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

K. M. Miller,^{*a} P. H. Hart,^a N. H. de Klerk,^a E. A. Davis^b and R. M. Lucas^c

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.

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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.

^aTelethon Kids Institute, 100 Roberts Road, Subiaco, Western Australia 6008, Australia. E-mail: kate.tan@telethonkids.org.au

^bPrincess Margaret Hospital, Roberts Road, Subiaco, Western Australia 6008, Australia

^cNational Centre for Epidemiology and Population Health, The Australian National University, Canberra 2600, Australia



Type 1 diabetes

T1D results from the progressive autoimmune destruction of insulin-producing beta (β) cells in the pancreas^{16,17} sufficient to decrease and ultimately cease insulin production.¹⁸ Although genetic, serological and biochemical testing can be used to identify at-risk individuals,¹⁹ there is no established mechanism for slowing or preventing the progression of β -cell destruction in humans.²⁰ Similarly, there is no cure for T1D.²¹

T1D can present at any age, but onset is typically in childhood or adolescence.²² Children with T1D commonly present to health services with acute clinical symptoms including polydipsia and polyuria, nocturia, enuresis and weight loss (which can occur despite polyphagia), and blurred vision, or more severe conditions such as diabetic ketoacidosis, severe dehydration or shock.¹⁹ These symptoms occur only once the majority (90%) of the β -cells are destroyed or significantly compromised, rendering the child insulin dependent.¹⁸

The incidence of T1D among children aged 0–14 years varies significantly worldwide with rates ranging from 0.6 per 100 000 in China to 62.3 per 100 000 in Finland.² In 2015, the International Diabetes Federation reported that over half a million children worldwide are living with T1D, and an additional 86 000 are estimated to be diagnosed each year.²

Children with T1D are more likely to develop a range of disabling and life threatening conditions when compared to children without the disease,² of which diabetic ketoacidosis is most common.¹ Many of the complications associated with T1D are the result of chronic hyperglycaemia and are now being observed at a younger age as a consequence of the earlier onset of the disease.¹ In addition to the physical health problems, T1D also places psychological pressure on both those diagnosed and their families.¹ Following a diagnosis significant vigilance is required to minimise glucose variability and sustain optimal diabetes management.²³ A recent large population based cohort study out of Sweden has shown that children with T1D were 2.1 times more likely to be diagnosed with a psychiatric disorder and 1.7 times more likely to attempt suicide when compared with controls.²⁴ There are also considerable financial costs associated with having the disease. Direct costs associated with treatment, monitoring and health service utilisation, in addition to more indirect costs resulting from carer support, disability and loss of productivity are shared by individuals, their families, health services and government bodies.² Globally the cost of treating and managing T1D in 2015 was estimated to be between USD 673–1197 billion.²

Children diagnosed with T1D have a 2–4 fold increased mortality^{25–28} most commonly the result of acute complications, such as diabetic ketoacidosis or other diabetes related events, death due to unnatural causes and sudden unexpected death. Despite improvements in the treatment and care provided to those with T1D, the average life expectancy of those diagnosed in childhood is still reported to be approximately

10 years less than that observed in the general population.^{18,29} This is infinitely better than the prognosis without insulin therapy, and a vast improvement from the 20 year reduction in life expectancy reported some 50 years ago.³⁰

The aetiology of T1D is complex. Both genetic and environmental factors are involved and possibly gene-environment interactions. The rapid rate at which T1D is increasing worldwide and the trend towards earlier onset point to a significant role for environmental influences.^{1,31,32}

Multiple genes have been linked with increased susceptibility to T1D; most of the risk loci are associated with immune cells and their function.³³ Major susceptibility genes are in the HLA region.¹⁸ From this region *DR3-DQ2* and *DR4-DQ8* haplotypes are most strongly associated, evident in up to 90% of those with T1D.³⁴ However, less than 10% of genetically susceptible individuals go on to develop T1D³⁵ and only 13–34% of monozygotic twins are pairwise concordant for the disease.^{36–40}

Putative non-genetic/environmental risk factors include maternal and child enterovirus infection, increased maternal age, rapid growth in early childhood, obesity, stress³² and vitamin D deficiency in early life.⁴¹ The idea that modifiable risk factors may be involved in the pathogenesis of T1D introduces great hope and purpose for preventative efforts. There is mounting evidence that non-genetic and environmental factors may act *in utero*, in early infancy and/or in childhood to modify the risk of T1D, particularly for those who are not genetically susceptible.⁴²

T1D development is associated with hyperactivity of the adaptive immune system (T and B lymphocytes) leading to autoimmune destruction of insulin-secreting pancreatic β cells. Amongst the inflammatory cells of insulinitis lesions, CD8⁺ T lymphocytes are the most prominent, followed by macrophages, CD4⁺ T lymphocytes, B lymphocytes and plasma cells. The first three mentioned cell types can enhance immune responses by production of molecules, enzymes and cytokines with tissue destructive properties, as well as chemokines to attract further inflammatory cells. The demise of pancreatic β cells may be an off-target effect of activated cytotoxic CD8⁺ cells. Activation of B lymphocytes and plasma cells is already evident at the time of first diagnosis of T1D. As reviewed elsewhere,⁴³ ninety percent of newly diagnosed patients with T1D already have antibodies reactive to at least one antigen expressed by β cells, namely to insulin (anti-insulin antibodies, IAA), glutamic acid decarboxylase (GADA), insulinoma-associated autoantigen 2 (IA-2) or zinc transporter 8. Of interest, T regulatory lymphocytes, an important cell type that would normally control over-zealous inflammatory responses, are rare in inflammatory regions of the pancreas.⁴³

While a wide range of environmental factors are implicated in risk of T1D, the immunopathology of the disease and the known immunomodulatory effects of vitamin D and other molecules produced in skin following exposure to UVR, make these exposures of particular interest as potential regulators of development and progression of T1D.



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)



Table 2 (Contd.)

Publication	Case/control	Gene	Allele	Association with T1D risk
Abd-Allah <i>et al.</i> (2014) ¹⁸⁴	120/120	VDR	Taq1 (TT)	Tt: AOR = 0.6 (0.4–1.2); tt: AOR = 0.7 (95% CI 0.4–1.3)
Lemos <i>et al.</i> (2008) ¹⁸⁷	207/249	VDR	Taq1	T allele (cf: C allele): OR = 0.90 (95% CI 0.69–1.18)
García <i>et al.</i> (2007) ¹⁸⁵	216/203	VDR	Taq1	<i>p</i> = NS for the difference in allele frequency between cases and controls
García <i>et al.</i> (2007) ¹⁸⁵	216/203	VDR	haplotype	BAT higher in cases (<i>p</i> = 0.002); BAT higher in controls (<i>p</i> = 0.003); AabbTT and aabbTT associated with higher levels of GADA, IA-2 and TGF- β
Cooper <i>et al.</i> (2011) ¹⁷⁶	8517/10438	CYP24A1	rs2296241	A allele (cf: G allele): OR = 1.00 (95% CI 0.95–1.05)
Bailey <i>et al.</i> (2007) ¹⁶¹	7854/8758	CYP24A1		Null (<i>p</i> = 0.23)
Thorsen <i>et al.</i> (2014) ¹⁷⁸	1467 trios (907 cases, 896 sibs)	CYP24A1	rs6013897	A allele (cf: T allele): OR = 1.03, <i>p</i> = 0.65
Lopez <i>et al.</i> (2004) ¹⁸³	187 families with one child with T1D	CYP27B1	–1260 C/A	Haplotype CT (–1260 A/2338 T) was significantly more often transmitted to affected offspring; AT (–1260 C/2838 T) was significantly less often transmitted (<i>p</i> = 0.02)
Miettinen <i>et al.</i> (2015) ⁶⁸	Mothers (<i>n</i> = 512), family members (<i>n</i> = 708), & child with T1D (<i>n</i> = 534) cf: healthy control children (<i>n</i> = 381), mothers (379) and fathers (<i>n</i> = 340)	VDR	2338 T/C (31 SNPs)	Maternal rs1544410 and rs731236 significant after adjustment for multiple testing. No difference in allele frequency between case and control children.

Unfortunately, the translation of these beneficial results from NOD mice to humans has been challenging due to the hypercalcaemic and/or bone effects resulting from the large doses that have been required to achieve disease protection.

Administration of 1,25(OH)₂D₃ can at least partially prevent the development of insulinitis and T1D in NOD mice.^{77–80} Daily oral administration of pharmacological doses (50 ng) from weaning to end of life completely prevented the development of the disease.⁷⁵ However, a lower calcaemic dose of 10 ng day^{–1} offered only partial protection, suggesting that hypercalcaemia may be required for complete protection. This was supported by another study that found reduced disease protection with lower calcium levels, when comparing outcomes of NOD mice treated with 1,25(OH)₂D₃ and its synthetic precursor 1 α OH D₃. Similar effects have been demonstrated in other autoimmune disorders such as experimental autoimmune encephalomyelitis (EAE, the animal model of MS).^{80,81}

Initially all studies that had shown beneficial effects of vitamin D on T1D incidence administered 1,25(OH)₂D₃ and the first study to administer the precursor, vitamin D₃, produced null results.⁸² This led to the belief that vitamin D₃ administration was ineffective in reducing disease risk; however, more recent studies have refuted this⁸³ and it is likely that the short regimen (*in utero* to 70 days) was insufficient to interfere with disease progression in NOD mice.

Different treatment windows for vitamin D supplementation for preventing the onset of T1D have been tested. So far, administration of vitamin D₃ during pregnancy or early life (up to 70 days), even at high doses (1000 IU per day) has had little to no effect on diabetes development in NOD mice or their offspring.^{82–84} It has been suggested that during this early period, training of immune cells is yet to occur and thus escapes the immunomodulatory actions of vitamin D.⁸³ In one study NOD mice were treated with 800 IU of vitamin D₃ during pregnancy, early life or lifelong. The study reported that vitamin D supplementation could only significantly prevent T1D development in both male and female NOD mice if administered lifelong. Lifelong supplementation resulted in a 58% and 52% reduction in diabetes incidence, respectively, when compared with control mice.⁸³ These findings were corroborated by Gysemans and colleagues (2005) suggesting that only lifelong treatment, initiated immediately after weaning, resulted in significant protection from developing T1D (66% risk reduction; *p* ≤ 0.0001).⁸⁵

A concern from many of the NOD mouse studies was that the doses required to elicit therapeutic benefit resulted in hypercalcaemia and/or bone demineralization. In response there have been concerted efforts worldwide to develop structural analogs of 1,25(OH)₂D₃ which offer more pronounced immunomodulatory effects without calcaemic side effects. KH1060, a structural analogue of 1,25(OH)₂D₃, administered at doses of 200 ng or 400 ng KH1060 per kg intraperitoneally in 0.05 ml arachis oil on alternate days, delayed and prevented onset of insulinitis and T1D in NOD mice,⁷⁹ and in another study the 1(OH)D₃ analog provided greater disease protection when compared with 1,25(OH)₂D₃.⁸⁰ Of the two analogs, only

environmental and behavioural factors. However, migrant studies which show that the rates among those from low incidence countries assimilate quickly to those of the country they live in^{118,119} suggest that environmental effects may be more important than those of ethnicity.

Seasonal variation in month of birth

Many studies have examined the effect of month of birth on T1D risk. However, whether an association exists remains an open question. A significant difference in birth seasonality pattern in children who were later diagnosed with T1D compared with the seasonal pattern of total live births has been found in several registers worldwide, with most studies reporting an excess in late spring/summer^{120–128} and/or a trough in late autumn/winter,^{123,124} though this is not completely consistent.^{129–131} One study reported that a diagnosis of T1D was 30% more common among those born in spring than in winter.¹²⁷ Some studies have reported seasonal variation in sub-populations (such as males and homogenous populations) only,^{122,124,131–133} and many have reported no difference in seasonality pattern from that of the general population.^{31,134–137}

It has been suggested that the apparent seasonal month of birth pattern among those diagnosed with T1D is due to the well-established relationship between season and serum 25(OH)D levels observed in many countries throughout the world and the immunosuppressive effects of UVR and/or vitamin D.³⁵ An excess in late spring/summer births coincides with the lowest serum 25(OH)D levels being experienced in the second trimester of pregnancy. Maternal vitamin D insufficiency during the second trimester may affect in utero development of the pancreatic β -cells¹³⁸ or regulate the developing immune system in a way that increases the risk of T1D in childhood.³¹

Of note, in countries closest to the Equator, where cutaneous production of vitamin D is possible throughout the year, or countries where dietary intake of vitamin D is significant, there is little seasonal variation in 25(OH)D levels and no evidence of a birth-month effect for T1D. A number of nutritional studies have found that the commonly observed association between season and 25(OH)D levels was negated by increased supplement use¹³⁹ or greater consumption of fatty fish⁶ in winter months. If vitamin D status were in some way responsible for the seasonal variation in birth rate among those diagnosed with T1D then we would expect that not all studies, particularly those conducted in countries where 25(OH)D levels are relatively constant throughout the year, would observe an association.

An absence of birth-month effect is more commonly reported in countries of low incidence of T1D,^{132,137} likely somewhat the result of small sample sizes but also pointing to ethnicity as a potential confounding factor.¹⁰⁶ A large cohort study spanning Europe (37–53°N), Australia (34°N), USA (39–40°N) and Israel (31°N) was one of several studies that found no seasonal pattern in month of birth in low incidence countries, reporting a seasonal pattern only in ethnically homo-

genous populations that had a medium to high incidence rate of T1D.¹³² It has been suggested by others that ethnicity may be a modifier due to differences in circulating vitamin D binding protein between ethnic groups,¹⁴⁰ in addition to differences in sun protective behaviours and diet. Beyond mechanisms of UVR and/or subsequent vitamin D production, alternative explanations for a birth-month effect are thought to involve seasonal exposures to viruses and infections.¹²¹ Overall the evidence supporting an association between month of birth and T1D is unclear. Despite some consistency across studies, factors such as ethnicity, latitude and background incidence appear to be important, making it difficult to draw conclusions in relation to an effect of sun exposure or vitamin D.

Seasonality of diagnosis of T1D

Seasonality of diagnosis of T1D has been extensively studied and reported in both northern^{114,122,133,141–145} and southern hemisphere studies.^{31,141,146–151} In many countries, seasonality of diagnosis follows a cyclic, sinusoidal pattern¹⁴⁹ with a peak occurring in cooler winter months,^{143,148–150,152–154} or late autumn/winter^{122,141,155} and a nadir in spring and summer.^{122,143,155} Unlike month of birth, seasonality of diagnosis is now a well-recognised feature of T1D in childhood.¹⁴¹

Seasonal variation in diagnosis is commonly more marked in boys,^{133,141,142,144,145,148,152} older children (5–14 years)^{122,141–144,152,156} and in countries which experience higher incidence and prevalence of T1D.¹⁴¹ Larger studies offering greater statistical power are required to confirm these findings¹⁰⁷ and identify the underlying reasons for the observed differences between sub populations.

While somewhat consistent, the source or sources of the above associations are not clear. It is likely that the birth month effect and seasonal variation in diagnosis of T1D arise by different mechanisms. Given the time that it takes for T1D to manifest, it is more likely that an excess in diagnosis of T1D in cooler winter months is more indicative of a season-dependent precipitating factor that triggers the presentation of symptoms or acceleration of β -cell destruction among those at an advanced stage of disease development, rather than a cause-of-disease relationship.¹⁸ Viral infections, temperature, low UVR exposure, or low 25(OH)D levels are likely influencing factors as they follow similar cyclic patterns to disease onset and have also been linked with the pathogenesis of T1D. However, we need more precisely characterised risk factors to demonstrate that there is variation in disease onset between seasons that persists after adjusting for confounding factors, and to allow a better understanding of the underlying reasons for the association.

Temperature

Temperature is one of the possible mechanisms underpinning the observed seasonal variation in diagnosis, and recent human observational studies have reported an independent



association between temperature and T1D onset.^{145,157} Cooler temperatures affect insulin production by triggering the release of norepinephrine which increases hepatic and peripheral insulin resistance.¹⁵⁷ The increase in demand for insulin places significant load on the remaining operating β -cells causing them to fail. In the advanced stages of disease development when β -cells are limited, this acute overload may lead to the presentation of symptoms and subsequent diagnosis in cooler months.

Ambient UVR

One environmental factor that varies by both latitude and season is ambient UVR. Latitude, season, and sunshine hours have been used as proxies for UVR, however technological advances have allowed more recent studies to measure ambient UVR. To our knowledge no study has measured personal exposure to UVR and examined its independent effect on T1D risk so here we review studies that have used ambient UVR as a marker of sun exposure.

Data on ambient UVR collected from various satellites have been linked with registers in Australia, Canada, and Sweden and from the Diabetes Mondial Project Group. Two Australian and one Canadian study found that T1D incidence/prevalence was significantly and independently inversely correlated with levels of ambient UVR.^{31,110,153} One of the Australian studies showed that the relationship was non-linear and moreover, the inverse relationship appeared to be restricted to low population density areas and with an apparent positive relationship in areas of high population density.³¹ It was hypothesised that the relationship between ambient UVR and actual exposure may be lost in cities (high population density) because of tall buildings and a less outdoors lifestyle. Data from the Diabetes Mondial Project Group showed that there was an inverse association between incidence of T1D and UVB irradiance which persisted after controlling for per capita health expenditure.⁶ However, while there was an overall inverse trend based on the combined data, there were a number of discrepant regions. These recent studies provide some evidence of a statistically significant inverse correlation between T1D and UVB irradiance.

The idea that UVR may in some way have a role in the development of the disease seems well supported in the epidemiological literature; however the mechanisms remain unknown. It is unclear whether UVR exposure impacts T1D risk through its subsequent production of vitamin D or whether other immunoregulatory molecules produced in skin following exposure to UVR are responsible. Equally, it may be that the combination of exposure to UVR and vitamin D sufficiency is necessary to reduce risk.¹⁵⁸

25(OH)D levels and T1D

As both dietary vitamin D and exposure to UVR increase blood 25(OH)D levels, 25(OH)D levels are not only an indication of

vitamin D status but also recent UV exposure. For this reason, 25(OH)D levels are addressed here in this perspective. The modulating effects of high 25(OH)D levels during the antenatal period on T1D risk maybe specific to the third trimester. Serum 25(OH)D levels of newborn babies taken at the time of birth were not associated with later risk of T1D in two studies.^{140,159} In a recent study, Sorenson and colleagues (2016) measured the serum 25(OH)D levels of mothers at multiple time points throughout pregnancy and subsequent risk of T1D in their offspring up to the age of 15 years.¹⁰ The results were consistent with findings of two previous studies in the field, indicating a significant association between low 25(OH)D levels during late pregnancy and increased T1D risk¹¹ and no association with levels in early pregnancy.¹⁶⁰ It is not known whether the apparent reduction in risk with high levels of 25(OH)D during the third trimester is associated with the large transfer of 25(OH)D through the placenta that occurs during this time¹⁶¹ or some other unknown biological mechanism unique to the third trimester.

In the absence of a single study that has quantified maternal supplement use and measured 25(OH)D levels at multiple time points throughout pregnancy, it is not possible to uncover why there is an apparent association between reduced risk of T1D and high levels of 25(OH)D during the third trimester but not with maternal intake of vitamin D supplements during pregnancy. It is possible that higher doses of supplementation than are currently recommended may be required to achieve a protective effect in offspring, or that benefits of high levels of 25(OH)D in the third trimester are due to factors other than vitamin D *per se*, such as exposure to UVR.

Studies that measure 25(OH)D levels in addition to both dietary intake of vitamin D and sun exposure are required to determine whether any reduction in risk is the result of the actions of vitamin D specifically, or exposure to UVR more generally.¹⁶² In a related autoimmune disease, multiple sclerosis (MS), UV irradiation inhibits development of symptoms of EAE with only a small, transient rise in 25(OH)D levels. Human studies also show that higher exposure to UVR decreased first demyelinating event risk (a common precursor to MS) independently of vitamin D, suggesting that supplementation alone may be insufficient to reduce the risk of MS.¹⁶³

Mechanisms of potential control of T1D by sun exposure and vitamin D

To study the immunoregulatory mechanisms of UVR exposure, it is preferable to irradiate an intact epidermal surface. This is usually the skin of a shaved experimental rodent although changes within human skin biopsies have also contributed to our understanding. Skin dendritic cells are affected both directly and indirectly by UVR exposure.^{164,165} It is necessary that they reach the draining lymph nodes where they can present antigens in such a way as to stimulate reduced, less proliferative responses. UVR-exposed dendritic cells (possibly with UVR-induced pyrimidine dimers) secrete down-regulatory



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