(OH)D ≥30 nmol/l</b> <b>n=34 (49%)</b>	
APAI	AA	15 (42)	14 (41)	1.00
	Aa	17 (47)	16 (47)	
	aa	4 (11)	4 (12)	
TAQI	TT	11 (31)	14 (41)	0.69
	Tt	18 (50)	15 (44)	
	Tt	7 (19)	5 (15)	
BSMI	BB	14 (39)	12 (36)	0.63
	Bb	12 (33)	9 (26)	
	bb	10 (28)	13 (38)	
FOKI	FF	<b>11 (31)</b>	<b>21 (62)</b>	<0.01
	Ff	21 (58)	11(32)	
	ff	4 (11)	2 (6)	
TRU9I	UU	23 (64)	18 (53)	0.59
	Uu	12 (33)	14 (41)	
	uu	1 (3)	2 (6)	

519

520 Data are presented as absolute value (percentage). Significant differences are  
 521 presented in bold.

522 Abbreviations: VDR: Vitamin D receptor; 25(OH)D: 25-hydroxy-vitamin D.

523

524 **Table 6.** Association between maternal FokI genotypes and risk of maternal and  
525 neonatal Vitamin D deficiency [25(OH)D <30 nmol/l].

<b>Outcome</b>	<b>Maternal genotypes</b>	<b>OR</b>	<b>CI</b>	<b>p-value</b>
Neonatal deficiency	FF vs. Ff	0.27	0.10 - 0.77	0.01
Maternal deficiency	FF vs. Ff	0.30	0.09 - 0.92	0.03
Maternal deficiency	FF vs. ff	0.12	0.02 - 0.78	0.03

526

527 Abbreviations: OR: odds ratio, CI: confidence interval

528