



Weekday sunlight exposure, but not vitamin D intake, influences the association between vitamin D receptor genotype and circulating concentration 25-hydroxyvitamin D in a pan-European population: the Food4Me study

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Keywords:	25-hydroxyvitamin D, Food4Me, environment-gene interaction, sunlight, diet, vitamin D receptor gene

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Title

Weekday sunlight exposure, but not vitamin D intake, influences the association between vitamin D receptor genotype and circulating concentration 25-hydroxyvitamin D in a pan-European population: the Food4Me study

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Running title: Vitamin D status, lifestyle and genetics

Abbreviations: 25-hydroxyvitamin D (25(OH)D), Body mass index (BMI), Food frequency questionnaire (FFQ), Healthy eating index (HEI), Mediterranean diet (MD); Moderate and vigorous physical activity (MVPA); Physical activity level (PAL), Personalized Nutrition (PN), Randomized controlled trial (RCT), Sedentary behavior (SB), Waist circumference (WC)

Trial registration: Clinicaltrials.gov NCT01530139

(<http://clinicaltrials.gov/show/NCT01530139>)

Key Words: 25-hydroxyvitamin D, Food4Me, diet, vitamin D receptor gene, environment-gene interaction, sunlight

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3 1 **Abstract**
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6 2 **Scope**
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9 3 Little is known about diet- and environment-gene interactions on 25-hydroxyvitamin D
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11 4 (25(OH)D concentration. This cross-sectional study aimed to investigate i) predictors of
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13 5 25(OH)D concentration and relationships with vitamin D genotypes and ii) whether dietary
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15 6 vitamin D intake and sunlight exposure modified these relationships.
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20 7 **Methods and results**
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23 8 Participants from the Food4Me study (n=1312; age 18-79) were genotyped for vitamin D
24
25 9 receptor (*VDR*) and vitamin D binding protein at baseline and a genetic risk score was
26
27 10 calculated. Dried blood spot samples were assayed for 25(OH)D concentration and dietary
28
29 11 and lifestyle information collected. Circulating 25(OH)D concentration was lower with
30
31 12 increasing genetic risk score, lower in females than males, higher in supplement users than
32
33 13 non-users and higher in summer than winter. Carriage of the minor *VDR* allele was
34
35 14 associated with lower 25(OH)D concentration in participants with the least sunlight
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37 15 exposure. Vitamin D genotype did not influence the relationship between vitamin D intake
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39 16 and 25(OH)D concentration.
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45 17 **Conclusion**
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48 18 Age, sex, dietary vitamin D intake, country, sunlight exposure, season and vitamin D genetic
49
50 19 risk score were associated with circulating 25(OH)D concentration in a pan-European
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52 20 population. The relationship between *VDR* genotype and 25(OH)D concentration may be
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54 21 influenced by weekday sunlight exposure but not dietary vitamin D intake.
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22 INTRODUCTION

23 Vitamin D deficiency, defined by the Institute of Medicine as a circulating concentration of
24 25-hydroxyvitamin D (25(OH)D) <30 nmol/L (1), is highly prevalent (2). In addition to
25 increased incidence of skeletal health outcomes such as rickets (3) and osteoporosis (4),
26 research suggests that inadequate vitamin D intake is linked with greater risk of
27 cardiovascular disease (5), obesity (6), diabetes (7) and cancer (8).

28 Determinants of 25(OH)D concentrations include skin exposure to sunlight and dietary
29 vitamin D intake. Vitamin D synthesis occurs in the skin during exposure to ultraviolet (UVB)
30 radiation (290–320 nm), which converts 7-dehydrocholesterol to vitamin D (9). Dietary
31 sources include vitamin D₂, derived from fungi, and vitamin D₃, derived from fish and other
32 animal sources (9). Vitamin D undergoes a series of enzymatic conversions in the liver to
33 form 25(OH)D, which is the accepted blood-based biomarker for vitamin D status (10).

34 Although estimates differ, approximately 25% of the variability in 25(OH)D concentrations is
35 attributable to dietary and environmental influences (11), whereas twin and family studies
36 have shown heritability of vitamin D status to be as high as 80% (11, 12). A recent genome-
37 wide association study (13) and a systematic review (14) identified that variation in the
38 vitamin D binding protein gene, *GC*, and the vitamin D receptor gene, *VDR*, modulated
39 circulating concentrations of 25(OH)D. These findings have been replicated in a subsequent
40 cross-sectional study in 201 healthy Danish adults, where polymorphisms in *GC* and *GC* risk
41 score were associated with lower total 25(OH)D concentration (15).

42 While previous studies have evaluated the relationship between 25(OH)D concentrations,
43 vitamin D genotypes and sociodemographic characteristics (16-20), few studies have

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3 44 comprehensively evaluated interactions between genotype and dietary and sunlight
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5 45 exposure on circulating vitamin D concentrations.
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9 46 The present study uses baseline data from the Food4Me study, a pan-European randomized
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11 47 controlled trial, designed to investigate the effect of personalized nutrition (PN) advice on
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13 48 changes in diet and physical activity after a 6-month intervention. Our study had two aims
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15 49 to i) identify predictors of circulating 25(OH)D concentration in healthy European adults and
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17 50 investigate relationships with specific vitamin D genotypes and ii) to determine whether
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19 51 these relationships are modified by dietary vitamin D intake and sunlight exposure.
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54 **METHODS**

55 **Study population**

56 This analysis used baseline data from the Food4Me study (21), which was a 6-month, 4-
57 arm, internet-based, randomized controlled trial (RCT) conducted in 7 European countries
58 (22). A total of 1607 participants (age range 18 to 79) were randomized into the study and
59 recruited between August 2012 and August 2013 from the following centers: University
60 College Dublin (Ireland), Maastricht University (The Netherlands), University of Navarra
61 (Spain), Harokopio University (Greece), University of Reading (United Kingdom, UK),
62 National Food and Nutrition Institute (Poland) and Technical University of Munich
63 (Germany). The Research Ethics Committees at each University or Research Center
64 delivering the intervention granted ethical approval for the study. The Food4Me trial was
65 registered as a RCT (NCT01530139) at Clinicaltrials.gov. All participants expressing an
66 interest in the study were asked to sign online consent forms at two stages in the screening

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3 66 process. These consent forms were automatically directed to the local study investigators to
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5 67 be counter-signed and archived (22).
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10 69 **Blood sample collection**

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13 70 Participants collected finger-prick blood samples using a pack provided by Vitas Ltd, Oslo,
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16 71 Norway and DSM, Kaiseraugst, Switzerland. Pre-treated filter cards were packed in an
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18 72 airtight aluminium bag (Whatman Foil Bags, item no. 10534321, Whatman Inc., Sanford,
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20 73 ME) with a drying agent (Sorb-it, item no. 10548234, Süd-Chemie, Germany) and stored at
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22
23 74 room temperature until use. To aid collection, participants had access to an online video
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25 75 with instructions and frequently asked questions in their local language. Participants were
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27
28 76 asked to fill two dried blood spot (DBS) cards (500 μ L of blood in total), each requiring five
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30 77 spots, and were instructed to dry the cards at room temperature for at least 2 hours, but
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32
33 78 not longer than four hours, before returning by post to the corresponding recruiting center.
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35 79 Centers then shipped the DBS cards to DSM (DSM Nutritional Products Ltd. Switzerland) for
36
37 80 measurements of 25(OH)D₃ and 25(OH)D₂. No results are presented for 25(OH)D₂ due to
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39 81 all values being below the lower limit of quantification (<25 nmol/L) and detection (8
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41
42 82 nmol/L) of the analytical method and thus all results for 25(OH)D presented in the
43
44 83 manuscript refer to 25(OH)D₃. Further details of the blood spot preparation and analysis,
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46 84 including method validation and performance parameters, are provided elsewhere (23).
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51 52 86 **Genotyping**

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55 87 Participants collected buccal cell samples at baseline using Isohelix SK-1 DNA buccal swabs
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57 88 and Isohelix dried-capsules and posted samples to each recruiting center for shipment to
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89 LCG Genomics (Hertfordshire, United Kingdom). LCG Genomics extracted DNA and
90 genotyped specific loci using KASP™ genotyping assays to provide bi-allelic scoring of single
91 nucleotide polymorphisms (SNPs) of the genes encoding the vitamin D receptor (*VDR*)
92 *rs1544410* and *rs2228570* and the vitamin D binding protein (*GC*) *rs2282679*, *rs4588* and
93 *rs7041*. These 5 SNPs were selected based on findings from Genome Wide Association
94 Studies (13).

95

96 **Dietary intake, sunlight exposure and confounders**

97 Participants completed an online food frequency questionnaire (FFQ), which was developed
98 and validated for the Food4Me Study (24, 25), and included 157 food items consumed
99 frequently in each of the 7 recruitment countries. Usual intakes of total vitamin D from
100 foods and supplements at baseline were computed using a food composition database
101 based on McCance & Widdowson's "The composition of foods" (26) and a dichotomous
102 variable created to indicate if an individual was taking a vitamin D supplement.

103 Weekend and weekday sunlight exposure was estimated by asking participants if they spent
104 "Less than 20 min", "Between 20 and 45 min", "Between 45 min and 1 hour", "Between 1
105 and 2 hours" or "More than 2 hours" outside during daylight hours on a typical week day
106 and on a weekend day during the sunny months of the year (i.e. April to September) (27).

107 Responses were collapsed into "Less than 20 min", "Between 20 min and 2 hour" and "More
108 than 2 hours". Season was defined as spring (March to May), summer (June to August),
109 autumn (September to November) and winter (December to February) for the purpose of
110 descriptive statistics. Season was operationalized as a function of seasonal variation
111 estimated by $\sin(\text{SampleYear} \times 2 \times \pi)$ and $\cos(\text{SampleYear} \times 2 \times \pi)$ for inclusion as a covariate
112 in linear regression models. Full details are described elsewhere (23).

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3 113 Physical activity level (PAL, ratio between total energy expenditure and basal metabolic rate
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5 114 (BMR)), moderate and vigorous PA (MVPA) and time spent in sedentary behaviors (SB) were
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7 115 directly estimated from triaxial accelerometers (TracmorD, Philips Consumer Lifestyle, The
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9 116 Netherlands). Participants self-reported occupations, which were grouped according to the
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11 117 European commission list of occupations (28, 29). To facilitate data analysis according to
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13 118 geographic location, participant centers were grouped according to the United Nations
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15 119 Composition of geographical regions, geographical sub-regions, and selected economic and
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17 120 other groupings (30). Thus, Ireland and United Kingdom were grouped as “British Isles”,
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19 121 Germany and The Netherlands as “Western Europe”, Poland as “Eastern Europe”, and Spain
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21 122 and Greece as “Southern Europe”. Body weight (kg) and height (m) were self-measured and
22
23 123 self-reported with the aid of information sheets and online video instructions in their own
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25 124 language. Body mass index (BMI; kg/m²) was estimated from body weight and height and
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27 125 anthropometric measurements showed a high degree of reliability (31). Further details on
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29 126 these variables are provided elsewhere (22).
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128 **Statistical analyses**

129 Participants with complete data for vitamin D-related genotype, diet, vitamin D status from
130 DBS and covariates at baseline were included in the present analyses. Data were analyzed
131 using Stata (version 14; StataCorp, College Station, TX, USA). Variables were tested for
132 skewness and kurtosis, and if not normally distributed, were log transformed. Thus
133 circulating 25(OH)D concentrations, vitamin D intake from food and from food and
134 supplements, PAL and MVPA were log transformed prior to analyses. A total genetic risk
135 score was created by summing the minor alleles for *GC rs2282679 (CC)*, *GC rs4588 (AA)*, *GC*

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3 136 *rs7041 (TT)*, *VDR rs2228570 (TT)*, *VDR rs1544410 (AA)*, which were coded as 0, 1 and 2.
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5 137 Individuals were then grouped according to whether they carried 0-2, 3 to 5 or 6 or more
6
7 138 minor alleles. Additive allele coding (coded as 0, 1 and 2 for no copies, one copy and two
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9 139 copies of the minor allele respectively) was used throughout, with the exception of tests for
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11 140 interactions, where the dominant model (coded as 0 and 1 for no copies and one or two
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13 141 copies respectively) was used to increase statistical power. To address our first aim, multiple
14
15 142 linear regression tested for differences in 25(OH)D concentration (dependent variable)
16
17 143 according to sociodemographic characteristics (independent variables) and logistic and
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19 144 multiple linear regression were used to test for significant differences in baseline
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21 145 sociodemographic characteristics (dependent variable) between genotypes (independent
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23 146 variable) for categorical and continuous variables, respectively. Analyses were adjusted for
24
25 147 age, sex, BMI, ethnicity, country, season, vitamin D intake (food only) and vitamin D
26
27 148 supplementation. Models were adjusted for season by inclusion of two estimates of
28
29 149 seasonal variation as described earlier and detailed elsewhere (23). Physical activity
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31 150 outcomes were further adjusted for the length of time that the accelerometers were worn
32
33 151 (32). To investigate our second aim, interactions between vitamin D-related genotype and
34
35 152 diet or sunlight exposure on baseline 25(OH)D concentrations were tested by including an
36
37 153 interaction term in the model. To estimate any moderating effects on this interaction,
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39 154 alcohol intake (operationalized as meeting alcohol guidelines of less than 24g of alcohol/day
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41 155 or not), smoking habits (current smoker or not current smoker) and occupation
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43 156 (professional or managerial, intermediate, manual or routine, student and retired as defined
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45 157 by EU classifications of occupations (28)) were included in the model as a sensitivity analysis.
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47 158 In addition, to investigate any effect of linkage disequilibrium on the total risk score, linkage
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49 159 disequilibrium (LD) between all SNPs was calculated. If LD was observed then a revised total
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3 160 genetic risk score was derived excluding any relevant SNPs. Data were considered
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5 161 statistically significant at $P<0.05$.
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11 163 **RESULTS**

14 164 Of the 1607 participants who were randomized into the study, 1312 individuals had
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16
17 165 complete data for concentrations of 25(OH)D, *GC* and *VDR* genotype, dietary intakes and
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19 166 covariates (**Figure 1**). Mean 25(OH)D concentration was 60.6 (SD 26.4) nmol/L. A total of
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21 167 22% of participants took vitamin D supplements. Mean vitamin D3 intake from foods only
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23 168 was 159 (SD 100) IU/d and 327 (SD 1675) IU/d from foods and supplements. The mean
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25 169 proportion of individuals spending more than 2 hours in the sun on a weekday was 12% and
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27 170 40% during the weekend (**Supplemental Table 1**). The majority of individuals were tested
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29 171 during the spring months (56%), with 28% tested in winter, 9% in summer and 7% in
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31 172 autumn. No significant deviation from the Hardy-Weinberg Equilibrium was observed for
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33 173 any genotype (Table 2).
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42 175 **Associations with 25(OH) vitamin D concentration**

44 176 Concentrations of 25(OH)D were higher with increasing age ($P=0.002$), PAL ($P<0.001$), MVPA
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46 177 ($P=0.002$), total vitamin D intake ($P<0.001$) and vitamin D intake from food only ($P=0.010$)
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48 178 and lower with increasing time spent sedentary ($P=0.011$; **Table 1**). Concentrations of
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50 179 25(OH)D were also lower in females than males ($P<0.001$), highest in Western Europe
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52 180 ($P=0.001$) compared with the British Isles, and higher in supplement users than non-
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54 181 supplement users ($P<0.001$). In addition, concentrations of 25(OH)D were higher when
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3 182 measured in spring, summer and autumn compared with winter ($P<0.001$) and with
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5 183 increasing time spent in the sunlight during the weekend (P -trend=0.007) and weekday (P -
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7 184 trend=0.006).

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11 185 Concentrations of 25(OH)D were lower with each additional copy of the minor allele of *GC*
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13 186 *rs2282679*, *GC rs4588* and *GC rs7041* ($P<0.001$) but did not vary with *VDR* genotype (**Table**
14
15 187 **2**). When assessed according to a total genetic risk score, concentrations of 25(OH)D were
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17 188 lower with increasing number of minor alleles (P -trend<0.001; **Figure 2**). Baseline socio-
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19 189 demographic characteristics of participants according to *VDR* and *GC* genotype are
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21 190 presented in Supplemental Table 1 and **Supplemental Table 2**.

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26 27 28 192 **Dietary vitamin D and sunlight exposure interactions**

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31 193 The relationship between *VDR rs2228570* genotype and 25(OH)D concentration was
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33 194 modulated by time spent in the sunlight during the week (P -interaction=0.009; **Figure 3**).
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35 195 Individuals with no copies of the *VDR rs2228570* minor allele (*TT*) and reporting less than 20
36
37 196 minutes of sunlight exposure per day had 14.6 nmol/L lower concentrations of 25(OH)D
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39 197 than participants reporting more than 2 hours of sunlight exposure (P -trend=0.036).
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41 198 Individuals with one or more copies of the minor allele had 5.7 nmol/L higher
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43 199 concentrations (P -trend=0.06; **Figure 3**). When total sunlight exposure (weekdays plus
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45 200 weekend days) was considered, the interaction with *VDR rs2228570* remained significant
46
47 201 but evidence for the interaction was weaker ($P=0.045$) (See **Supplemental Figure 1**). No
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49 202 significant interactions were observed between genotype and dietary vitamin D intake from
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51 203 foods and supplements alone or combined on vitamin D status.
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3 205 **Sensitivity analyses**
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6 206 Inclusion of alcohol consumption, smoking and occupation as covariates in statistical models
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8 207 did not affect the pattern of results significantly. LD was observed between *GC rs2282679*
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10 208 and *GC rs4588* ($r^2=0.96$) and *GC rs7041* ($r^2=0.69$) and between *GC rs4588* and *GC rs7041*
11
12 209 ($r^2=0.72$). VDR SNPs showed no LD. Removal of two *GC* SNPs (*GC rs4588* and *GC rs7041*) in
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14 210 the generation of the total risk score attenuated slightly the observed relationship between
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16 211 total risk score and 25(OH)D concentration but did not change the pattern of significant
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18 212 results (P -trend=0.005).
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25 214 **DISCUSSION**
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28 215 **Main findings**
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30 216 Our main findings were that 25(OH)D concentration was associated with both modifiable
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32 217 (e.g. vitamin D intake, physical activity and sunlight exposure) and non-modifiable (e.g. age,
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34 218 sex, country and season) predictors, and that concentrations were lower in carriers of the
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36 219 minor alleles of vitamin D risk genotypes. Moreover, the relationship between time spent in
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38 220 the sunlight during the week and 25(OH)D was modulated by the *VDR rs2228570* genotype
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40 221 in a dose-wise manner. Larger studies are warranted to confirm these findings.
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48 223 **Comparison with other studies**
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50 224 For the first time in a pan-European setting, we observed that concentrations of 25(OH)D
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52 225 were higher in individuals who were more physically active, which is likely to be a proxy for
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54 226 time spent outdoors, and in participants with higher vitamin D intakes from food or
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56 227 supplements, which confirms findings from previous single country studies (13, 15, 33). Our
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3 228 findings are in agreement with the latest National Diet and Nutrition Survey (NDNS) in the
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5 229 UK, where 25(OH)D concentration increased with age in adults up to 70 years of age (34).
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7 230 However, over the age of 70 years, 25(OH)D concentrations decrease due to impaired skin
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9 231 photosynthesis and renal conversion to the active form (2). We observed that circulating
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11 232 25(OH)D concentrations were lower in females than males, which confirms previous findings
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13 233 among adults (35) and children (36). A proposed explanation for this is that women are
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15 234 more diligent in applying sunscreen and in covering skin for social reasons and that men
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17 235 spend more time outdoors during physical activity and in manual occupations.
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22 236 Our study complements findings from a genome-wide association study designed to identify
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24 237 common genetic predictors of vitamin D insufficiency (13). Participants in the highest
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26 238 quartile of genotype score (combining *GC*, *DHCR7* and *CYP2R1* variants were at increased
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28 239 risk of having 25(OH)D concentrations lower than 50 nmol/L (OR: 1.92, 95% CI: 1.70–2.16,
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30 240 $P < 0.001$) compared with participants in the lowest quartile (13). Potential mechanisms for
31
32 241 this genotype-related lower vitamin D status include impaired synthesis of 25(OH)D due to
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34 242 lower activity of the enzyme 7-dehydrocholesterol reductase by *DHCR7* variants (13) and an
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36 243 impairment of *CYP2R1* activity, which may be a microsomal enzyme responsible for 25-
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38 244 hydroxylation of vitamin D in the liver (37). In addition, *GC* encodes DBP, a protein
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40 245 synthesized in the liver that binds and transports vitamin D and its metabolites (38), and
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42 246 studies have suggested that alterations in DBP could influence proportions of free,
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44 247 circulating 25(OH)D, thereby being a rate limiting factor in the production of the vitamin D
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46 248 metabolite 1,25(OH)₂D (13, 17).
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53 249 For the first time, we observed that weekday sunlight exposure modulated the relationship
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55 250 between vitamin D-related genes and circulating vitamin D concentrations. We observed
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57 251 that greater weekday sunlight exposure mitigated the detrimental effect of carriage of the
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3 252 minor allele of *VDR rs2228570 (TT)* on 25(OH)D concentration. We anticipated that total sun
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5 253 exposure (weekday + weekend) would be the best overall measure of exposure. However,
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7 254 when considering total sun exposure, we found that the interaction with *VDR* genotype on
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9 255 circulating 25(OH)D concentration remained significant but evidence for the interaction was
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11 256 weaker for total sun exposure compared with when only weekday exposure was considered.
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14 257 Our data do not allow us to determine the reason for this unexpected finding but we
15
16 258 hypothesise that it may be related to the differences in the precision of measurement of sun
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18 259 exposure on weekdays v. weekend days. For most people, assessing time spent outdoors
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20 260 may be easier (and, therefore made with greater precision) during weekdays than at
21
22 261 weekends because of the more regular activity patterns (for both work and leisure) during
23
24 262 weekdays. Given the role of *VDR rs2228570* genotype in enhancing risk of vitamin D
25
26 263 deficiency (12, 13), the present study demonstrates the importance of sunlight exposure
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28 264 (both adventitious and recreational) in obviating the genotype-based risk. Previous research
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30 265 has shown that the vitamin D intake required to maintain circulating 25(OH)D
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32 266 concentrations in late winter above 25 nmol/L in older Irish adults was 492, 352, and 288
33
34 267 IU/d for individuals who avoided sunlight exposure, those who had some exposure, and
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36 268 those who enjoyed exposure, respectively (39). Cumulatively, this evidence emphasizes the
37
38 269 importance of considering both genotype and sunlight exposure when developing policies
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40 270 to ensure vitamin D adequacy (40). The molecular mechanism responsible for this
41
42 271 interaction is not known. As a first step, it would be helpful to confirm our findings in larger,
43
44 272 independent, studies with a wider range of sunlight exposures. If the interaction is
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46 273 confirmed, this would provide a robust foundation for future investigations of the
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48 274 underlying mechanism. Inter-personal differences in efficiency of endogenous vitamin D
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50 275 synthesis, as well as clothing and sunscreen use and inter- and intra-country variations in
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3 276 latitude, altitude, cloud cover and air pollution, mean that setting population-wide
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5 277 recommendations for sunlight exposure is difficult (41). However, PN has been highlighted
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7 278 as an important feature of current and future nutrition research (42, 43) and current
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10 279 findings should stimulate research to address these challenges within the context of PN. Our
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12 280 evidence concerning interactions between vitamin D-related genotype and vitamin D
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14 281 sources (diet and sunlight exposure) on vitamin D concentration may help in designing
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17 282 future PN interventions to improve vitamin D status in Europeans.
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21 22 284 **Strengths and limitations**

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25 285 The present study had a number of strengths. Our participants were drawn from 7 European
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27 286 countries, facilitating the comparison of vitamin D concentrations between European
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29 287 countries. Vitamin D status was determined using a relatively novel approach in which
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31
32 288 25(OH)D concentrations were measured in dried blood spots self-collected by the study
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34 289 participants. Although performance characteristics did not meet those of current gold
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36 290 standard liquid chromatography in plasma, they were found to be suitable for status-level
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38 291 determination under field conditions (23) and this approach offers a novel and likely cost-
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40 292 effective method for collecting blood samples remotely for 25(OH)D quantification.

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44 293 A limitation of our study is that data were self-measured and self-reported via the Internet,
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46 294 which may have introduced measurement error. Nonetheless, the accuracy of internet-
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48 295 based, self-reported anthropometric data is high (44), which this has been confirmed in our
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50 296 study (31). Dietary intakes were estimated by a FFQ which is subject to misreporting error
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52 297 (45) but this was minimized by prior validation against a 4-day weighed food record (25).
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55 298 Furthermore, our measure of sunlight exposure was limited to an indirect measurement
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57 299 that may be subject to self-reporting bias (27, 46), however, similar measures of indirect
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3 300 sunlight exposure have been used successfully previously (47, 48). Small sample size limited
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5 301 our power to investigate the effect of individual gene variants in the present study.
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7 302 Moreover, 97% of our study participants were Caucasians and thus further research in wider
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9 303 ethnicity groups is required to generalize our findings to other population groups. Our
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11 304 sample is a self-selected group of individuals, who may be more health-conscious than the
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13 305 general population. However, characterization of our participants suggests that they were
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15 306 similar to most European adults and would benefit from improved diet and physical activity
16
17 307 (49). For technical reasons, we were unable to evaluate the role of 25(OH)D₂, which may
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19 308 have influenced estimates of total vitamin D concentration (50), but research suggests that
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21 309 genetic variation in the vitamin D binding protein influences responsiveness to 25(OH)D only
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23 310 (51). Future studies should consider the complex interplay between lifestyle factors and
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25 311 other genetic variants, such as *rs12785878*, located near the 7-dehydrocholesterol
26
27 312 reductase *DHCR7* gene, *rs10741657*, located near the *CYP2R1* gene and variations in
28
29 313 *CYP24A1*, which encodes the kidney 24-hydroxylase enzyme (13) on vitamin D status. In
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31 314 addition, information on factors influencing individual level sun exposure including weather,
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33 315 skin type and clothing habits should be collected. Finally, although the present analysis is in
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35 316 a large pan-European populations, we may have been under-powered to detect some
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37 317 significant interactions.
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319 **Implications of findings**

320 We demonstrate the importance of dietary and environmental influences on circulating
321 vitamin D concentration in a pan-European setting. Our findings also show that the
322 relationship between genes related to vitamin D metabolism and vitamin D status is
323 modified by sunlight exposure. These findings have important implications for the design of

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3 324 PN interventions targeted towards vitamin D and for the development of vitamin D-related
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5 325 policy on sunlight exposure. However, larger studies are needed to understand the clinical
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7 326 relevance of these findings for the health of European (and other) adults.
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11 12 13 328 **Conclusions**

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15 329 Our findings identify several cross-sectional dietary (vitamin D intake and supplement use)
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17 330 socio-demographic (age and sex) and environmental (country, season, sunlight exposure,
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19 331 physical activity and vitamin D axis genes) predictors of circulating 25(OH)D concentration in
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21 332 a pan-European population. We observed that carriage of the minor allele for *VDR*
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23 333 *rs2228570* genotype was associated with lower circulating 25(OH)D concentrations and that
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25 334 this relationship may be modulated by weekday sunlight exposure.
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31 32 33 336 **Author contributions**

34
35 337 YM, IT, CAD, ERG, LB, JAL, JAM, WHS, PW, HD, MG, TH and JCM contributed to the research
36
37 338 design. JCM was the Food4Me Proof of Principle study leader. CCM, ERG, LB and JCM
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39 339 contributed to the developing the Standardized Operating Procedures. CCM, CPL, GM, ALM,
40
41 340 RF and JCM conducted the intervention. CCM and WHS contributed to physical activity
42
43 341 measurements. UH, MB, IB, FFR, KG and JB contributed towards the vitamin D
44
45 342 measurements. KML and CCM wrote the paper and performed the statistical analysis and
46
47 343 are joint first authors. All authors contributed to a critical review of the manuscript during
48
49 344 the writing process. All authors approved the final version to be published.
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54 55 56 346 **Acknowledgement**

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4
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6
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11
12 351 **Conflict of interest statement**

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14
15 352 UH, MB, IB, FFR and PW are employed by DSM Nutritional Products. No other authors had a
16
17 353 personal or financial conflict of interest.
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21 354 **REFERENCES**

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Table 1 Associations between circulating 25(OH) vitamin D concentrations, socio-demographic and lifestyle characteristics

	25 OH- Vitamin D	P value
Age (year)	0.004 (0.002, 0.005)	0.002
Female (ref male)	-0.176 (-0.224, -0.127)	<0.001
Caucasian (ref other)	0.074 (-0.060, 0.207)	0.28
Country (ref British Isles)		
Western Europe	0.272 (0.208, 0.336)	<0.001
Southern Europe	0.190 (0.123, 0.258)	<0.001
Eastern Europe	0.185 (0.108, 0.261)	<0.001
Physical activity ³		
PAL	0.537 (0.281, 0.792)	<0.001
MVPA (MET/min/week)	0.047 (0.017, 0.077)	0.002
Sedentary behavior	-0.001 (-0.001, -0.001)	0.011
Vitamin D		
Supplement user (ref non-user)	0.242 (0.184, 0.300)	<0.001
Total Vitamin D intake (IU/d)	0.091 (0.056, 0.126)	<0.001
Vitamin D intake (food; IU/d)	0.055 (0.013, 0.096)	0.010
Weekday sunlight exposure (ref < 20 min/d)		
20 min to 2 h/d	0.038 (-0.028, 0.104)	0.225
> 2 h/d	0.133 (-0.042, 0.224)	0.004
Weekend sunlight exposure (ref < 20 min/d)		
20 min to 2 h/d	0.780 (-0.045, 0.201)	0.21
> 2 h/d	0.136 (0.009, 0.262)	0.036
Season (ref winter)		
Spring	0.155 (0.100, 0.210)	<0.001
Summer	0.263 (0.170, 0.356)	<0.001
Autumn	0.306 (0.197, 0.414)	<0.001

1, Values represent log-transformed (ln) coefficients (95% CI); PAL, physical activity level; MVPA, Moderate and vigorous physical activity; PA, physical activity.

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2, Multiple linear regression was used to test for significant differences between groups. Analyses were adjusted for age, sex, ethnicity, season, country, BMI (where appropriate), vitamin D intake from foods (where appropriate) and vitamin D supplementation.
3, PA measures were available in 1161 participants only

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Table 2 Associations between vitamin D-related genotype and circulating 25(OH) vitamin D concentrations

SNP	Minor allele frequency	Hardy Weinberg equilibrium (P value)	M/m	Gt	N	25 OH- vitamin D (nmol/L, 95% CI) ¹	P ²
VDR rs1544410	40.2	0.811	G/A	GG	467	54.9 (52.8, 57.0)	0.82
				AG	635	54.9 (53.1, 56.7)	
				AA	210	54.4 (51.4, 57.6)	
VDR rs2228570	38.8	0.654	C/T	CC	487	54.9 (52.9, 57.0)	0.74
				CT	631	55.0 (53.2, 56.8)	
				TT	194	54.0 (50.9, 57.3)	
GC rs2282679	27.2	0.259	A/C	AA	704	58.1 (56.3, 59.9)	<0.001
				CA	503	51.7 (49.8, 53.6)	
				CC	105	49.1 (45.3, 53.1)	
GC rs4588	28.6	0.912	C/A	CC	670	58.9 (57.1, 60.8)	<0.001
				CA	534	51.1 (49.3, 52.9)	
				AA	108	49.6 (45.9, 53.6)	
GC rs7041	43.0	0.824	G/T	GG	429	59.5 (57.2, 61.9)	<0.001
				TG	639	53.4 (51.7, 55.2)	

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TT 244 50.7 (48.1, 53.4)

1, Values represent adjusted means and 95% CI. SNP, single nucleotide polymorphism; Gt, genotype
2, Multiple linear regression was used to test for significant differences between genotype groups. Analyses were adjusted for age, sex, ethnicity, season, country, BMI, vitamin D intake from foods and vitamin D supplementation.

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3 **FIGURE LEGENDS**
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8 **Figure 1** Consort diagram of participants randomized into the Food4Me Proof of Principle
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10 Study. * Total number of participants reporting one or more exclusion criteria. SNP, single
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12 nucleotide polymorphism
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17 **Figure 2** Concentration of 25(OH) vitamin D according to vitamin D genetic risk score. Minor
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19 alleles for *GC rs2282679*, *GC rs4588*, *GC rs7041*, *VDR rs2228570*, *VDR rs1544410* were coded
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21 as 0, 1 and 2 and summed to generate the risk score and grouped into 0-2, 3 to 5 or 6 or
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23 more minor alleles.
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26 Values represent adjusted means and SE. Analyses were adjusted for age, sex, ethnicity,
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28 season, country, BMI, vitamin D intake from foods and vitamin D supplementation.
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33 **Figure 3** Impact of time spent in sunlight during a weekday on the relationship between
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35 Vitamin D receptor (VDR) gene SNP *VDR rs2228570* (*TT*) and 25(OH) vitamin D
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37 concentrations.
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40 Values represent adjusted means and SE. Analyses were adjusted for age, sex, ethnicity,
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42 season, country, BMI, vitamin D intake from foods and vitamin D supplementation. SNP,
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44 single nucleotide polymorphism
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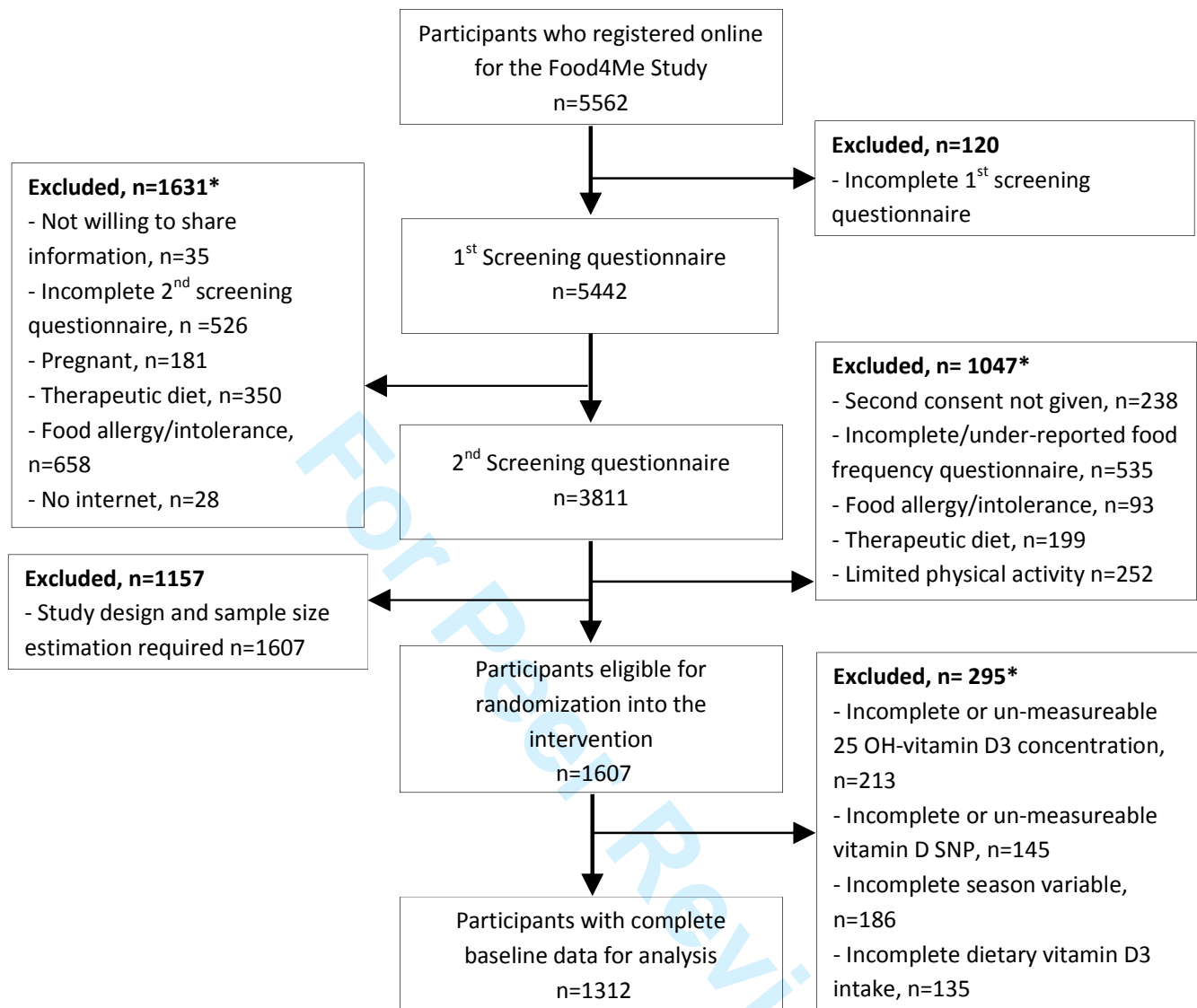


Figure 1

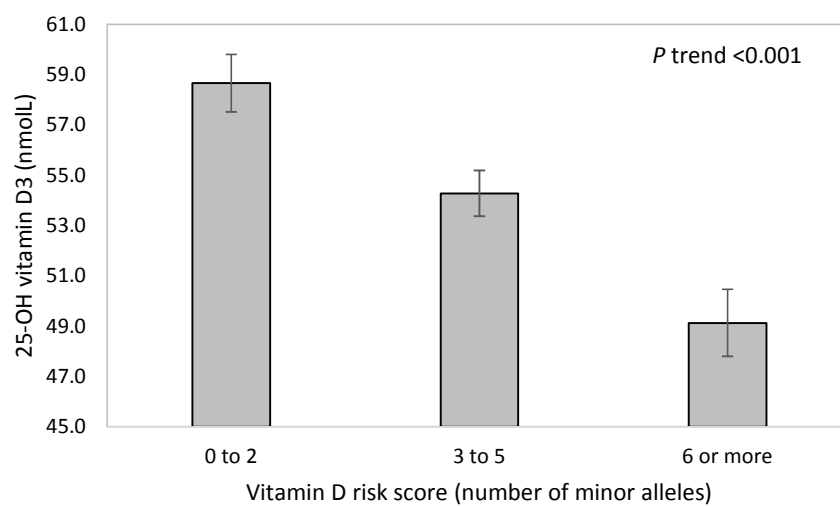


Figure 2

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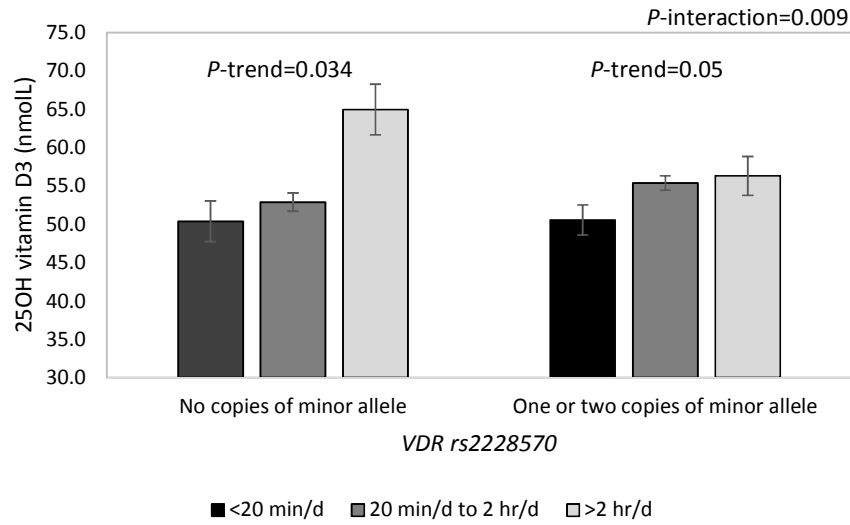


Figure 3

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