



Cite this: *Photochem. Photobiol. Sci.*, 2017, **16**, 381

Are low sun exposure and/or vitamin D risk factors for type 1 diabetes?

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The global variation in type 1 diabetes (T1D) incidence rates is one of the most significant observed for any non-communicable disease. Geographical patterns in incidence suggest that low sun exposure may contribute to the wide disparity, with incidence rates generally increasing with distance from the Equator. T1D development is associated with hyperactivity of the adaptive immune system leading to autoimmune destruction of insulin-secreting pancreatic β cells. Both exposure to ultraviolet radiation (UVR) and vitamin D, with their known immunosuppressive effects, have the potential to delay or inhibit the disease. Efforts to confirm the role of UVR by vitamin D dependent and independent pathways in the pathogenesis of T1D have been challenged by inconsistent results among studies. Human observational studies and animal and *in vitro* experiments indicate that at least some of the benefits of sun exposure come from improved vitamin D status. There is no evidence of benefit for T1D risk of vitamin D supplementation during pregnancy at current recommended levels (400 IU per day); but some evidence supports that higher sun exposure and/or vitamin D sufficiency in pregnancy, or supplementation in early life, decreases T1D risk. Further research is required to confirm an association between UVR exposure and T1D and clarify the mechanisms involved.

Received 10th August 2016,
Accepted 27th November 2016

DOI: 10.1039/c6pp00294c

rscl.li/ppp

Introduction

Type 1 diabetes (T1D) is the most common autoimmune disease of childhood.¹ The incidence of T1D is increasing worldwide – an annual increase of approximately 3%² – and there is evidence of a trend towards earlier age of onset.^{2–5} These relatively rapid changes implicate alterations in exposure to environmental agents as risk factors for the disease. Geographic patterns of increased incidence with greater distance from the Equator (higher latitude) provide a clue to environmental influences that may be important.⁶ Among several factors that vary according to latitude, levels of ultraviolet radiation (UVR) have been of particular interest because UV irradiation of the skin is the primary source of vitamin D. The active form of this pre-hormone has known effects on immune function that make vitamin D deficiency a plausible candidate risk factor for T1D.⁷ Increasing prevalence of vitamin D deficiency in children,⁸ occurring in parallel with

increasing incidence of T1D, and seasonal variation in the onset of T1D, have provided additional evidence that vitamin D deficiency may be a risk factor for T1D.

While observational studies support a link between vitamin D deficiency during pregnancy, typically measured as a serum/plasma level of serum 25-hydroxyvitamin D (25(OH)D) less than 50 nmol L⁻¹,⁹ and risk of T1D,^{10,11} trials of vitamin D supplementation have returned largely null results.¹² The most important determinant of serum 25(OH)D levels is sun exposure (including time in the sun and the intensity of UVR).^{13,14} This has led more recently to considerations of whether the 25(OH)D levels measured in observational studies (and related to T1D risk) are specific for vitamin D status or a proxy for recent sun exposure. UVR exposure of the skin suppresses adaptive immunity in ways that are similar to those of vitamin D.¹⁵ Thus exposure to UVR may modulate immune function relevant to the onset of T1D through both vitamin D and non-vitamin D pathways. Vitamin D supplementation improves only the former.

Here we review the evidence that low sun exposure, or vitamin D deficiency specifically, are associated with an increased risk of T1D. We restrict our analysis to T1D in children as the risk factors in this age group are likely to be clearer, with less effect of risk exposures later in life. We begin with an overview of the clinical, genetic and immune characteristics of T1D.

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Vitamin D as a risk factor for T1D

In this section we begin by providing an overview of the vitamin D metabolic pathway and then review the evidence of a link between vitamin D and risk of T1D, beginning with human observational studies and then exploring possible pathways of action using animal and *in vitro* studies. As associations with serum 25(OH)D levels do not allow differentiation of specific vitamin D effects rather than those of sun exposure; the focus of this section is therefore on vitamin D intake, including supplementation and evidence from associations between disease risk and polymorphisms in genes of the vitamin D pathway. The effects of environmental exposures on disease risks may depend on the age of exposure; we thus consider exposure at different age groups separately.

The vitamin D metabolic pathway

Endogenous synthesis of vitamin D in the skin begins with the photoconversion of 7-dehydrocholesterol (7-DHC) to pre-vitamin D. Single nucleotide polymorphisms within the *DHCR7* gene that encodes 7-dehydrocholesterol reductase (that converts 7-DHC to cholesterol) and the *NADSYN1* gene that produces NADPH that is required in this reaction, influence serum 25(OH)D levels.⁴⁴ Vitamin D is converted in the liver to 25(OH)D by the 25-hydroxylase enzyme encoded mainly in humans by the *CYP2R1* gene, with conversion to the active metabolite, 1,25(OH)₂D catalysed by the 1 α -hydroxylase enzyme encoded by *CYP27B1*. Within the blood, vitamin D metabolites are mainly tightly bound to a vitamin D binding protein, with a smaller proportion loosely bound to albumin or circulating as the free metabolite. The vitamin D binding protein gene (*GC*) has been repeatedly shown to be a major determinant of 25(OH)D levels,⁴⁴ as has the *CYP24A1* gene that encodes the breakdown enzyme, 24 hydroxylase. The active metabolite exerts its effects on gene transcription through a nuclear vitamin D receptor (*VDR*); polymorphisms within this gene may affect its activity and therefore its downstream functions. The most commonly studied *VDR* polymorphisms are *Fok1*, *Bsm1*, *Apa1* and *Taq1*. Their alleles are commonly referred to as letters (or as nucleotides), such as *Fok1* F/f (rs2228570 C/T, previously rs10735810), *Bsm1* B/b (rs1544410, A/G), *Taq1* T/t (rs731236, T/C) and *Apa1*, A/a (rs7975232, G/T). The *Fok1* “F” allele is more transcriptionally active than the “f” allele resulting in higher *VDR* production;^{45–48} the *Taq1* “t” allele may confer increased responsiveness to 1,25(OH)₂D.^{45–47,49}

Maternal dietary intake of vitamin D during pregnancy

In human observational studies, maternal intake of vitamin D supplements does not reduce risk of T1D in offspring. A protective association between maternal consumption of cod liver

oil supplements (400 IU vitamin D) and T1D was found in one pilot population-based case-control study⁵⁰ but was not replicated in a larger follow-up study.⁵¹ In both studies the use of vitamin D supplements was not significantly associated with a reduced risk of T1D and the authors were unable to establish whether the association found in the pilot study was a direct result of the 400 IU of vitamin D or the n-3 fatty acids (eicosapentaenoic acid and docosahexaenoic acid) found in the cod liver oil, or the combination of both. Consistent with these findings, a Finnish study⁵² based on a cohort of participants with genotypes for T1D conferring moderate or high risk, found no association between self-reported supplement use of vitamin D and presence of multiple autoantibodies/clinical diabetes. Similar findings were reported in a recent Swedish study.⁵³ However, in both studies the vitamin D intake reported from supplements was low, with only 15% of mothers consuming the recommended daily dose of 400 IU in the Finnish study and in the Swedish study the highest dose reported was 300 IU. For both studies, the baseline 25(OH)D levels of mothers were not known. The absence of such data limits the interpretation of the results. For example, if the majority of participants were deficient at baseline, then the small supplement doses reported would be unlikely to increase 25(OH)D levels to sufficient levels and an effect may not be expected. Equally, if mothers already had sufficient levels of 25(OH)D at baseline, then supplementation would equally not be expected to reduce risk. A meta-analysis of three studies confirmed that there was inadequate evidence to indicate an association between intake of vitamin D supplements in pregnancy and risk of T1D in the offspring.¹²

Higher maternal dietary intake of vitamin D *via* food or supplements during pregnancy has however been weakly associated with a decreased risk of developing T1D-related autoimmunity (positive for one or more of the three islet autoantibodies (GADA, IA-2, IAA)) in offspring in two cohort studies, though both reported notable discrepancies in their findings. The ABIS study indicated that the mothers' use of vitamin D-containing supplements (400 IU) during pregnancy reduced the odds of their offspring developing T1D-related autoimmunity by 29% (OR 0.71; 95% CI 0.52–0.96, *n* = 8694) at 1 year of age, but this effect was no longer apparent at 2.5 years (OR 1.25; 95% CI 0.09–1.73, *n* = 7766).⁵⁴ The authors were unable to explain why the association was lost at 2.5 years. The other study, of 233 American mothers (cases = 16), showed that the risk of T1D-related autoimmunity in offspring halved for each increase of 155.6 IU of vitamin D consumed during pregnancy *via* food per day (adjusted HR 0.37; 95% CI 0.17–0.78), but not supplements.⁵⁵ As the intake of vitamin D *via* food was considerably lower than the recommended adequate daily intake in the affected group, it is possible that it is deficiency that increases risk of autoimmunity rather than sufficiency being protective. Curiously, the results indicated an apparent 3-fold increase in risk among offspring of mothers who consumed vitamin D (400 IU) *via* supplements, which the authors did not discuss further. The data suggest that maternal supplement use increased the risk of offspring devel-



Table 1 The association between vitamin D supplementation in early life and subsequent risk of T1D

Publication	Supplementation	Age range	Dose	Frequency	Duration	Association
EURODIAB Study (1999) ¹⁶	Yes vs. no	Supplements first year of life	Not specified	Not specified	Less than a year vs. more than 1 year	Yes; OR = 0.67 (95% CI 0.53–0.86)
Stene <i>et al.</i> (2000) ⁵⁰	Yes vs. no (less than once per week)	Supplements first year of life	Speculated 400 IU	Less than once a week, 1–4 times a week, nearly everyday	Not specified	Cod liver oil – no; OR = 0.82 (95% CI 0.47–1.42) Other vitamin D supplements no; OR = 1.27 (95% CI 0.70–2.31)
Hypponen <i>et al.</i> (2001) ⁵⁹	Regular vs. none Irregular vs. none	Supplements first year of life	<2000 IU, within 2000 IU, >2000 IU	Regular, irregular and none	Not specified	Irregular – yes; OR = 0.16 (95% CI 0.04–0.74) Regular – yes; OR = 0.12 (95% CI 0.03–0.51)
Stene <i>et al.</i> (2003) ⁵¹	Yes vs. no (less than once per week)	Supplements first year of life	Speculated 400 IU Vit D supplements and cod liver oil supplements	Less than once a week, 1–4 times a week, nearly everyday	Not specified	Cod liver oil 1–4 times per week – no; OR = 0.81 (95% CI 0.55–1.19) Cod liver oil ≥5 times per week – yes; OR = 0.74 (95% CI 0.56–0.99) Other vitamin D supplements 1–4 times per week – no; OR = 0.99 (95% CI 0.69–1.42) Other vitamin D supplements 1–4 times per week – no; OR = 0.97 (95% CI 0.73–1.29)
Visalli <i>et al.</i> (2003) ⁵⁸	Yes vs. no	Supplementation during “early years” – no time period provided	Not specified	Not specified	Not specified	No; OR = 1.22 (95% CI 0.82–1.83)
Tenconi <i>et al.</i> (2007) ⁵⁷ Ahadi <i>et al.</i> (2011) ¹⁷⁵	Yes vs. no Yes vs. no	Vitamin D during lactation Supplements first year of life	Not specified Not specified	Not specified Not specified	Not specified Not specified	Yes; OR = 0.33 (95% CI 0.14–0.81) Yes; lack of vitamin D supplementation OR = 3.78 (95% CI 1.60–8.89)





Table 2 Vitamin D pathway genes and risk of T1D

Publication	Case/control	Gene	Allele	Association with T1D risk
Cooper <i>et al.</i> (2011) ¹⁷⁶ Frederiksen <i>et al.</i> (2013) ¹⁷⁷	8517/10438 1708 high genetic risk; 148 IA; 62 IA and T1D	<i>DHCR7</i> <i>DHCR7/NADSYN1</i>	rs12785878	G allele (<i>cf.</i> T allele): OR = 1.07 (95% CI 1.02–1.13) Increased risk of IA but not T1D. HR = 1.36 (95% CI 1.08–1.73) for each additional minor allele
Thorsen <i>et al.</i> (2014) ¹⁷⁸ Cooper <i>et al.</i> (2011) ¹⁷⁶ Hussein <i>et al.</i> (2012) ¹⁷⁹ Ramos-Lopez <i>et al.</i> (2007) ⁷⁰	1467 trios (907 cases, 896 sibs) 8517/10438 120/120 203 simplex T1D families (<i>n</i> = 609)	<i>DHCR7</i> <i>CYP2R1</i> <i>CYP2R1</i> <i>CYP2R1</i>	rs12785878 rs12794714 rs10741657 rs10741657	G allele (<i>cf.</i> T allele): OR = 0.93, <i>p</i> = 0.21 T allele (<i>cf.</i> C allele): OR = 1.04 (95% CI 1.00–1.09) GG associated with increased risk Variant G more often transmitted to affected offspring and more frequent in cases than controls
Thorsen <i>et al.</i> (2014) ¹⁷⁸ Blanton <i>et al.</i> (2011) ¹⁸⁰ Blanton <i>et al.</i> (2011) ¹⁸⁰ Thorsen <i>et al.</i> (2014) ¹⁷⁸ Cooper <i>et al.</i> (2011) ¹⁷⁶ Cooper <i>et al.</i> (2011) ¹⁷⁶ Cooper <i>et al.</i> (2011) ¹⁷⁶ Bailey <i>et al.</i> (2007) ¹⁸¹ Bailey <i>et al.</i> (2007) ¹⁸¹ Fichna <i>et al.</i> (2010) ¹⁸²	1467 trios (907 cases, 896 sibs) 203/153/116 first degree relatives 203/153/116 first degree relatives 1467 trios (907 cases, 896 sibs) 8517/10438 8517/10438 8517/10438 7854/8758 7854/8758 215/236	<i>CYP2R1</i> <i>GC</i> <i>GC</i> <i>GC</i> <i>GC</i> <i>CYP27B1</i> <i>CYP27B1</i> <i>CYP27B1</i> <i>CYP27B1</i>	rs12794714 rs10741657 rs4588 rs7041 rs2282679 rs4588 rs7041 rs10877012 rs10877012 rs4646536	A allele (<i>cf.</i> G allele): OR = 1.01, <i>p</i> = 0.86 A allele (<i>cf.</i> C allele): OR = 1.05 (95% CI 0.91–1.20) T allele (<i>cf.</i> G): OR = 1.07 (95% CI 0.92–1.24) C allele (<i>cf.</i> A allele): OR = 1.01, <i>p</i> = 0.90 A allele (<i>cf.</i> C allele) OR = 0.95 (95% CI 0.91–1.00) T allele (<i>cf.</i> G allele): OR = 0.98 (95% CI 0.93–1.03) A allele (<i>cf.</i> G allele): OR = 0.93 (95% CI 0.89–0.98) C allele (<i>cf.</i> A): OR = 1.07 (95% CI 1.02–1.13) T allele (<i>cf.</i> C allele): OR = 1.08 (95% CI 1.02–1.14) <i>p</i> = 0.67 for difference in allele frequency between cases and controls
Hussein <i>et al.</i> (2012) ¹⁷⁹ Lopez <i>et al.</i> (2004) ¹⁸³	120/120 252/320	<i>CYP27B1</i> <i>CYP27B1</i>	rs10877012 rs10877012	CC associated with increased risk T1D associated with allelic variation in the promoter (rs10877012) polymorphism (<i>p</i> = 0.003) but not the intron 6 (rs4646536) polymorphism
Frederiksen <i>et al.</i> (2013) ¹⁷⁷	1708 high genetic risk; 148 IA; 62 IA and T1D	<i>CYP27B1</i>	rs4646536 rs4646536	Increased risk of IA but not T1D HR = 0.59, 0.39–0.89 for A/G <i>cf.</i> G/G
Thorsen <i>et al.</i> (2014) ¹⁷⁸ Abd-Allah <i>et al.</i> (2014) ¹⁸⁴	1467 trios (907 cases, 896 sibs) 120/120	<i>CYP27B1</i> <i>VDR</i>	rs4646536 Bsm1; (BB)	C allele (<i>cf.</i> T allele): OR = 0.96, <i>p</i> = 0.48 Bb: AOR = 2.1 (95% CI 1.1–3.2); bb: AOR = 1.7 (95% CI 1.0–1.9)
Cooper <i>et al.</i> (2011) ¹⁷⁶ Garcia <i>et al.</i> (2007) ¹⁸⁵	8517/10438 216/203	<i>VDR</i> <i>VDR</i>	Bsm1 Bsm1 (BB)	A allele (<i>cf.</i> G allele): OR = 1.00 (95% CI 0.95–1.05) Frequency of b allele and bb genotype significantly lower in T1D cases, <i>p</i> < 0.04
Thorsen <i>et al.</i> (2014) ¹⁷⁸ Capoluongo <i>et al.</i> (2006) ¹⁸⁶	1467 trios (907 cases, 896 sibs) 246/246	<i>VDR</i> <i>VDR</i>	Bsm1 Bsm1 (BB)	T allele (<i>cf.</i> C allele): OR = 0.94, <i>p</i> = 0.22 Bb: OR = 1.01 (95% CI 0.64–1.59); bb: OR = 0.92 (95% CI 0.54–1.57)
Lemos <i>et al.</i> (2008) ¹⁸⁷ Abd-Allah <i>et al.</i> (2014) ¹⁸⁴	207/249 120/120	<i>VDR</i> <i>VDR</i>	Bsm1 Fok1; (FF)	G allele (<i>cf.</i> A allele): OR = 1.01 (95% CI 0.78–1.31) Ff: AOR = 1.7 (95% CI 1.0–2.7); Ff: AOR = 3.8 (95% CI 1.2–9.4)
Hamed <i>et al.</i> (2013) ¹⁸⁸ Thorsen <i>et al.</i> (2014) ¹⁷⁸ Cooper <i>et al.</i> (2011) ¹⁷⁶ Capoluongo <i>et al.</i> (2006) ¹⁸⁶	132/40 1467 trios (907 cases, 896 sibs) 8517/10438 246/246	<i>VDR</i> <i>VDR</i> <i>VDR</i> <i>VDR</i>	Fok1 Fok1 Fok1 Fok1 (FF)	f allele (<i>cf.</i> F allele): OR = 1.08 (95% CI 0.64–1.85) T allele (<i>cf.</i> C allele): OR = 0.99, <i>p</i> = 0.85 A allele (<i>cf.</i> G allele): OR = 0.99 (95% CI 0.95–1.04) Ff: OR = 0.90 (0.60–1.35); ff: OR = 1.64 (95% CI 0.91–2.97)
Lemos <i>et al.</i> (2008) ¹⁸⁷ Abd-Allah <i>et al.</i> (2014) ¹⁸⁴	207/249 120/120	<i>VDR</i> <i>VDR</i>	Fok1 Apa1 (AA)	T allele (<i>cf.</i> C allele): OR = 0.93 (95% CI 0.71–1.22) Aa: AOR = 0.6 (95% CI 0.5–1.0); aa: AOR = 0.6 (95% CI 0.2–1.1)
Thorsen <i>et al.</i> (2014) ¹⁷⁸ Garcia <i>et al.</i> (2007) ¹⁸⁵	1467 trios (907 cases, 896 sibs) 216/203	<i>VDR</i> <i>VDR</i>	Apa1 Apa1	C allele (<i>cf.</i> A allele): OR = 0.99, <i>p</i> = 0.92 <i>p</i> = NS for the difference in allele frequency between cases and controls
Lemos <i>et al.</i> (2008) ¹⁸⁷	207/249	<i>VDR</i>	Apa1	T allele (<i>cf.</i> G allele): OR = 1.01 (95% CI 0.77–1.31)

the low dose of KH1060 mediated some protection without eliciting significant hypercalcaemic effects (60% decrease in incidence). As seen with $1,25(\text{OH})_2\text{D}_3$, the increased protection provided by the higher dose of KH1060 (400 ng) and the $1(\text{OH})\text{D}_3$ analog coincided with elevated serum calcium levels.

In addition to reducing the incidence of T1D, non-hypercalcaemic analogs of $1,25(\text{OH})_2\text{D}_3$ inhibited the progression of insulinitis to clinical diabetes and prevented recurrence of T1D following islet transplantation. An experiment using NOD mice tested the effectiveness of MC1288 (a non-hypercalcaemic analog of $1,25(\text{OH})_2\text{D}_3$) with and without a short course of the anti T cell immunosuppressant, cyclosporine A (CyA), in inhibiting the progression of insulinitis to clinical diabetes. Monotherapy of CyA or MC1288 was ineffective in preventing the progression to overt disease at 200 days, however, the combined treatment of MC1288 with CyA, significantly reduced diabetes incidence to 35% when compared with the control group (65%).⁸⁶ In another study, a $1,25(\text{OH})_2\text{D}_3$ analog, BXL-219 (formerly RO 26-2198), given to adult NOD mice, prevented progression to overt disease by 38 weeks (90% in the control group *vs.* 16% in the NOD mice treated for 16 weeks). In mice that did not progress to overt disease, there was an increase in the frequency of T regulatory cells (CD4+ CD25+) in pancreatic lymph nodes that specifically inhibited T cell responses to the autoantigen IA-2.⁸⁷ BXL-219 can downregulate proinflammatory chemokine production by pancreatic islets leading to reduction in T-cell recruitment.⁸⁸

KH1060 also delayed recurrence of T1D in NOD mice following islet transplantation. Islet survival was significantly prolonged in NOD mice treated with KH1060 or cyclosporine1 (60 days and 50 days, respectively, *versus* 9.5 days in controls; $p < 0.001$ and $p < 0.0001$); however mice treated with sub-therapeutic doses of both drugs also showed prolonged graft survival (48 days; $p < 0.0001$). Furthermore, 80% of the mice that remained normoglycemic to 60 days post-transplantation remained disease free for more than 15 days after the cessation of all treatment.⁸⁹

Streptozotocin (STZ), an antibiotic that causes destruction of pancreatic β cells, can be used to create rodent models of T1D. The model has been useful in elucidating associations between insulin and various stages in the vitamin D metabolic pathway. In the STZ diabetic rat, induction of T1D was associated with reduction of $1,25(\text{OH})_2\text{D}$ levels that was reversible with insulin therapy.⁹⁰ Furthermore, induced diabetes resulted in a decrease in vitamin D binding protein concentration, so that total $1,25(\text{OH})_2\text{D}$ was reduced. Notably, the concentration of free $25(\text{OH})\text{D}$ and $1,25(\text{OH})_2\text{D}$ were unchanged.⁹¹

In this model, insulin has a direct stimulatory effect on the 1α -hydroxylase enzyme that converts $25(\text{OH})\text{D}$ to the active form, and also alters the responsiveness of renal 1α -hydroxylase to the usual cue of lowered phosphate levels and changes in PTH.^{92,93} In the chronic insulin-deficient state there is significantly reduced 1α -hydroxylase activity and enhanced renal 24 -hydroxylase activity. Similar actions in humans could explain the lower $1,25(\text{OH})_2\text{D}$ levels and increased $24,25(\text{OH})_2\text{D}$ levels that are commonly seen in chil-

dren with T1D,^{94,95} *i.e.* changes in vitamin D metabolism are the result of insulin deficiency, rather than the cause.⁹⁶

In summary, $1,25(\text{OH})_2\text{D}_3$, vitamin D_3 or their structural analogs can delay or inhibit the development of T1D when administered at pharmacological doses from weaning to end of life. Daily oral administration appears to be more effective in preventing the onset of T1D than intraperitoneal administration given on alternate days. Elevated calcium levels may be required for maximal benefit from vitamin D supplementation. Structural analogues provided alone or in conjunction with an immunosuppressant agent can reduce progression to overt disease after the development of autoantibodies, and recurrence after islet transplantation.

In vitro studies

Human pancreatic tissue expresses 1α -hydroxylase and VDR⁹⁷ and a vitamin D response element has been identified in the human insulin receptor gene promoter.⁹⁸ Suspensions of rat islet cells can convert $25(\text{OH})\text{D}$ to $1,25(\text{OH})_2\text{D}$ and acute application of high dose $1,25(\text{OH})_2\text{D}$ (50 nmol L^{-1}) causes a marked rise in calcium levels, confirming that the cells are responding to $1,25(\text{OH})_2\text{D}$.⁹⁷ That is, pancreatic tissue can use circulating $25(\text{OH})\text{D}$ to make $1,25(\text{OH})_2\text{D}$ locally, and pancreatic cells are responsive to $1,25(\text{OH})_2\text{D}$.

Immune cells from people with T1D have been cultured with $25(\text{OH})\text{D}$ or $1,25(\text{OH})_2\text{D}$. The doses used are often much higher than is seen in serum; it is proposed that local production of vitamin D metabolites can lead to high local levels. In monocytes from patients with T1D and healthy controls, culture with $25(\text{OH})\text{D}$ (125 nmol L^{-1}) significantly inhibited the differentiation of monocytes into dendritic cells, increasing the number of intermediate cells (CD11c+ CD14+ CD83+ CD123–/low). These cells had a similar phenotype to previously described DC-10 cells, which produce IL-10 and have tolerogenic characteristics because they can induce T regulatory cells.⁹⁹ This increase in response to $25(\text{OH})\text{D}$ depended on the VDR genotype: smaller increase in intermediate cells in genotype bb (of BsmI) than BB and Bb; and a smaller increase for genotype TT compared to tt (TaqI).⁷² In T-helper (Th) cells isolated from patients with T1D and healthy controls that were stimulated with $25(\text{OH})\text{D}$ (125 nmol L^{-1}), there was significantly lower VDR expression in cells from patients with T1D compared to those from healthy controls. There were significantly fewer CD4+ cells in $25(\text{OH})\text{D}$ - and $1,25(\text{OH})_2\text{D}$ -stimulated Th cells from T1D patients carrying the FF genotype of FokI compared to those with Ff/ff ($p = 0.02$). This suggests that vitamin D supplementation may be more effective for prevention and/or management of T1D in those carrying the FF genotype, through promoting a more regulatory T-cell milieu.¹⁰⁰ However, in another study, in isolated CD4+ memory cells, FOXP3 expression was higher in cells from Estonian compared to Finnish children with T1D even in those who were vitamin D sufficient. The authors concluded that the findings did not support a crucial role for $25(\text{OH})\text{D}$ as



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