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The impact of vitamin D pathway genetic variation and circulating 25-hydroxyvitamin D on cancer outcome: systematic review and meta-analysis

P G Vaughan-Shaw¹, F O'Sullivan², S M Farrington¹, E Theodoratou^{1,3}, H Campbell^{1,3}, M G Dunlop¹ and L Zgaga^{*,2}

¹MRC Human Genetics Unit, Institute of Genetics and Molecular Medicine, University of Edinburgh, Edinburgh EH42XU, UK; ²Department of Public Health and Primary Care, Trinity College Dublin, Dublin 24, Republic of Ireland and ³Centre for Population Health Sciences, University of Edinburgh, Edinburgh EH164UX, UK

Background: Vitamin D has been linked with improved cancer outcome. This systematic review and meta-analysis investigates the relationship between cancer outcomes and both vitamin D-related genetic variation and circulating 25-hydroxyvitamin D (25OHD) concentration.

Methods: A systematic review and meta-analysis of papers until November 2016 on PubMed, EMBASE and Web of Science pertaining to association between circulating vitamin D level, functionally relevant vitamin D receptor genetic variants and variants within vitamin D pathway genes and cancer survival or disease progression was performed.

Results: A total of 44 165 cases from 64 studies were included in meta-analyses. Higher 25OHD was associated with better overall survival (hazard ratio (HR = 0.74, 95% CI: 0.66-0.82) and progression-free survival (HR = 0.84, 95% CI: 0.77-0.91). The rs1544410 (Bsml) variant was associated with overall survival (HR = 1.40, 95% CI: 1.05-1.75) and rs7975232 (Apal) with progression-free survival (HR = 1.29, 95% CI: 1.02-1.56). The rs2228570 (Fokl) variant was associated with overall survival in lung cancer patients (HR = 1.29, 95% CI: 1.0-1.57), with a suggestive association across all cancers (HR = 1.26, 95% CI: 0.96-1.56).

Conclusions: Higher 25OHD concentration is associated with better cancer outcome, and the observed association of functional variants in vitamin D pathway genes with outcome supports a causal link. This analysis provides powerful background rationale to instigate clinical trials to investigate the potential beneficial effect of vitamin D in the context of stratification by genotype.

The importance of vitamin D for bone health is well established, but the role of vitamin D beyond the skeletal system has been under debate for decades (Theodoratou *et al*, 2014). In recent years, it has become apparent that the vitamin D receptor (VDR) is expressed in most cells, and that multiple tissues have the ability to convert the primary circulating form of vitamin D into the active form (Bouillon *et al*, 2013), implying that extra-skeletal effects of vitamin D are likely. While typically thought of as 'vitamin', it may be more appropriate to regard the primary circulating form, 25-hydroxyvitamin D (25OHD), as a pre-hormone and the primary active form, 1,25-dihydroxyvitamin D (1,25(OH)₂D), as a *hormone*. It has been previously recognised that mutations in genes involved in response to hormones, their metabolism or actions may affect the prognosis of disease and thus act as modifiers. Correspondingly, 1,25(OH)₂D binds to the VDR (a ligand-dependent

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© The Author(s) named above

^{*}Correspondence: Dr L Zgaga; E-mail: zgagal@tcd.ie

transcription factor) and polymorphisms in the VDR gene have been shown to modify the activity of this VitD-VDR complex (Anderson et al, 2003): for example, rs11568820 is situated in the VDR promoter region and can influence transcriptional activity (Yamamoto et al, 1999), while rs2228570 affects the translational start site (Uitterlinden et al, 2004). Therefore, it is hypothesised that not only vitamin D status but also expression and structure of VDR determine molecular actions, and can potentially modify cancer risk and survival (Flugge et al, 2007; Li et al, 2007; Zgaga et al, 2014). The VitD-VDR complex has the ability to exert downstream biological effects; amongst others, it can regulate the expression of multiple target genes, including several with antitumour properties (Ramagopalan et al, 2010). Moreover, polymorphisms in the VDR gene have been linked to cancer risk, including prostate (Taylor et al, 1996), breast (Lowe et al, 2005), skin and bowel (Ingles et al, 2001; Xu et al, 2014; Serrano et al, 2016), and VDR expression has been linked to survival in prostate and breast cancer (Berger et al, 1991; Hendrickson et al, 2011; Ditsch et al, 2012). Unlike highly variable environmental exposures (sunlight, diet and supplements) or seasonally varying 25OHD levels (Kelly et al, 2015), genetic variants are constant, present since conception and cannot be modified by the disease; thereby removing reverse causation concerns.

Three aspects further strengthen the case for understanding the associations between vitamin D and cancer outcomes: first, cancer incidence and mortality are increasing (CRUK, 2015); second, vitamin D deficiency is common worldwide among otherwise healthy individuals (Holick, 2007; Zgaga *et al*, 2011), and particularly among cancer patients (Crew *et al*, 2009; Fakih *et al*, 2009; Shanafelt *et al*, 2011; Vrieling *et al*, 2011; Zgaga *et al*, 2014), and third, vitamin D deficiency is a modifiable risk factor; based on the studies that report an association between vitamin D deficiency and poorer cancers outcomes. Unsurprisingly, it has been proposed that vitamin D may have potential value as an adjuvant chemotherapeutic agent, particularly since vitamin D supplements are cheap, safe and readily available (Newton-Bishop *et al*, 2009, 2015; Drake *et al*, 2010; Hatse *et al*, 2012; Zgaga *et al*, 2014).

Here we present a systematic review and meta-analysis examining the role of vitamin D on cancer progression and survival. We conducted a comprehensive evaluation of the literature that examines the associations between cancer outcomes and genetic factors involved in the vitamin D pathway, in addition to circulating 25OHD concentration. Focus on vitamin D-related genetic variation allowed us to partially mitigate against potential confounding or reverse causation, biases that typically limit implications of findings from observational vitamin D studies.

MATERIALS AND METHODS

Literature search. We performed a systematic literature review and meta-analysis following PRISMA guidelines (Moher et al, 2009). The electronic databases PubMed (NCBI, 2015), EMBASE (EMBASE, 2015), and Web of Science (JISC, 2015) were searched up to week 3, November 2015. We searched for studies that examined the association between cancer outcomes and (i) measured vitamin D levels and (ii) genetic factors known to affect vitamin D metabolism or pathways. A list of search terms was compiled using a number of core papers in the field. For cancer outcomes, we included a combination of terms: cancer, neoplasm, malignant, malignancy with survival, outcome, prognosis, mortality, death, recurrence. For vitamin D levels, we included terms: 25-hydroxyvitamin D, calcidiol and 25OHD; for vitamin D receptor, and for commonly studied variants, we searched for: vitamin D receptor, VDR, rs1544410, BsmI, rs10735810, rs2228570, FokI, rs7975232, ApaI, rs11568820, Cdx2, rs2282679, rs12785878, rs10741657 and rs6013897. Finally, we also included variation in genes related to vitamin D synthesis, transport or metabolism: 1- α -hydroxylase, *CYP27B1*, 25-hydro-xylase, *CYP2R1*, 24-hydroxylase, *CYP24A1*, vitamin D binding protein, 27-hydroxylase and *CYP27A1*. Genetic variants beyond those explicitly searched for were only included if previously shown to affect vitamin D metabolism. We considered all human research full text articles, with no restriction on language or article type. Bibliographies of retrieved papers and previous reviews were hand-searched to identify other relevant studies.

Selection criteria and selection of relevant studies. Study inclusion 'PICO' criteria were as follows: (i) participants: individuals of any age who received a diagnosis of cancer; (ii) intervention/Exposures: assessment of vitamin D status or genetic factors known to affect vitamin D concentration, metabolism or pathways; (iii) comparators: study reports a quantitative association between cancer outcome and either vitamin D status (e.g., concentration, quartiles, low/high levels) sampled at most 1 year prior to the diagnosis, or any germline genetic variation or gene expression in normal tissue; and (iv) Outcome: cancer-specific or all-cause mortality, or disease progression (e.g., disease-free survival, local recurrence or metastasis). Observational retrospective and prospective cohorts were included.

In relation to patients, exclusion criteria were: (i) pre-cancerous lesions, and (ii) mixed-cancer cohort without site-specific reporting; in relation to exposures: (iii) vitamin D intake and supplementation, (iv) acquired non-germline mutations or tumour gene expression, and (v) predicted vitamin D status; in relation to outcomes: (vi) prognostic markers such as Prostate Specific Antigen or Breslow thickness, (vii) population cancer mortality rates; in relation to study/publication type: (viii) ecological studies, and (ix) reviews, editorials, case reports, conference abstracts and nonclinical publications. If the same patient cohort was reported on more than once, we used the highest quality, largest sample size or most recent publication. Article titles and abstracts were screened for eligibility, independently by two authors (PVS and LZ or FOS). Disagreements were resolved by discussion and review of full text.

Data extraction. The data extraction was performed by a single investigator (PVS or FOS) using the predefined data fields and extraction was cross-checked by a second investigator in its

Box 1. Conversion of continuous HR and 95% CI estimate to per 10 ng ml^{-1} HR estimates.

To achieve this, we raised the continuous HR (or 'HR per 1 ng ml^{-1}) to the power of 10 to get [A], per 10 ng ml⁻¹ HR (e.g., continuous HR, 0.89, hence per 10 ng ml⁻¹ HR: 0.89^10 = 0.3118). In order to calculate the confidence intervals we first found the exp(s.e.(beta)) = [B]; the standard errors (s.e.) were calculated using the formula below. We then calculated 1.96*(x - xbar) = [C], where x was fixed at 10 ng ml⁻¹ and xbar was the median of all cohort means from the rest of the studies included in the meta-analysis. We found this to be 23 ng ml⁻¹, therefore 1.96*(10 – 23) = -25.872 = [C]. [B] was then raised to the power of [C], to get [D]. Finally, the HR per 10 ng ml⁻¹[A] was multiplied or divided by [D] in order to derive the upper and lower 95% confidence intervals. Therefore, the resulting HR was A and 95% CI: (A*D) to (A/D).

(Continuous HR)^10 = [A] = per 10 ng/ml HR $SE = \frac{\log(upper 95\% \text{ CI}) - \log(\text{HR})}{1.96}$

exp(s.e.(beta)) = [B] $1.96^{*}(x - xbar) = [C]$ $B^{*}C = [D]$ $A^{*}D = lower 95\%$ Cl A/D = upper 95% Cl. entirety (FOS or PVS). The data from eligible studies were extracted using a tailored data extraction form that included the following information: first author, publication year, location or ethnicity of patients, sample size, mean age, gender, cancer site (subtype/histology where relevant), cancer stage, any interventions (e.g., chemotherapy), vitamin D exposure studied and important meta-data (time of sampling, mean/median 25OHD values or range for categories being compared; SNP position, name and rs ID, genotypes compared and model: additive, recessive or dominant), covariates considered, details of outcomes studied,

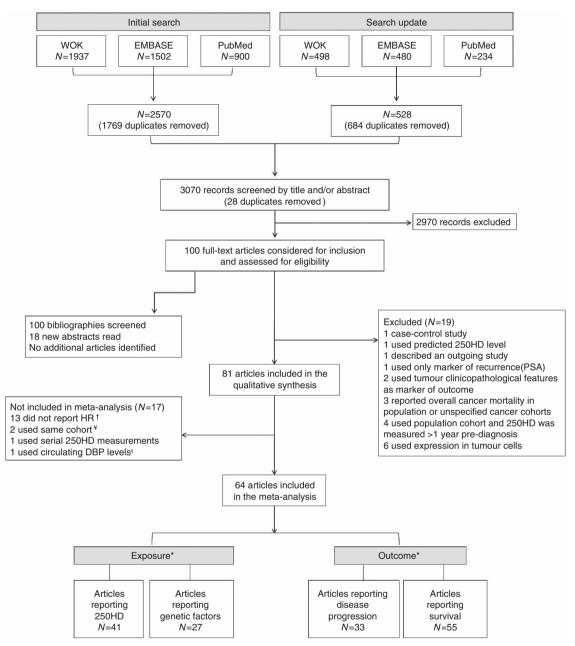


Figure 1. PRISMA Flowchart of the study selection process. Two studies used the same prostate cancer cohort but one reported on circulating 25OHD and the other on genetic variants, and so both were retained. (Holt *et al*, 2010, 2013) Three publications used the same initial cohort of lung cancer patients but two reported on different subpopulations of patients (according to disease stage) and so were retained, (Zhou *et al*, 2007; Heist *et al*, 2008), while a third reported on different exposures to the first two and so was also retained (Zhou *et al*, 2006). Finally, four studies reported on the same melanoma patient cohort (Newton-Bishop *et al*, 2009, 2015; Field *et al*, 2013; Davies *et al*, 2014) (one paper scored lower in NOS scoring was excluded (Field *et al*, 2013), while the remaining three, which reported different exposure or outcomes were retained. [§]Only a single study reported impact of circulating vitamin D-binding protein levels on outcome and so could not be included in the meta-analysis. *Includes only exposures and outcomes included in MA. Articles may report on multiple exposure-outcome pairs hence the sum of the pairs is greater than the number of articles included. For example, several papers studied the effect of more than one SNP for example, Zgaga *et al*, (Zgaga *et al*, 2015). However, where multiple estimates were extracted, no patient was included more than once for a certain exposure or outcome. † Study authors were contacted to provide HR, RR or OR when not reported; 13 did not respond. ‡ One study (Vrieling *et al*, 2011) used the same breast cancer cohort as a later, larger study (Vrieling *et al*, 2014) and as both had the same NOS score, the newer study was included. 25OHD: 25-hydroxyvitamin D; DBP: vitamin D binding protein; HR: hazard ratio; PSA: prostate specific antigen; WOK: Web of Knowledge.

Table 1. Characteristics of studies (N = 64) included in the meta-analysis

| | | | | | | | | Varia | ble | Outc | ome |
|--------------------------------|--------------------------------------|-------|----------------|---------|-------------------|--|-----|----------|---------|------------------|----------|
| First author, year | Cancer (subtype) | HR/OR | Sample Size | Site | Follow- up (m) | Events | NOS | 25OHD | Genetic | Progr- ession | Survival |
| Anic <i>et al</i> (2012) | Brain (glioma) | HR | 320 | USA | 28 | 248 cancer deaths | 5 | | ~ | | CS |
| Lim <i>et al</i> (2015) | Breast | HR | 491 | Korea | 86 | 32 recurrences; 22 cancer deaths | 8 | 🛩 BT | | DFS | CS |
| Lohmann <i>et al</i> (2015) | Breast | HR | 934 | Canada | 112 | Not given | 4 | 🛩 BT | | RFS | OS |
| Clark et al (2014) | Breast | HR | 82 | USA | >36 | 23 relapses or deaths | 5 | 🛩 BT | | RFS | |
| Vrieling et al (2014) | Breast | HR | 2177 | Germany | 64 | 206 cancer deaths, 241 recurrences or deaths | 7 | ⊭ 66% BT | | DFS | CS |
| Mishra et al (2013) | Breast | OR | 232 | USA | NA | Not given | 5 | | ~ | DFS | |
| Pande et al (2013) | Breast | HR | 1029 | USA | 114 | 266 recurrences or deaths | 6 | | - | DFS | |
| Perna et al (2013a) | Breast | HR | 498 | Germany | 60 | 48 cancer deaths | 7 | | ~ | | CS |
| Villasenor et al (2013) | Breast | HR | 585 | USA | 110 | 48 cancer deaths | 7 | M AT | | | CS |
| Hatse et al (2012) | Breast | HR | 1800 | Belgium | 56 | 118 relapses; 64 cancer deaths | 5 | ⊯ BT | | DFI | CS |
| Jacobs et al (2011) | Breast | OR | 512 | USA | 88 | Not given | 5 | 🖊 AT | | R | OS |
| Kim et al (2012) | Breast | HR | 310 | Korea | 23 | 33 metastases or deaths | 7 | ⊯ BT | | DFS | |
| Goodwin <i>et al</i> (2009) | Breast | HR | 512 | Canada | 139 | 116 recurrences; 106 deaths | 7 | 🛩 BT | | R | OS |
| Goode <i>et al</i> (2002) | Breast | HR | 721 | UK | NA | 200 deaths | 6 | | 1 | | OS |
| Lundin <i>et al</i> (1999) | Breast | RR | 111 | Sweden | 67 | 44 deaths | 4 | | 1 | | OS |
| Tretli <i>et al</i> (2012) | Breast, colon, lung, and lymphoma | HR | 658 | Norway | >60 | 343 cancer deaths | 7 | 🛩 BT | | | CS |
| Wesa et al (2015) | Colorectal | HR | 250 | USA | NA | 153 deaths | 5 | 🛩 BT | | | OS |
| Zgaga et al (2014) | Colorectal | HR | 1598 | UK | 107 | 363 cancer deaths | 8 | 🛩 AT | ~ | | CS |
| Perna <i>et al</i> (2013b) | Colorectal | HR | 1397 | Germany | 60 | 336 cancer deaths | 6 | | 1 | | CS |
| Szkandera <i>et al</i> (2013) | Colorectal | HR | 264 | Austria | 53 | 45 recurrences | 5 | | 1 | R | |
| Fedirko et al (2012) | Colorectal | HR | 1202 | Europe | 73 | 444 cancer deaths | 8 | 🛩 BD | 1 | | CS |
| Ng et al (2011) | Colorectal | HR | 515 | USA | 61 | 440 progression; 475 deaths | 5 | 🛩 BT | | TTP | OS |
| Mezawa et al (2010) | Colorectal | HR | 257 | Japan | 32 | 30 cancer deaths; recurrences not given | 5 | ► NS | | DFS | CS |
| Ng et al (2008) | Colorectal | HR | 304 | USA | 78 | 96 cancer deaths | 7 | 🛩 BD | | | CS |
| Ren <i>et al</i> (2012) | Gastric | HR | 197 | China | >60 | 106 deaths | 5 | 🛩 BT | | | OS |
| Lee et al (2014) | Haematological (AML) | HR | 97 | USA | 16 | 55 relapses; 51 deaths | 4 | 🛩 BT | | R | OS |
| Shanafelt <i>et al</i> (2011) | Haematological (CLL) | HR | 543 | USA | 118 | 201 progression; 96 deaths | 8 | MNS NS | | TTT | OS |
| Aref <i>et al</i> (2013) | Haematological (CLL, NHL) | HR | 195 | Egypt | 60 | 118 deaths | 5 | 🛩 BT | | | OS |
| Drake et al (2010) | Haematological (DLBCL) | HR | 983 | USA | 35 | 404 events; 168 cancer deaths | 6 | 🛩 66% BT | | EFF | CS |
| Pardanani <i>et al</i> (2011) | Haematological (PMF, MDS) | HR | 321 | USA | 34 | 36 progression; 171 deaths | 4 | ⊯ BT | | LFS | OS |
| Bittenbring et al (2014) | Haemotological (BCL) | HR | 359 | Germany | 49 | Not given | 4 | 🛩 AT | | EFF | OS |
| Kelly et al (2015) | Haemotological (FL) | HR | 423 | USA | 65 | 193 progression; 58 deaths | 5 | 🛩 BT | | PFS | OS |
| Azad <i>et al</i> (2013) | Head and neck | HR | 522 | Canada | >53 | 214 deaths | 8 | | ~ | | OS |
| Zeljic <i>et al</i> (2012) | Head and neck | OR | 110 | Serbia | 28–100 | Not given | 5 | | ~ | | CS |
| Meyer et al (2011) | Head and neck | HR | 540 | Canada | 96 | 119 recurrences; 223 deaths | 8 | ⊯ BT | | R | OS |
| Gugatschka <i>et al</i> (2011) | Head and neck (SCC) | RR | 88 | Austria | NA | 31 progression; 29 deaths | 4 | ₩ BT | | DFS | OS |
| Hama <i>et al</i> (2011) | Head and neck (SCC) | HR | 204 | Japan | 34 | 103 progression or deaths | 6 | | ~ | DFS | |
| Finkelmeier et al (2014) | Liver (HCC) | HR | 200 | Germany | 11 | 60 deaths | 6 | 🛩 BT | | | OS |
| Zhou <i>et al</i> (2007) | Lung | HR | 447 | USA | 72 | 126 cancer deaths | 7 | 🛩 BT | | | CS |
| Liu et al (2011) | Lung (AC, SCC) | HR | 568 | China | 19 | 311 deaths | 6 | 🛩 NS | 1 | | OS |
| Heist <i>et al</i> (2008) | Lung (AC, SCC) | HR | 294 | USA | 42 | 233 deaths | 6 | 🛩 NS | ~ | | OS |

Table 1. (Continued)

| | | | | | | | 1 | Varia | ble | Outo | ome |
|------------------------------|---------------------|-------|----------------|---------------|-------------------|---|-----|----------|---------|------------------|---------|
| First author, year | Cancer (subtype) | HR/OR | Sample Size | Site | Follow- up (m) | Events | NOS | 25OHD | Genetic | Progr- ession | Surviva |
| Zhou <i>et al</i> (2006) | Lung (AC, SCC) | HR | 373 | USA | 71 | 186 deaths | 7 | | ~ | | OS |
| Xiong et al (2013) | Lung (NSCC) | HR | 755 | China | NA | Not given | 4 | | ~ | PFS | OS |
| Newton-Bishop et al (2015) | Melanoma | HR | 2182 | UK | NA | Not given | 6 | 🛩 NS | | | CS |
| Davies et al (2014) | Melanoma | HR | 3137 | Various | 96 | 653 deaths | 7 | | 4 | | OS |
| Orlow et al (2014) | Melanoma | HR | 3566 | World wide | 91 | 254 cancer deaths | 7 | | ~ | | CS |
| Newton-Bishop et al (2009) | Melanoma | HR | 872 | UK | 56 | 173 relapses | 5 | 🛩 NS | | DFS | |
| Halsall et al (2004) | Melanoma | HR | 171 | UK | 75 | 18 metastases | 4 | | ~ | М | |
| Webb <i>et al</i> (2015) | Ovarian | HR | 670 | Australia | >60 | 491 progression; 435 deaths | 7 | 🛩 BT | | PFS | OS |
| Tamez <i>et al</i> (2009) | Ovarian | HR | 101 | Japan | 85 | 28 cancer deaths; total deaths not given | 7 | | ~ | | OS |
| Van Loon <i>et al</i> (2014) | Pancreatic | HR | 256 | Europe | 35 | progression not given; 254 deaths | 4 | 🛩 BT | | PFS | OS |
| Cho <i>et al</i> (2013) | Pancreatic | HR | 178 | USA | 33 | 82 deaths | 5 | 🛩 BT | | | OS |
| Gupta et al (2015) | Prostate | HR | 125 | USA | 31 | 49 deaths | 7 | 🛩 BT | | | OS |
| Trummer et al (2015) | Prostate | HR | 702 | Austria | 73–91 | 93 metastases; 123 deaths | 6 | | ~ | М | OS |
| Holt et al (2013) | Prostate | HR | 1476 | USA | 130 | 325 progression; 95 cancer deaths | 7 | M NS | | Ρ | CS |
| Pao <i>et al</i> (2013) | Prostate | HR | 601 | Taiwan | 60–120 | 415 progression; 101 cancer deaths | 8 | | ~ | Ρ | CS |
| Fang <i>et al</i> (2011) | Prostate | HR | 1822 | USA | 120 | 166 cancer deaths | 8 | 🛩 BD | | | CS |
| Holt <i>et al</i> (2010) | Prostate | HR | 1294 | USA | 102 | 139 recurrences; 57 cancer deaths | 8 | | ~ | R | CS |
| Penney et al (2010) | Prostate | OR | 1292 | USA | >60 | Not given | 5 | | - | | OS |
| Tretli <i>et al</i> (2009) | Prostate | HR | 160 | Norway | 44 | 52 cancer deaths | 6 | 🛩 77% BT | | | CS |
| Williams et al (2004) | Prostate | HR | 728 | USA | 60–120 | Not given | 7 | | ~ | DFS | |
| | Renal | HR | 630 | Europe | 30 | 152 cancer deaths | 8 | 🛩 BT | | | CS |
| Obara et al (2007) | Renal (RCC) | RR | 135 | Japan | >60 | Not given | 5 | | ~ | | CS |
| Samimi <i>et al</i> (2014) | Skin (Merkel cell) | HR | 89 | France | NA | 33 metastases; 19 deaths | 6 | ► NS | | М | CS |

250HD = assayed before diagnosis; BT 250HD = assayed before treatment; CML = chronic myeloid leukaemia; CS = cancer-specific survival; DFI = disease-free interval; DFS = disease-free survival; DLBCL diffuse large B-cell lymphoma; EFF = event-free survival; FL = follicular lymphoma; HCC = Hepatocellular carcinoma; LFS = leaukaemia-free survival; m months; M = metastasis; MDS = myelodysplastic syndrome; NA = not available; NHL = Non-Hodgkins lymphoma; NOS = Newcastle-Ottawa score; NS = Timing of 250HD not specified/variable; NSCC = non-small-cell lung carcinoma; OS = overall-survival; P = progression not otherwise specified; PFS = progression-free survival; PMF = primary myelofibrosis; R = recurrence or relapse not otherwise specified; RCC = renal cell carcinoma; TTP = time to progression; TTT = time to treatment.

and follow-up time. Finally, hazard ratios (HR) and 95% confidence intervals (95% CIs) adjusted for the maximum number of confounding variables were extracted. We preferentially focused on cancer-specific mortality, but if these data were not available, all-cause mortality was used instead. Relative risk estimates (RR) or adjusted odds ratios (OR) were extracted where HR were not given and used in meta-analysis (Symons and Moore, 2002). Study authors were contacted to provide additional information where needed.

Quality assessment. The methodological quality of all studies included in the systematic review was performed using the Newcastle-Ottawa Quality Assessment Scale (NOS; Wells *et al*, 2000). Two investigators (PVS and FOS) applied predefined NOS criteria to each study to generate summary quality judgement. The risk of bias was considered 'low' for studies with score of 7 or 8; 'unclear' for score of 5 or 6, and 'high' for score of 4 or lower.

Exposure assessment. The association between circulating 25OHD and outcomes was summarised in meta-analyses by comparing the risk in the highest to the lowest reported category. The majority of studies used vitamin D categories such as quartiles

or tertiles. To enable inclusion of studies that used 25OHD as a continuous variable, we sought to transform the 'continuous HR' into a 'HR per 10 ng ml^{-1} ' (Box 1).

Genetic factors. For SNPs, the rs number naming convention was typically used in the paper and some recoding was needed to ensure that uniform reference system was followed. For example, where a restriction fragment length polymorphism was referenced, the mutation and risk allele were recoded (e.g., *FokI f* allele was converted to the rs2228570 *T* allele). The genome browser ENSEMBL (80 GRCh38.p2) was used to determine if alias names existed (e.g., *FokI*, rs10735810 and rs2228570 are the same variant). HR values were inverted where needed, so that the same allele acted as the reference. Where additive models were used, the HR value for comparison between two homozygotes.

Statistical analysis. We conducted meta-analyses for a range of exposure-outcome pairs by cancer site and across all sites. A meta-analysis was performed if at least two studies considered the same exposure-outcome pair. The same study may have been included multiple times in different meta-analyses if it reported on multiple

subpopulations, outcomes, and/or exposures. The extracted HRs and 95% CIs were used to calculate the pooled HR estimates. The standard errors (s.e.) were used to calculate weighting for each study. The DerSimonian and Laird random-effects model was used to calculate pooled HR because of the *a priori* expected heterogeneity between studies, due to differences among populations and methodological dissimilarities between studies; most notably, different definition of 25OHD categories. All analyses were performed in R (R Core Team, 2013), and the R-package 'metafor' was used for meta-analyses (Viechtbauer and Cheung, 2010). *P*-value <0.05 was considered statistically significant.

In order to assess the impact of study quality on results, metaanalyses were rerun (i) after exclusion of studies at high risk of bias, (ii) limited to studies at low risk of bias only, (iii) limited to studies that looked at cancer-specific mortality, (iv) excluding studies that used 25OHD as a continuous variable, and (v) excluding studies that reported RR or OR. The I^2 statistic was calculated to quantify the degree of heterogeneity between studies and assess impact on the meta-analysis (Higgins *et al*, 2003). To further explore this issues arising due to the striking differences in 25OHD category definition, we conducted a stratified analysis (Cochrane, 2011) according to: (vi) the difference in mean/median 25OHD between

| | | | | | | Vari | able | Outco | ne | | |
|--|---|-------|-------------------|--|-----|-------|---------|-------------|----------|--|-----------------------------------|
| First author, year | Cancer (subtype) | Size | Follow- up (m) | Events | NOS | 25OHD | Genetic | Progression | Survival | Author conclusion | Reason exclude |
| Obermannova <i>et al</i> (2015) | Colorectal | 84 | 24 | Not given | 4 | ~ | | PFS | OS | Consistently low 25OHD (always <16 ng/ml) associated with worse PFS and OS | Serial 250HD |
| Turner <i>et al</i> (2013) | Lung (NSCC) | 142 | 52 | Not given | 7 | | | | CS | Low serum DBP levels predicted lung cancer-specific death ($P = 0.04$) | Only paper reporting DBP |
| Turna <i>et al</i> (2012) | Lung(NSCC) | 62 | NA | Not given | 5 | | لمع | | OS | Haplotype analysis revealed rs731236 (Taql)—rs2228570 (Fokl) TTFf/TtFf haplotype associated with reduced OS ($P=0.04$) | No individual SNP HR |
| Bade <i>et al</i> (2014) | Melanoma | 324 | NA | Not given | 6 | ~ | | | OS | Increased 25OHD (Q4 v Q1) associated with increased OS 195 months v 80 months ($P = 0.049$) | No HR |
| Der et al (2014) | Prostate | 16535 | 60 | 4613 deaths | 5 | ~ | | | OS | Vitamin D deficiency significantly associated with reduced survival (<0.001) | No HR |
| Dickinson <i>et al</i> (2010) | Haematological (CML) | 228 | NA | 55 relapses; 84 deaths | 5 | | ~ | R | OS | No data provided on impact of VDR variants | No HR |
| Furuya <i>et al</i> (1999) | Prostate | 66 | NA | Not given | З | | ~ | PFS | | Taql TT genotype associated with shorter PFS ($P = 0.07$) | No HR |
| Hansson <i>et al</i> (2014) | Haematological (AML, ALL, CML, MDS) | 123 | 96 | 29 relapses; 31 deaths | 6 | 14 | | R | OS | 25OHD < 20 ng/ml associated with reduced OS ($P=0.01$) and increased relapse ($P=0.03$) | No HR |
| Kim <i>et al</i> (2012) | Haematological | 100 | 105 | 12 relapses; 4 deaths | 4 | | 1 | EFS | OS | VDR rs2228570 Fokl genotype did not impact survival in paediatric ALL | No HR |
| Numberg et al (2009) | Melanoma | 205 | NA | 118 metastases | 4 | ~ | | М | | 25OHD > 20 ng/l associated with increased time to distant metastatic disease ($P = 0.64$) | No HR |
| Peiris <i>et al</i> (2013) | Bladder | 4126 | NA | 2025 deaths | 6 | ~ | | | OS | 25OHD < 20 ng/ml associated with reduced OS (X2 = 10.44; P = 0.001) | No HR |
| Silvagno <i>et al</i> (2010) | Ovarian (Epithelial) | 26 | NA | Not given | 2 | | ~ | | OS | Increased platelet VDR expression (> 50 fMol) associated with increased OS (P=0.12) | No HR |
| Walentowicz- Sadlecka <i>et al</i> (2012) | Ovarian | 72 | 60 | 45 deaths | 6 | ~ | | | OS | 25OHD <10 ng/ml associated with reduced OS (P <0.04) | No HR |
| Yagmurdur et al (2009) | Breast | 56 | 60 | 5 recurrences | 3 | | | R | | rs1544410 (Bsml) genotype not associated with local recurrence or metastasis P>0.05 | No HR |
| Yiallourou <i>et al</i> (2014) | Breast | 87 | 60 | Not given | 3 | | ~ | PFS | OS | rs2228570 Fokl ff associated with reduced PFS 35 months vs >54 months (P =0.08) | No HR |
| Field et al (2013) | Melanoma | 795 | 56 | 137 cancer deaths | 4 | ~ | | | CS | 8 ng/ml incremental increase in 25OHD associated with improved DFS (P =0.02) and MSS (P =0.05) | Duplicate patient cohort |
| Vrieling <i>et al</i> (2011) | Breast | 1295 | 70 | 182 recurrence or metastases; 183 deaths | 7 | ~ | | DFS | OS | Low 25OHD significantly associated with worse DFS and OS | Duplicate patient cohort |

nucleotide polymorphism.

'high' and 'low' categories compared (below or $\ge 20 \text{ ng ml}^{-1}$), and (vii) the degree of deficiency in 'low' category (mean/median 250HD concentration below or $\ge 12.5 \text{ ng ml}^{-1}$). Publication and selection bias was investigated by checking for asymmetry in the funnel plots and running the Egger's regression test (Sterne and Egger, 2001).

RESULTS

A flowchart illustrating study selection is shown in Figure 1. After removal of duplicates, the search yielded 3070 potential articles. Irrelevant articles were eliminated after screening titles (N = 2708)

| Author | Comparision 0 | 12.5 | 25 | 37.5 | 0HD (ng ml ⁻¹) 50 | 62.5 | 75 | 87.5 | 10 |
|----------------------|--|--------------|------------------------------|--------------------|----------------------------------|-----------------------------|-----|------|----|
| Mezawa | Quartile 4 vs Quartile 1 | | | - | | | | | |
| Zgaga | Tertile 3 vs Tertile 1 | — — | | | - | | | | |
| Bittenbring | < >8 ng ml ⁻¹ | -88 | | | | - | | | |
| Webb | 20–30 <i>vs</i> < 10 ng ml ⁻¹ | • • | | | | | | | |
| Finkelmeier | >20 vs 10 ng ml ⁻¹ | | | | | _ | - | | |
| Liu | Quartile 4 vs Quartile 1 | | > | | | | | | |
| Zhou | Quartile 4 vs Quartile 1 | | | | | | | | |
| Holt | 20–59.9 vs < 12 ng ml ⁻¹ | • • | | | _ | - | | | |
| Heist | Quartile 4 vs Quartile 1 | | | | | | | | |
| Ng (2011) | Quartile 4 vs Quartile 1 | | | | | | | | |
| Vrieling | Tertile 3 vs Tertile 1 | | | | | | | | |
| Fedirko | Quintile 4 vs Quintile 1 | | | 1 | | | | | |
| ohmann | >20 and < 16 ng ml ⁻¹ | | | | | | | | |
| Tretli (2012)B | Quartile 4 vs Quartile 1 | | | 1 | | | | | |
| Tretli (2012)L | Quartile 4 vs Quartile 1 | | | | | | | | |
| Tretli (2012)Ly | Quartile 4 vs Quartile 1 | | | | | | | | |
| Meyer | Quartile 4 vs Quartile 1 | | | | _ | | | | |
| Samimi | <>20 ng ml ⁻¹ | | _ | | _ | | | | |
| Lim | <>20 ng ml ⁻¹ | | | | | | | | |
| | < >20 ng ml ⁻¹ | | _ | | | | | | |
| Ren | <>20 ng ml ⁻¹ | _ | | | | | | | |
| lacobs | < >20 ng ml ⁻¹ | | | | | | | | |
| Aref | <>20 ng ml ⁻¹ | - | | | | | | | |
| Cho Newton-Bishop | | | | | | | | | |
| 2015) | 20–60 vs <20 ng ml ⁻¹ | | | | | - | | | |
| Goodwin | >29 <i>vs</i> <20 ng ml ⁻¹ | | • | - | | | - | | |
| Kim | >30 <i>vs</i> <20 ng ml ⁻¹ | _ | · · | - | | | | | |
| Villasenor | >30 <i>vs</i> <20 ng ml ⁻¹ | | • | - | | | - | | |
| .ee | >32 vs <20 ng ml ⁻¹ | | - | - | | _ | | | _ |
| Tretli (2009) | >32 vs <20 ng ml ⁻¹ | | - | | -0- | _ | | | |
| Gupta | >32 vs <20 ng ml ⁻¹ | | | • | | | | | |
| Tretli (2012)C | Quartile 4 vs Quartile 1 | | | | | | | | |
| Ng (2008) | Quartile 4 vs Quartile 1 | | | | | | • | | |
| Drake | < >25 ng ml ⁻¹ | | - | | | | | | |
| Pardanani | < >25 ng ml ⁻¹ | | | | _ | _ | | | |
| Shanafelt | < >25 ng ml ⁻¹ | | | | | | - | | |
| Vesa | < >30 ng ml ⁻¹ | | | - | | | | | |
| Hatse | <>30 ng ml ⁻¹ | | | - | _ | | | _ | |
| Clark | Continuous (range NA) | | | | | | | | |
| Gugatschka | Continuous | | | 200 (| | | | | |
| Vewton-Bishop | Continuous | | | | | | | | |
| (2009) Van Loon | | | | | | | | | |
| | Continuous | NA | | | | | | | |
| Kelly | Tertile 3 vs Tertile 1 | NA | | | | | | | |
| =ang | Quartile 4 vs Quartile 1 | NA | | | | | | | |
| Muller | Quartile 4 vs Quartile 1 | | | | | | | | |
| | 0 | 12.5 | 25 | 37.5 25-hydoxyv | 50 itamin D level r | 62.5 Ig ml ⁻¹ | 7.5 | 87.5 | 10 |
| | | Catagory/col | Legen nort median (report | | | | | | |

Range of category values
 Cut off of category values (range not given)

250HD continuous

or abstracts (N = 262). One hundred full-texts were considered for inclusion and assessed for eligibility and 19 were excluded. Finally, 81 articles were kept for the systematic review and 64 of these were included in the meta-analysis. The main characteristics of included studies are summarised in Table 1 and Table 2.

Assessment of included studies. The risk of bias assessment revealed that 35 studies (43%) had a low risk of bias, 35 (43%) had an uncertain, and 11 (14%) had a high risk of bias. The risk of bias assessment summary per each domain is shown in Supplementary Figure S1 and individual study scores in Supplementary Figure S2. Sixty-four studies were included in the meta-analysis, with a total of 44165 patients. Most studies were conducted in the USA (N=24) and Europe; breast cancer was most commonly studied (N=15), followed by nine studies (each) on prostate cancer and colorectal cancer. In total, 157 HR estimates for a range of exposure-outcome pairs were included in meta-analyses: 77 estimates (from 41 studies) for association with 25OHD, and 80 estimates (from 27 studies) relating to genetic factors. Separate estimates were extracted for different patient subgroups (e.g., different type of haematological malignancy (Drake et al, 2010)), different exposures (e.g., multiple polymorphisms (Zgaga et al, 2014)), or different outcome (i.e., survival or disease progression (Lohmann et al, 2015)). No patients were included more than once in meta-analysis, as separate meta-analyses have been conducted for each exposure-outcome pair. Very large differences were observed in definition of vitamin D categories being compared. For example, the median 25OHD concentration was 18.26 ng ml^{-1} in the 'high' category in one study, (Zgaga et al, 2014) yet this was actually lower than the median $(19.7 \text{ ng ml}^{-1})$ in the 'low' category in another study (Hatse et al, 2012). The variety of vitamin D categories, cutoffs and means/medians used are presented in Figure 2 and Supplementary Figure S3.

Meta-analysis of 25OHD studies

Circulating vitamin D and survival. Forty-eight estimates from 38 studies were included in the meta-analysis of 25OHD and survival (17 studies (45%) examined cancer-specific mortality), comprising in total 24013 cancer patients. Twelve cancer types were represented: breast, haematological, head and neck, colorectal, lung, prostate, skin, pancreatic liver, gastric, kidney and ovarian cancers. Overall, a significantly reduced risk of death was observed when comparing those with high to those with low vitamin D levels; meta-analysis HR = 0.74, 95% CI = 0.66 to 0.82 (Figure 3). The same significant trend was also observed in subgroup meta-analysis for breast (HR = 0.75, 95% CI = 0.56–0.95), haematological (HR = 0.59, 95% CI = 0.42–0.77) and colorectal cancers (HR = 0.75, 95% CI = 0.60–0.90). There was also a non-significant

trend towards better survival with increased 25OHD observed in the subgroup analysis for prostate, skin, head and neck cancers. Virtually no change in direction or significance in overall effect was observed in sensitivity analyses when excluding studies at high risk of bias (HR = 0.73, 95% CI = 0.65-0.80), focusing on the studies at low risk of bias only (HR = 0.72, 95% CI = 0.63-0.81), excluding studies that used continuous 25OHD (HR = 0.73, 95% CI = 0.65-0.80), limited to cancer-specific mortality studies only (HR = 0.75, 95% CI = 0.65-0.84), or to studies that strictly reported HR (HR = 0.74, 95% CI = 0.66-0.82). The same was true after selection of studies where the difference in mean/median between high and low categories being compared was over or below 20 ng ml⁻¹; (HR = 0.70, 95% CI = 0.60-0.81, and HR = 0.71, 95% CI = 0.55-0.87, respectively), or when stratifying by the lower category mean/ median below or greater than 12.5 ng ml⁻¹ (HR = 0.76, 95% CI = 0.64 - 0.88, and HR = 0.61, 95% CI = 0.47 - 0.75, respectively) (for sensitivity analysis please see supplementary material).

The relationship between circulating vitamin D level and disease progression. Twenty-three studies investigated the association between circulating 25OHD and disease progression; from these studies 29 estimates were included in our meta-analysis comprising in total 14 307 patients with breast, haematological, head and neck, colorectal, prostate, skin, pancreatic, or ovarian cancer. Higher circulating vitamin D was associated with a significant reduction in disease progression for all cancers combined (HR = 0.84, 95% CI = 0.77 - 0.91; Figure 4); this was also observed in subgroup metaanalysis of breast (HR = 0.66, 95% CI = 0.45-0.88), haematological (HR = 0.75, 95% CI = 0.61 - 0.88) and skin cancer (HR = 0.77, 95% CI = 0.61 - 0.88)CI = 0.58-0.97). Findings remain fundamentally unchanged after exclusion of studies at high risk of bias (HR = 0.82, 95% CI = 0.74-0.90), limited to studies at low risk of bias only (HR = 0.80, 95%CI = 0.70 - 0.90) or excluding studies that used continuous 25OHD (HR = 0.81, 95% CI = 0.73-0.90), or limited to studies that strictly reported HR (HR = 0.84, 95% CI = 0.77-0.91). The same was true after selection of studies where the difference in mean/median between high and low categories being compared was over or below 20 ng ml^{-1} ; (HR = 0.81, 95% CI = 0.72-0.90, and HR = 0.75, 95% CI = 0.55-0.95, respectively), or when stratifying by the lower category mean/median below or greater than 12.5 ng ml^{-1} (HR = 0.84, 95% CI = 0.71-0.97, and HR = 0.77, 95% CI = 0.62-0.92, respectively) (for sensitivity analysis please see supplementary material).

Vitamin-D-related genetic variation

VDR and other vitamin D pathway SNPs and survival. Twentyone studies investigated the association between vitamin-D-related genetic variation and survival; 10 (48%) examined cancer-specific

Figure 2. Large variation in definition of vitamin D categories in studies included in systematic review. Where not given in the paper, median 25OHD concentration for categories compared was requested from study authors and if not subsequently available was approximated. Approximation of the median for each category was performed using the cohort and/or category range where available. For categories defined by numerical cutoffs of 25OHD, the median for the lower category was approximated as the lowest reported 25OHD value (or 0 if category range not given) added to the midpoint of the category upper cutoff minus the lowest reported 25OHD value. For example, Bittenbring et al (Bittenbring et al, 2014) reported outcome according to 25OHD < >8 ng ml⁻¹ groups and reported a study cohort range of 4–61.9. The median of the lower category ($< 8 \text{ ng ml}^{-1}$) was approximated as the lowest value in the range plus the midpoint of the category that is, 4 + ((8 - 4)/2) = 6. The upper category median was approximated as the category cutoff (that is, the lowest value in that category) added to the midpoint of the lower category. for example, in the Bittenbring et al, paper the median of the upper category was approximated as 8 + ((8 - 4)/2) = 10. Where the compared categories were tertiles, quartiles or quintiles, the median of the lower category and upper categories was the midpoint of the difference between upper cutoff of the lower category compared and the lower cut-off of the higher category compared divided by the number of groups between two categories compared, either subtracted from the upper cutoff of the lower category or added to the lower cutoff of the higher category, respectively. For example, Bade et al, (Bade et al, 2014) grouped patients by quartile of 25OHD and report a cohort range of 4–59.6 ng ml⁻ Q1 is given as 250 HD < 9.86 ng ml⁻¹ and Q4 > 24.4 ng ml⁻¹. Therefore, the medians of Q1 and Q4 were approximated as follows: Q1(median) = 9.86 - (((24.4 - 9.86)/2)/2) = 6.225 and Q4(median) = 24.4 + (((24.4 - 9.86)/2)/2) = 28. Insufficient data were reported in three studies to allow graphical illustration of categories or approximation of median. NA = data not reported; For Tretli et al, study: B = breast; C = colon; L = lung; Ly = lymphoma.

| Author and Year | Population | N.O. score | N | Exposure (ng ml-1) | | Weight | HR (95% CI) |
|--|-------------------|------------|-------------|---|--|--------------|--|
| Breast | | | | | | | · · · · · · |
| Hatse, 2012 | Belgium | 5 | 1800 | >30 <i>vs</i> <30 | | 1.6% | 0.49 (0.27 , 0.89) |
| Villasenor, 2013 | USA | 7 | 585 | >30 vs <30 | | 0.9% | 1.21 (0.52, 2.80) |
| Goodwin, 2009 | Canada | 7 | 512 | >29 <i>vs</i> <20 | —— ———————————————————————————————— | 2.2% | 0.63 (0.38 , 1.04) |
| Tretli, 2012 | Norway | 8 | 251 | Quartile 4 vs Quartile 1 | ⊢■ −−−1 | 1.3% | 0.42 (0.21 , 0.82) |
| Jacobs, 2011 | USA | 7 | 512 | >30 vs <10 | ∎ | 2.5% | 0.88 (0.56, 1.39) |
| Lohmann, 2015 | Canada | 5 | 934 | >20 <i>vs</i> <16 | ⊢ | 1.4% | 1.07 (0.57, 2.02) |
| Lim, 2015 | Korea | 8 | 491 | >20 <i>vs</i> <20 | ⊢ ∎———————————————————————————————————— | 0.6% | 0.46 (0.17, 1.22) |
| Vrieling, 2014 | Germany | 7 | 2177 | Tertile 3 vs tertile 1 | H | 3.3% | 0.79 (0.53, 1.16) |
| Subtotal (breast) | | | | | \diamond | | 0.75 (0.56 , 0.95) |
| Haemotological | | | 07 | | | 4.40/ | 0.04 (0.40, 0.70) |
| Lee, 2014 | USA | 4 | 97 | >32 vs <20 | | 1.1% | 0.34 (0.16, 0.72) |
| Shanafelt, 2011 | USA | 8 | 543 | >25 <i>vs</i> <25 >25 <i>vs</i> <25 | | 3.0% | 0.68 (0.45, 1.03) |
| Drake (PostFL), 2010 | USA USA | 7 7 | 109 285 | >25 vs <25 >25 vs <25 | | 0.3% | 0.36 (0.08, 1.72) |
| Drake (FL), 2010 | USA | 7 | 71 | >25 vs <25 | | 0.7% | 1.11 (0.29, 4.35) 0.74 (0.29, 1.89) |
| Drake (MCL), 2010 Drake (TCL), 2010 | USA | 7 | 70 | >25 vs <25 | | 0.9% | 0.44 (0.19, 1.01) |
| Drake (DLBCL), 2010 | USA | 7 | 370 | >25 vs <25 | | 2.3% | 0.46 (0.28 , 0.75) |
| Drake (other), 2010 | USA | 7 | 78 | >25 vs <25 | | 0.5% | 0.58 (0.19, 1.72) |
| Tretli, 2012 | Norway | 8 | 145 | Quartile 4 vs Quartile 1 | | 1.0% | 0.39 (0.18, 0.83) |
| Aref (NHL), 2013 | Egypt | 6 | 120 | >20 vs <20 | H H | 0.3% | 0.19 (0.04 , 0.86) |
| Aref (CLL), 2013 | Egypt | 6 | 75 | >20 vs <20 | ⊨■ →1 | 0.4% | 0.24 (0.07 , 0.85) |
| Pardanani (MDS), 2011 | USA | 5 | 74 | >25 vs <25 | | 1.2% | 0.71 (0.37, 1.43) |
| Pardanani (PMF), 2011 | USA | 5 | 247 | >25 vs <25 | +=1 | 3.0% | 0.83 (0.63 , 1.45) |
| Kelly (Swog cohort), 20 | | 7 | 183 | Tertile 3 vs Tertile 1 | | 0.4% | 0.35 (0.10 , 1.16) |
| Kelly (Lysa cohort), 201 | | 7 | 240 | Tertile 3 vs Tertile 1 | -■ | 0.3% | 0.19 (0.04 , 0.90) |
| Bittenbring, 2014 | Germany | 5 | 359 | >8 <i>vs</i> <8 | H | 2.4% | 0.56 (0.30, 0.90) |
| Subtotal,(Haemotologic | | Ū | 000 | | \diamond | 2.170 | 0.59 (0.42 , 0.77) |
| Colorectal | | | | | | | |
| Wesa,2015 | USA | 5 | 250 | >30 <i>vs</i> <20 | ⊢∎ —(| 2.4% | 0.61 (0.38,0.98) |
| Fedirko et al, 2012 | Europe | 8 | 1202 | Quartile 5 vs Quartile 1 | HEH | 4.8% | 0.69 (0.50 , 0.93) |
| Tretli, 2012 | Norway | 8 | 52 | Quartile 4 vs Quartile 1 | H=H | 0.2% | 0.20 (0.04 , 1.10) |
| Ng, 2011 | USA | 6 | 515 | Quartile 4 vs Quartile 1 | H ai -I | 5.5% | 0.94 (0.72 , 1.23) |
| Zgaga,2014 | UK | 8 | 1598 | Tertile 3 vs Tertile 1 | HEH | 5.2% | 0.68 (0.50 , 0.90) |
| Ng, 2008 | USA | 7 | 304 | Quartile 4 vs Quartile 1 | | 1.3% | 0.61 (0.31 , 1.19) |
| Mezawa,2010 Subtotal (Colorectal) | Japan | 5 | 257 | Per 10 ng ml ⁻¹ increase | \diamond | 0.4% | 0.82 (0.23 , 2.94) 0.75 (0.60 , 0.90) |
| Lung Tretli,2012 | Norway | 8 | 210 | Quartile 4 vs Quartile 1 | | 2.4% | 0.18 (0.11 , 0.29) |
| Liu,2011 | China | 6 | 568 | Quartile 4 vs Quartile 1 | · · · | ▶ 0.7% | 2.54 (1.01,6.41) |
| Zhou,2007 Heist,2008 | USA USA | 7 7 | 447 294 | Quartile 4 vs Quartile 1 Quartile 4 vs Quartile 1 | | 3.2% 3.5% | 0.74 (0.50 , 1.10) 1.08 (0.75 , 1.57) |
| Subtotal (Lung) | 00,1 | | 201 | quanto i vo quanto i | | 0.070 | 1.07 (0.17 , 1.98) |
| Head and Neck | | | 5.40 | | | 0.001 | 0.95 (0.57 1.98) |
| Meyer, 2011 Gugats chka, 2011 | Canada Austria | 8 4 | 540 88 | Quartile 4 vs Quartile 1 Per 10 ng ml ⁻¹ increase | | 3.0% 0.5% | 0.85 (0.57 , 1.28) 0.31 (0.10 , 0.97) |
| Subtotal (Head and Nec | | - | 00 | i ei io ig illi illoicase | $\langle \rangle$ | 0.070 | 0.79 (0.40 , 1.17) |
| Prostate | | _ | | | | | 0.02/0.07 0.00 |
| Holt, 2013 Gupta 2015 | USA USA | 7 7 | 1476 125 | >20 <i>vs</i> <12 >32 <i>vs</i> <20 | | 0.8% 1.2% | 0.83 (0.37, 2.00) 0.71 (0.31, 1.42) |
| Gupta, 2015 Tretli, 2009 | Norway | 6 | 160 | >32 vs <20 >32 vs <20 | | 0.5% | 0.09 (0.03, 0.27) |
| Fang, 2011 | USA | 8 8 | 1822 | Quartile 4 vs Quartile 1 | | 6.2% | 0.90 (0.72, 1.15) |
| Subtotal(Prostate) | | | | | | | 0.84 (0.63 , 1.06) |
| Pancreatic | USA | 5 | 178 | > 20, 1/2, < 20, | | 1.8% | 0.50 (0.29 , 0.87) |
| Cho, 2013 Van Loon, 2014 | USA | 5 4 | 256 | >20 <i>vs</i> <20 Per 10 ng ml ⁻¹ increase | | 10.7% | 1.00 (0.88, 1.14) |
| Subtotal (Pancreatic) | 00/1 | - | 200 | i ei io ig illi illoicase | $\langle \dot{\langle} \rangle$ | 10.7 /0 | 0.83 (0.36 , 1.29) |
| Skin | | | | | | | |
| Samimi, 2014 | France | 6 | 89 | >20 <i>vs</i> <20 | H= | 0.2% | 0.19 (0.03 , 1.30) |
| Newton-Bishop, 2015 Subtotal (Skin) | UK | 6 | 1557 | >8 <i>vs</i> <8 | | 2.6% | 0.66 (0.41 , 1.03) 0.64 (0.20 , 1.07) |
| | | | | | | | |
| Liver Finkelmeier, 2014 | Germany | 6 | 200 | >10 <i>vs</i> <10 | += | 1.8% | 0.50 (0.28 , 0.88) |
| Castria | | | | | | | |
| Gastric Ren, 2012 | China | 6 | 197 | >20 <i>vs</i> <20 | ⊢ ∎ | 2.8% | 0.59 (0.37 , 0.91) |
| Kidney Muller, 2015 | Czech Republic | 8 | 630 | Quartile 4 vs Quartile 1 | | 1.7% | 0.70 (0.39 , 1.24) |
| | | 0 | 030 | Quartile 4 VS QUARTILE | | 1.770 | 0.70 (0.08 , 1.24) |
| Ovarian Webb, 2015 | Australia | 8 | 670 | >20 <i>vs</i> <10 | He-I | 4.8% | 0.69 (0.51 , 0.93) |
| OVERALL CANCER | | | | | • | | 0.74 (0.66 , 0.82) |
| | | | | | | | |
| | | | | | 0.00 1.50 3.00 | 4.50 | |

Figure 3. Cancer survival and 25-hydroxyvitamin D concentration: meta-analysis of adjusted hazard ratios. HR are sorted by cancer site and the difference in median between 'high' and 'low' vitamin D categories compared. Acute myeloid leukaemia (AML), Chronic Lymphoid Leukaemia (CLL), and subtypes of non-Hodgkin's lymphoma (NHL) (large B-cell lymphoma (DLBCL), T-cell lymphoma (TCL), Follicular Lymphoma (FL) and mantle cell lymphoma (MCL)) Myelodysplastic syndrome (MDS) and primary myelofibrosis (PMF). *I*² = breast: 0, haematological: 0, colorectal: 0.91, prostate: 0.68, head and neck: 0, pancreatic: 0.66, lung: 0.93, skin: 0, overall cancer: 0.18. Approximated Median in studies using quartiles/tertiles (ng ml⁻¹): Tretli breast (lower: 12.9, upper: 33.9), Tretli Haematological: (lower:14.3, upper: 34.1), Tretli colorectal: (lower:16.4, upper: 38), Tretli lung: (lower:14.3, upper: 34.1), Vrieling: (lower: 10.6, upper: NA), Kelly (NA), Fedirko: (lower:11.8, upper: 33.4), Ng *et al* (2011): (lower:9.6, upper: 30.7), Zgaga: (lower:4.4, upper: 18.3), Ng *et al* (2008): (lower:21, upper: 30.6), Liu: (lower:7, upper: 25.4), Zhou: (lower:7.4, upper: 24.5), Heist: (lower:10.4, upper: 23.9), Meyer: (lower:16.2, upper: 34.2), Fang: (NA), Muller: (NA).

Adjusted Meta-analysis: Disease progression for circulating 25(OH)D

| Author(s) and Year | Population | N.O. Score | Ν | Exposure (ng ml ⁻¹) | | Weight | HR (95% CI) |
|--|--|---|--|--|----------------|--|---|
| Breast Hatse, 2012 Goodwin, 2009 Kim, 2011 Jacobs, 2011 Lohmann, 2015 Lim, 2015 Vrieling, 2014 Clark, 2014 Subtotal (Breast) | Belgium Canada Korea USA Canada Korea Germany USA | 5 7 7 5 8 7 5 | 1800 512 310 512 934 491 2177 82 | >30 vs <30 >29 vs <20 >30 vs <20 >30 vs <10 >20 vs <16 >20 vs <20 Tertile 3 vs Tertile 1 >20 vs <20 | <pre></pre> | 0.5% 1.9% 0.8% 1.1% 1.7% 0.2% 4.3% 0.8% | $\begin{array}{c} 0.25 & (\ 0.09 & , 0.70 \\ 0.58 & (\ 0.35 & , 0.98 \\ 0.25 & (\ 0.11 & , 0.56 \\ 0.88 & (\ 0.43 & , 1.75 \\ 1.14 & (\ 0.65 & , 1.99 \\ 0.45 & (\ 0.25 & , 1.99 \\ 0.57 & (\ 0.40 & , 0.81 \\ 0.77 & (\ 0.34 & , 1.75 \\ 0.66 & (\ 0.45 & , 0.88 \\ \end{array}$ |
| Haemotological Lee, 2014 Shanafelt, 2011 Drake (PostFL), 2010 Drake (FL), 2010 Drake (MCL), 2010 Drake (MCL), 2010 Drake (DLBCL), 2010 Drake (DLBCL), 2010 Pardanani (MDS), 2011 Pardanani (MDS), 2011 Kelly (Lysa cohort), 2015 Kelly (Lysa cohort), 2015 Bittenbring, 2014 Subtotal (Haemotological) | USA USA USA USA USA USA USA USA USA USA | 4 8 7 7 7 7 7 7 5 5 7 7 5 5 7 7 5 | 97 543 109 285 71 70 370 78 74 247 183 240 359 | >32 vs <20 >25 vs <25 >25 vs <25 Tertile 3 vs Tertile 1 Tertile 3 vs Tertile 1 >8 vs <8 | | $\begin{array}{c} 1.1\% \\ 6.7\% \\ 1.2\% \\ 3.0\% \\ 1.4\% \\ 1.4\% \\ 4.0\% \\ 1.1\% \\ 0.2\% \\ 3.1\% \\ 1.3\% \\ 1.6\% \\ 2.7\% \end{array}$ | $\begin{array}{c} 0.39 & (0.19 & 0.78) \\ 0.68 & (0.51 & 0.90) \\ 1.02 & (0.53 & 1.96) \\ 0.93 & (0.62 & 1.41) \\ 0.92 & (0.50 & 1.69) \\ 0.52 & (0.28 & 0.96) \\ 0.71 & (0.49 & 1.02) \\ 0.87 & (0.43 & 1.75) \\ 0.77 & (0.25 & 3.30) \\ 0.83 & (0.63 & 1.25) \\ 0.76 & (0.40 & 1.45) \\ 0.65 & (0.37 & 1.16) \\ 0.71 & (0.47 & 1.10) \\ 0.75 & (0.61 & 0.88) \end{array}$ |
| Head and Neck Meyer, 2011 Gugatschka, 2011 Subtotal (Head and Neck) | Canada Austria | 8 4 | 540 88 | Quartile 4 <i>vs</i> Quartile 1 Per 10 ng ml ⁻¹ increase | | 1.8% 0.2% | 1.12(0.65,1.93) 0.20(0.04,0.98) 0.98(0.33,1.63) |
| Colorectal Ng, 2011 Mezawa, 2010 Subtotal (Colon) | USA Japan | 8 5 | 515 257 | Quartile 4 <i>vs</i> Quartile 1 Per 10 ng ml ⁻¹ increase | | 6.6% 0.5% | 1.07(0.81,1.42) 1.22(0.45,3.30) 1.08(0.81,1.35) |
| Prostate Holt, 2013 | USA | 7 | 1476 | >20 <i>vs</i> <20 | - | 13.0% | 0.91 (0.83 , 1.10) |
| Skin Samimi, 2014 Newton-Bishop, 2009 Subtotal (Skin) | France UK | 6 5 | 89 872 | >20 vs <20 per 8 increase | | 0.5% 13.8% | 0.34 (0.12 , 0.97) 0.79 (0.64 , 0.96) 0.77 (0.58 , 0.97) |
| Pancreatic Van Loon, 2014 | USA | 4 | 256 | Per 10 ng ml ⁻¹ increase | r ≖ ⊣ | 15.8% | 1.00 (0.82 , 1.20) |
| Ovarian Webb, 2015 | Australia | 8 | 670 | >20 <i>vs</i> <10 | ⊢ ∎-1 | 7.7% | 0.87 (0.60 , 1.13) |
| OVERALL CANCER | | | 14,307 | | • | | 0.84 (0.77 , 0.91) |
| | | | | | 0.00 1.00 2.00 | | |
| | | | | | Hazards ratio | | |

Figure 4. Cancer progression and 25-hydroxyvitamin D concentration: meta-analysis of adjusted hazard ratios. HR are sortd by the difference in median between high and low vitamin D levels compared. Acute myeloid leukaemia (AML), Chronic Lymphoid Leukaemia (CLL), and subtypes of non-Hodgkin's lymphoma (NHL) (large B-cell lymphoma (DLBCL), T-cell lymphoma (TCL), follicular lymphoma (FL) and mantle cell lymphoma (MCL)). I2 = breast: 0, haematological: 0, colorectal: 0, head and neck: 0, skin: 0 overall cancer: 0. Approximated median in studies using quartiles/ tertiles (ng ml⁻¹): Vrieling: (lower: 10.6, upper: NA), Kelly (NA), Meyer: (lower:16.2, upper: 34.2), Ng et al, (2011): (lower:9.6, upper: 30.7).

mortality. By far, the most commonly studied were polymorphisms in VDR gene, particularly rs2228570 (FokI), rs1544410 (BsmI), rs731236 (TaqI), rs11568820 (Cdx2), and rs7975232 (ApaI). In meta-analysis, rs1544410 TT/TC genotypes were associated with worse survival compared to CC genotype (HR = 1.40, 95%) CI = 1.05 - 1.75; Figure 5). The same direction of the effect was observed in the sensitivity analyses after exclusion of studies with NOS<7 (Supplementary Figure S4) and those reporting on cancer-specific mortality, but the association was no longer significant (Supplementary Figure S5). In lung cancer patients, a poorer outcome was observed to be associated with rs2228570 TT/ TC carriers (HR = 1.29, 95% CI = 1.00-1.57) and a consistent albeit non-significant association was found across all cancers (HR = 1.26, 95% CI = 0.96–1.56). A significant association was observed with rs731236 (Taq1) variant when limited to studies at low risk of bias (NOS score \geq 7; HR = 0.79, 95% CI = 0.62–0.95, Supplementary Figure S4). Other genetic factors were investigated in at most three original studies and no other statistically significant results were observed.

VDR and vitamin D pathway SNPs and disease progression. Ten studies examined the effect of genetic variation on disease progression (Figure 6; for sensitivity analysis see Supplementary Figure S6). In meta-analysis of three studies with a total of 1588 patients, it was observed that *rs*7975232 *AA* carriers had significantly worse survival than CC carriers (HR = 1.29, 95% CI = 1.02–1.56). Additionally, a suggestive association was observed for vitamin D binding protein variant *rs*2282679 (HR = 1.22, 95% CI = 0.99–1.46) in meta-analysis of two studies.

Testing for publication bias and study heterogeneity. There was some evidence of heterogeneity between studies in meta-analysis of 25OHD and some evidence of publication bias (Supplementary Figures S7 and S8). A non-insignificant degree of heterogeneity and evidence of publication bias were observed in some subgroup analysis. Heterogeneity was observed for subgroup analysis of rs1544410, rs7975232, rs2228570 and rs731236, as well as for some individual cancer types while publication bias was observed for rs1544410, rs2228570 and rs731236 (Supplementary Figures S7 and S8).

Studies not included in meta-analysis. Seventeen papers were excluded from the meta-analysis, but their findings were none-theless considered (Table 2). Eight studies report improved overall and/or progression-free survival among those with higher 25OHD concentration (Vrieling *et al*, 2011; Walentowicz-Sadlecka *et al*, 2012; Peiris *et al*, 2013; Field *et al*, 2013; Bade *et al*, 2014; Der *et al*,

2014; Hansson *et al*, 2014; Obermannova *et al*, 2015) and one study found no association between 25OHD and incidence of metastases (Nurnberg *et al*, 2009). Seven studies investigated genetic variants and outcome (median sample size: 66). One study reported that the rs731236/rs2228570 (*TaqI-FokI*, *TTFf/TtFf*) haplotype was significantly associated with reduced overall survival (HR = 1.81, 95% CI = 1.23–3.48, P = 0.04) (Turna *et al*, 2012): suggestive associations were reported between progression-free survival and rs731236 (*AA*) genotype in prostate cancer (Furuya

et al, 1999) and rs2228570 *TT* genotype in breast cancer (Yiallourou *et al*, 2014), while there was no association found between rs2228570 and paediatric ALL (Kim *et al*, 2012). No association was observed between rs1544410 and breast cancer outcome (Yagmurdur *et al*, 2009). There was a suggestive association between platelet VDR expression and survival in ovarian cancer (Silvagno *et al*, 2010). Finally, low vitamin D binding protein (DBP) levels were found to be predictive of lung cancer death (Turner *et al*, 2013).

| hΔ | usted Meta-anal | usis: Survival for | Vitamin D rece | entor and nathwa | y polymorphisms |
|----|-----------------|--------------------|----------------|------------------|-----------------|
| | | | | | |

| Author and Year | Population | N.O. score | Ν | Exposure | | Weight | HR (95% CI) |
|--|--------------------|------------|--------------|------------------------------------|---------------|----------------|--|
| Rs7975232 (Apal) Prostate | Tahuan | | 004 | 14.00 | | 10.4% | |
| Pao, 2013 | Taiwan | 8 | 601 | AA vs CC | | 19.4% | 0.98 (0.46 , 2.11) |
| ∟ungi Kong, 2013 | China | 4 | 755 | AA vs CC | ⊢∎ | 21.4% | 2.84 (2.63 , 3.94) |
| lead anneck Zeljic, 2012 | Serbia | 6 | 110 | AC/AA vs CC | ⊢ ∎1 | 21.3% | 0.98 (0.68 , 1.40) |
| Colorectal Igaga, 2014 | Scotland | 8 | 1598 | AA vs CC | L | 21.1% | 1.01 (0.74 , 1.53) |
| Kidney Dbara, 2007 | Japan | 5 | 135 | AA vs AC/CC | | ▶ 16.8% | 3.30 (1.01 , 10.60) |
| SUBTOTAL | | | 3,199 | | | | 1.77 (0.79, 2.75) |
| Rs1544410 (Bsml) | | | -, | | | | |
| Prostate Pao,2013 Holt,2010 Subtotal (Prostate) | Taiwan USA | 8 8 | 601 1294 | TT vs CC TT vs CC | | ► 2.5% 7.6% | 5.04 (0.68 , 37.44) 0.80 (0.40 , 1.90) |
| ∟ung Heist,2008 | | 7 | 004 | TT up CC | | 10.0% | 100(074,150) |
| _iu,2011 | USA China | 7 6 | 294 568 | TT <i>vs</i> CC TT <i>vs</i> CC | | 12.3% 5.4% | 1.08 (0.74 , 1.56) 4.33 (1.34 , 14.00) |
| Zhou,2006 Subtotal (Lung) | USA | 7 | 373 | TT vs CC | | 12% | 1.35 (0.90 , 2.03) 2.05 (0.94 , 3.16) |
| lead and neck Zeljic,2012 | Serbia | 6 | 110 | TT/TC vs CC | F | 8.0% | 1.33 (0.77 , 2.99) |
| Colorectal | | | | | | | |
| gaga,2014 edirko,2012 subtotal (Colorectal) | Scotland Europe | 8 8 | 1598 1202 | TT <i>vs</i> CC TT <i>vs</i> CC | | 13.1% 12.9% | 1.09 (0.74 , 1.44) 1.18 (0.87 , 1.61) 1.13 (0.92 , 1.34) |
| Skin Drlow,2014 | Worldwide | 7 | 3566 | TT vs CC | H a -(| 13.7% | 0.79 (0.64 , 0.96) |
| Glioma Anic,2012 | USA | 5 | 320 | TT <i>vs</i> CC | · | 12.5% | 1.41 (0.96 , 2.00) |
| SUBTOTAL | | | 9,926 | | • | | 1.40 (1.05 , 1.75) |
| Rs11568820 (Cdx2) | | | 0,020 | | | | |
| Breast Pema,2013 | Germany | 6 | 498 | TT vs CC | F | ▶ 1.3% | 1.50 (0.30 , 6.60) |
| Prostate Penney,2010 | | 0 | 1000 | T 00 | L | 00/ | 4.00 (0.77, 0.40) |
| Penney,2010 Pao,2013 | USA Taiwan | 6 8 | 1292 601 | TT <i>vs</i> CC TT <i>vs</i> CC | | 8% 7.2% | 1.38 (0.77, 2.49) |
| Subtotal (Prostate) | Taiwaii | 0 | 001 | 11 15 00 | \diamond | <i>1.2</i> /o | 1.34 (0.72 , 2.50) 1.36 (0.93 , 1.79) |
| Lung Heist,2008 | | | | | | | |
| | USA | 7 | 294 | TT vs CC | | 12.9% | 0.98 (0.61, 1.56) |
| Zhou,2006 Subtatal (Lupa) | USA | 7 | 373 | TT/TC vs CC | | 7.7% | 0.92 (0.50 , 1.68) 0.96 (0.59 , 1.33) |
| Subtotal (Lung) | | | | | \sim | | 0.96 (0.59 , 1.33) |
| Colorectal Pema,2013 | Germany | 7 | 1379 | TT vs CC | | 9.4% | 1.02 (0.59 , 1.76) |
| Zgaga,2014 | Scotland | 8 | 1598 | TT vs CC | ⊢_ ∎ | 14.9% | 0.90 (0.41 , 1.39) |
| Subtotal (Colorectal) | | | | | \diamond | | 0.95 (0.61 , 1.29) |
| Skin Orlow,2014 | Worldwide | 7 | 3566 | TT vs CC | ⊦∎⊣ | 27.5% | 0.88 (0.68 , 1.21) |
| Glioma Anic,2012 | USA | 5 | 320 | TT <i>vs</i> CC | ⊢ | 11.3% | 0.98 (0.59 , 1.61) |
| SUBTOTAL | | | 9,921 | | + | | 1.00 (0.84 , 1.17) |

Figure 5. Cancer survival and vitamin D receptor polymorphisms and other vitamin D-related genetic factors: adjusted meta-analysis. *I*² for Apal: 0.95, Bsml prostate: 0.93, Bsml Lung: 0.93, Bsml colorectal: 0, Bsml All: 0.85, Cdx2 prostate: 0, Cdx2 lung: 0, Cdx2 colorectal: 0, Cdx2 All: 0, Fokl Prostate: 0, Fokl lung: 0, Fokl colorectal: 0, Fokl All: 0.83, Taql breast: 0.88, Taql skin: 0.46, Taql all: 0.86, Cyp24a1(1) all: 0.75, Cyp24a1(2) all: 0.67, GC all: 0, Rs2107301 all: 0, Rs4516035: 0, Rs2238135: 0.

| Breads Perna,2013 Germany 6 498 TT w CC 4.4% 0.70 (0.20, 2.10) Provaline Para,2013 Taiwan 8 601 TT w CC Provaline Biological Statute B% 0.81 (0.46, 1.44) Lang Macobia USA 7 294 TT w CC Provaline Biological Statute B% 0.81 (0.46, 1.44) Lang Macobia USA 7 294 TT w CC Provaline Biological Statute B% 0.81 (0.46, 1.44) Lang Macobia USA 7 294 TT w CC Provaline Biological Statute B% 0.81 (0.46, 1.44) Colorectal Perna,2013 Germany Scalecal Z Biological Statute 7 573 TT w CC Provaline Biological Statute 100% 0.96 (0.66, 1.32) Subbal Colorectal Perna,2012 Scale and Scalecal Biological Statute 101% 0.96 (0.66, 1.32) 0.96 (0.66, 1.32) Subbal Scalecal Biological Statute TT w CC 101% 0.96 (0.67, 1.13) Head and nock Subbal Scale Biological Statute 7 3566 TT w CC 9.3% 1.42 (0.96, 2.13) <th>Rs2228570 (Fokl)</th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> | Rs2228570 (Fokl) | | | | | | | |
|---|----------------------|-----------|---|--------|-----------------|------------------------|--------------|-----------------------|
| Pad. 2013 Taiwan 8 601 Tit vs CC Pad. 2010 8% 0.61 (0.64, 1.44) Bull 2010 USA 8 1282 Tit vs CC Pad. 2010 6.1% 0.80 (0.30, 1.60) Subball (Prostate) USA 7 234 Tit vs CC Pad. 2010 9.5% 1.41 (0.96, 207) Zhou 2006 USA 7 373 Tit vs CC Pad. 1.44 9.5% 1.31 (0.74, 1.74) Zagea 2014 Southal (Unot) 8 1538 Tit vs CC Pad. 1.44 9.5% 1.31 (0.74, 1.74) Subball (Colorectal) 8 1538 Tit vs CC Pad. 1.06 9.5% 1.72 (1.08, 2.70) Subball (Colorectal) 8 1282 Tit vs CC Pad. 1.07% 0.96 (0.77, 1.19) Head and nock 200/2012 Serbia 6 110 Tit vs CC Pad. 1.02 (0.06, 2.13) Skin Ordrow 2014 Worldwide 7 3566 Tit vs CC Pad. 1.02 (0.06, 0.77, 1.19) Glona Anic. 2012 USA 5 320 Tit vs CC Pad. 1.22 (0.06, 0.27, 1.19) Glona Anic. 2012 | Breast Perna,2013 | Germany | 6 | 498 | TT vs CC | ⊢ ∎ | 4.4% | 0.70 (0.20 , 2.10) |
| Pad. 2013 Taiwan 8 601 Tit vs CC Pad. 2010 8% 0.61 (0.64, 1.44) Bull 2010 USA 8 1282 Tit vs CC Pad. 2010 6.1% 0.80 (0.30, 1.60) Subball (Prostate) USA 7 234 Tit vs CC Pad. 2010 9.5% 1.41 (0.96, 207) Zhou 2006 USA 7 373 Tit vs CC Pad. 1.44 9.5% 1.31 (0.74, 1.74) Zagea 2014 Southal (Unot) 8 1538 Tit vs CC Pad. 1.44 9.5% 1.31 (0.74, 1.74) Subball (Colorectal) 8 1538 Tit vs CC Pad. 1.06 9.5% 1.72 (1.08, 2.70) Subball (Colorectal) 8 1282 Tit vs CC Pad. 1.07% 0.96 (0.77, 1.19) Head and nock 200/2012 Serbia 6 110 Tit vs CC Pad. 1.02 (0.06, 2.13) Skin Ordrow 2014 Worldwide 7 3566 Tit vs CC Pad. 1.02 (0.06, 0.77, 1.19) Glona Anic. 2012 USA 5 320 Tit vs CC Pad. 1.22 (0.06, 0.27, 1.19) Glona Anic. 2012 | Prostate | | | | | | | |
| Hold 2010 USA 8 1292 TT vs CC Image: Comparison of the state of the | | Taiwan | 0 | 601 | TT vc CC | | Q 0/. | 0.91 (0.46 1.44) |
| Subtotal (Prostate) 0.81 (0.34, 1.28) Lung Heisl2008 USA 7 294 TT vs CC 9.5% 1.41 (0.96, 2.07) Zhou,2006 USA 7 373 TT vs CC 9.5% 1.41 (0.96, 2.07) Zhou,2006 USA 7 373 TT vs CC 9.5% 1.41 (0.96, 2.07) Colorectal Ferm,2013 Germany 7 1379 TT vs CC 10% 0.96 (0.68, 1.32) Zepiga,2014 Sottand 8 1202 TT vs CC 10% 0.96 (0.67, 1.13) Valuation (Colorectal) 8 1202 TT vs CC 9% 1.72 (1.98, 2.70) Skin Orarian 10% 0.96 (0.67, 1.13) 0.94 (0.59, 1.24) 0.94 (0.59, 1.24) Gloma Anic 2012 USA 5 320 TT vs CC 10.7% 0.96 (0.77, 1.19) Gloma Anic 2012 USA 5 320 TT vs CC 9.3% 1.42 (0.96, 2.13) Ovarian Tame2.2009 Japan 7 101 TT/CT vs CC | , | | | | | | | |
| Lung Heiz, 2008 USA 7 294 TT vs CC 9.5% 1.41 (0.96, 2.07) Subtotal (Lung) USA 7 373 TT vs CC 9.5% 1.13 (0.74, 1.74) Subtotal (Lung) Germany 7 373 TT vs CC 9.5% 1.13 (0.74, 1.74) Colorectal Berma, 2013 Germany 7 1379 TT vs CC 10% 0.96 (0.58, 1.32) Zgega, 2014 Scotland 8 1598 TT vs CC 10% 0.94 (0.77, 1.28) Subtotal (Colorectal) B 1202 TT vs CC 9% 1.72 (1.08, 2.70) Skin 0 0.76 (0.77, 1.19) 0.95 (0.77, 1.19) 0.95 (0.77, 1.19) Skin 0 10.7% 0.96 (0.77, 1.19) 0.96 (0.77, 1.19) Skin 0 10.7% 0.96 (0.77, 1.19) 1.42 (0.96, 2.13) Orarian Tamez, 2009 Japan 7 101 TT/CT vs CC 3.6% 5.56 (1.64, 20.00) SUBTOTAL 11.334 126 (0.96, 1.56) 11.5% 0.90 (5.27, 1.52) 0.90 (5.27, 1.52) Perma, 2013 Germany 6 488 <td>,</td> <td>USA</td> <td>0</td> <td>1292</td> <td>11 13 00</td> <td></td> <td>0.176</td> <td></td> | , | USA | 0 | 1292 | 11 13 00 | | 0.176 | |
| Heic 2008 USA 7 294 TT vs CC 9.7% 1.141 (0.56, 2.07) Subtotal (Lung) USA 7 373 TT vs CC 9.1% 1.131 (0.74, 1.74) Subtotal (Lung) Germany 7 1379 TT vs CC 9.1% 1.131 (0.74, 1.74) Pena, 2013 Germany 7 1379 TT vs CC 10% 0.96 (0.68, 1.32) Zaga, 2014 Sobtand 8 1598 TT vs CC 10% 0.94 (0.59, 1.29) Subtotal (Colorectal) Bad and neck 2gin, 2012 Serbia 6 110 TT/TC vs CC 9% 1.72 (1.08, 2.70) Skin Orlow 2014 Worldwide 7 3566 TT vs CC 9% 1.72 (1.08, 2.70) Skin Orlow 2014 Worldwide 7 3566 TT vs CC 9.3% 1.42 (0.56, 2.13) Ovarian Tamez.2009 Japan 7 101 TT/CC vs CC 9.3% 1.42 (0.56, 2.13) Subtotal (freest) Japan 7 101 TT/CC vs CC 9.3% 0.50 (0.20, 1.50) Subtotal (freest) Japan 7 <td>Subtotal (Prostate)</td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td>0.81 (0.34, 1.28)</td> | Subtotal (Prostate) | | | | | | | 0.81 (0.34, 1.28) |
| Zhoi.2006 USA 7 373 TT vs CC 9.1% 1.13 (0.74, 174) Subtola Germany 7 1379 TT vs CC 14 10% 0.66 (0.68, 1.32) Zagap.2014 Scotland 8 1598 TT vs CC 14 10% 0.64 (0.68, 1.32) Zagap.2014 Scotland 8 1598 TT vs CC 14 10% 0.64 (0.68, 1.32) Fedirk.2012 Serbia 6 110 TT/TC vs CC 9% 1.72 (1.08, 2.70) Skin Orlow.2014 Worldwide 7 3566 TT vs CC 9% 1.72 (1.08, 2.70) Skin Orlow.2014 Worldwide 7 3566 TT vs CC 9.3% 1.42 (0.96, 2.13) Ovarian Anic.2012 USA 5 320 TT vs CC 9.3% 1.42 (0.96, 2.13) Ovarian Tamez.209 Japan 7 101 TT/CC vs CC 9.3% 1.42 (0.96, 2.13) Ovarian Scoole, 202 UK 8 721 (GG vs AA 12.5% 0.56 (1.64, 20.00) SUBTOTAL 11.334 128 (GG vs AA | 0 | | _ | | | | | |
| Subtrati (Lung) 129 (1.00, 1.57) Colorectal 100% 0.06 (0.68, 1.32) Jenaz 2013 Germany 7 1379 TT vs CC Jendito.2012 Europe 8 1202 TT vs CC Jendito.2012 Europe 8 1202 TT vs CC Jendito.2012 Serbia 6 110 TT/TC vs CC Subtrati (Colorectal) 9% 1.72 (1.08, 2.70) Head and neck Zelje.2012 Serbia 6 110 TT/TC vs CC Skin Orov. 2014 Worldwide 7 3566 TT vs CC H 10.7% 0.96 (0.77, 1.19) Glioma Anic. 2012 USA 5 320 TT vs CC H 10.7% 0.96 (0.77, 1.19) Glioma Tanez.2009 Japan 7 101 TT/CT vs CC Jack 5.56 (1.64, 20.00) SUBTOTAL 11.334 126 (0.36, 1.54) 12.3% 0.56 (0.35, 1.54) Lundn, 1989 Swaden 4 121 GG vs AA Sec% 3.6% 0.50 (0.20, 1.51) Subtroti (Greast) Germany 6 | | | | | | H | | |
| Colorectal Pena, 2013 Germany 7 1379 TI vs CC Pena, 2014 Scotland 8 1598 TI vs CC Pena, 2012 Europe 8 1202 TI vs CC Pediticul (Colorectal) Head and neck Zeljic. 2012 Serbia 6 110 TI/TC vs CC Pena, 2014 Worldwide 7 3566 TI vs CC Pena, 2012 USA 5 320 TI vs CC Pena, 2012 USA 5 320 TI vs CC Pena, 2012 USA 5 320 TI vs CC Pena, 2013 Germany 7 101 TI/CT vs CC Pena, 2013 Germany 6 498 Gi vs AA Hot, 2010 USA 8 1292 GG vs AA Hot, 2011 China 6 586 GG vs AA Hot, 2012 USA 5 320 GG vs AA Hot, 2013 Germany 7 1379 GG vs AA Hot, 2010 USA 8 1292 GG vs AA Hot, 2010 China 6 586 GG vs AA Hot, 2012 USA 5 320 USA | Zhou,2006 | USA | 7 | 373 | TT vs CC | | 9.1% | |
| Perma 2013 Germany 7 1379 TT vs CC Image: CC Image | Subtotal (Lung) | | | | | \diamond | | 1.29(1.00,1.57) |
| Zagaz.014 Scolland 8 1598 TT vs CC Image: Constraint of the state of | Colorectal | | | | | | | |
| Zapaz.2014 Scotland 8 1598 TT vs CC Image: Constraint of the sector of the sec | Perna,2013 | Germany | 7 | 1379 | TT vs CC | ⊢ ∎ <u>−</u> −1 | 10% | 0.96 (0.68, 1.32) |
| Fadinc.2012 Europe 8 1202 TT vs CC Image: Constraints of the sector of the sec | | | | | | | | |
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| | Zeljic, 2012 | | | | | -∎ | | |
| Subtotal 632 0.99 (0.44 , 1.55) | | Cananda | 8 | | AA <i>vs</i> GG | | 59.5% | |
| | Subtotal | | | 632 | | | | 0.99 (0.44 , 1.55) |

Figure 5. (Continued)

DISCUSSION

This is the first systematic review with meta-analysis that examines the relationship between cancer outcomes and variation in vitamin D pathway genes, and also by far the largest review on vitamin D status and cancer outcome. Our review suggests that higher circulating vitamin D in cancer patients is associated with a 26% lower rate of death and a 16% lower rate of disease progression. The clear association with survival was also observed in site-specific analyses of breast, haematological and colorectal cancers, while an association with reduction in disease progression was also found in those diagnosed with breast, haematological and skin cancer.

Establishing a causal relationship between vitamin D status and cancer progression is challenging because risk factors associated with cancer outcome are often also associated with vitamin D deficiency. For example, the association between 250HD and improved survival observed in the original studies might be due to 250HD being a marker of healthier lifestyle (i.e., healthier diet containing more fish; physical activity and spending time outdoors). However, evidence that genetic factors linked to vitamin D metabolism and pathways impact upon cancer survival may be used to counter such concerns and support a causal link. In our meta-analysis, we found evidence of an association between the *VDR* gene variants with functionally characterised effects and cancer outcome. Forty percent higher rate of death was observed in *TT* carriers at rs1544410 locus and 26% higher rate in *TT* carriers at rs2228570, while 29% increased risk of disease progression was observed in *AA* carriers at *rs*7975232 and 22% in *GG* carriers at GC locus.

Evidence from biological studies support a role for these polymorphisms in modulating vitamin D biology. For example, rs2228570 has been shown to affect the translational start site of $1,25(OH)_2D$ and hence its downstream effects (Uitterlinden *et al*, 2004), while rs1544410 and rs7975232 have been associated with changes in *VDR* messenger RNA expression (Staal *et al*, 1996; Uitterlinden *et al*, 2004). We hypothesise that interactions between mutations in the vitamin D pathway and vitamin D status exist,

| Rs6013897 (CYP24A1) | (2)) | | | | | | |
|------------------------------------|-----------|---|---------------|---------------------|----------------|--------|--|
| Prostate Penney, 2010 Glioma | USA | 6 | 1292 | AA <i>vs</i> GG | ⊢ 1 | 45% | 1.28 (0.69 , 2.31) |
| Anic, 2012 Subtotal | USA | 5 | 320 1,612 | AA <i>vs</i> GG | | 55% | 0.62 (0.40 , 0.96) 0.92 (0.28 , 1.55) |
| Rs2282679 (GC) | | | | | | | |
| Glioma Anic, 2012 | USA | 5 | 320 | AA <i>vs</i> CC | ⊢ | | 1.04 (0.69 , 1.54) |
| | UUA | 5 | 020 | AA 13 00 | | 22.8% | 1.04 (0.03 , 1.04) |
| Skin Davies, 2014 | Europe | 7 | 3137 | AA vs CC | ⊢ ∎ | | 1.19 (0.86 , 1.66) |
| Prostate | | | | | | 30.7% | |
| Trummer, 2014 | Austria | 6 | 702 | AA vs AC | ⊢ = ⊣ | 46.5% | 0.89 (0.68 , 1.17) |
| Subtotal | | | 4,159 | | + | | 1.02 (0.83 , 1.20) |
| Rs2107301 | | | | | | | |
| Prostate Penney, 2010 | USA | 6 | 1292 | CC vs TT | ⊢ | 11.1% | 1.09 (0.62 , 1.88) |
| Glioma Anic, 2012 | USA | 5 | 320 | CC vs TT | F=1 | 19.7% | 1.02 (0.67 , 1.54) |
| Skin | | - | 0500 | 00 · · · · T | H I H | 00.00/ | 0.00 (0.74 . 4.40) |
| Orlow, 2014 Subtotal | Worldwide | 7 | 3566 5,178 | CC vs TT | | 69.2% | 0.90 (0.74 , 1.12) 0.94 (0.76 , 1.13) |
| Rs4516035 (EcoRV) | | | 5,176 | | | | 0.34 (0.70 , 1.10) |
| Prostate | | | | | | | |
| Penney, 2010 | USA | 6 | 1292 | TT <i>vs</i> CC | ⊢ | 36.2% | 1.15 (0.67 , 1.93) |
| Glioma Anic, 2012 | USA | 5 | 200 | TT <i>vs</i> CC | | CO 09/ | 1.05 (0.00 , 1.05) |
| Subtotal | 05A | Э | 320 1,612 | TT VS CC | | 63.8% | 1.25 (0.86, 1.85) 1.22 (0.91, 1.53) |
| Rs2238135 | | | 1,012 | | | | |
| Glioma | | _ | 105 | ~~~~~ | | 10.00/ | |
| Anic, 2012 Skin | USA | 5 | 135 | CC vs GG | | 16.8% | 0.94 (0.58, 1.49) |
| Orlow, 2014 | Worldwide | 7 | 3566 | CC vs GG | ⊢ a -i | 83.2% | 0.96 (0.78 , 1.18) |
| Subtotal | | | 3,701 | | • | | 0.96 (0.77 , 1.14) |
| | | | | | | | |
| | | | | | | | |
| | | | | | 0.00 1.50 3.00 | 0 4.00 | |
| | | | | | | | |

Figure 5. (Continued)

and that this interaction could have a critical role in cancer prognosis. Indeed, Han *et al* (Han *et al*, 2007) have shown an interaction between vitamin D intake and rs1544410 polymorphism on cancer risk, and we and others have previously shown a modification of the relationship between vitamin D intake or status and cancer outcome by other *VDR* variants, thus suggesting an interaction of genetic and environmental factors (Li *et al*, 2007; Theodoratou *et al*, 2008; Anderson *et al*, 2011; Zgaga *et al*, 2014). In conjunction with the strong associations observed for vitamin D status, evidence from genetic studies further supports an important role of vitamin D in cancer progression.

Few studies to date have analysed the associations between *VDR* or vitamin D pathway genetic variants and cancer outcomes, and no meta-analyses have been published to date. A review by Kostner *et al* (Kostner *et al*, 2009) concluded that associations between *VDR* polymorphisms and cancer prognosis are strongest for prostate cancer (rs2228570), breast cancer (rs1544410, rs731236) malignant melanoma (rs1544410), and renal cell carcinoma (rs731236) but did not perform meta-analysis on these data.

Interestingly, Afzal *et al* (Afzal *et al*, 2014) have employed principles of Mendelian randomization in a study comprising 95 766 participants and found that variation in genes involved in vitamin D and 25OHD synthesis (*DHCR7* and *CYP2R1*) were associated with both all-cause and cancer mortality, supporting a causal role of vitamin D. To date, there are no published findings from randomised controlled trials (RCT) assessing the effect of vitamin D supplementation on survival in cancer patients, although several ongoing trials (unfortunately only some of which have disease

progression as an outcome) were identified (ClinicalTrials.gov, 2016). Meanwhile, the data on cancer mortality from RCTs conducted in the general population can offer some insight; most notably, a Cochrane review of randomised studies comparing vitamin D supplements to placebo identified a significant reduction in cancer mortality in those taking vitamin D supplements (HR = 0.88, 95% CI = 0.78–0.98; Bjelakovic *et al*, 2014).

Hazard ratio

A major issue that is typically taken poor notice of in vitamin D meta-analyses-namely, a very large variability in vitamin D category definition amongst studies, is for the first time being highlighted and transparently shown in our review. Vitamin D categories differed in level as well as range-as a result, large heterogeneity in exposure definition occurred and study pointestimates are difficult to compare: it is, for example, unsurprising that the reported effect per 20 ng ml^{-1} is greater than effect per 5 ng ml^{-1} increase. Therefore, there is a need for a consensus in category definition and reporting of effect sizes: future original studies should report effect sizes using internationally agreed cutoffs, such as those given by the Institute of Medicine, solely or in addition to study-specific cutoff values chosen. Generally, variability in exposure categories results in a more heterogeneous estimates and is likely to increase statistical uncertainty and hence bias results towards the null. Nonetheless, our summary findings remain largely unchanged when the analysis was limited according to the difference in 25OHD between the compared groups.

There are some additional limitations of the present work. First, a number of relevant studies were published after the time limits stipulated in our search strategy and so are not included in our meta-analysis. Some such papers support the conclusions presented here (Brandstedt *et al*, 2016; Fang *et al*, 2016; Fanidi *et al*, 2016; Mondul *et al*, 2016; Orlow *et al*, 2016; Yao *et al*, 2016; Yuan *et al*, 2016), while others reported no association between circulating vitamin D and cancer outcome (Vashi *et al*, 2015; Ahn *et al*, 2016; Danilovic *et al*, 2016; McGovern *et al*, 2016).

| Adjusted meta-analysis: disease progression for vitamin [| D receptor and pathway polymorphisms |
|---|--------------------------------------|
|---|--------------------------------------|

| Author and Year | Population | N.o. score | N | Exposure | | Weight | HR (95% CI) |
|----------------------------|------------|------------|--------------|--------------------|---------------------------------------|---------|--|
| Rs7975232 (Apal) | | | | | | | |
| Breast | | | | | | | |
| Mishra, 2013 | USA | 6 | 232 | AA vs AC/CC | F | 16.5% | 1.08 (0.59 , 2.00 |
| Prostate Pao, 2013 | Taiwan | 8 | 601 | AA vs CC | H -I | 64.2% | 1.31 (1.94 , 1.84 |
| Lung | China | 4 | 755 | | | 10.0% | 1 40 (0 00 0 70 |
| Xiong, 2013 SUBTOTAL | China | 4 | 755 1,588 | AA vs CC | • | 19.2% | 1.43 (0.99 , 2.78 1.29 (1.02 , 1.56 |
| | | | ., | | · · · · · · · · · · · · · · · · · · · | | |
| Rs1544410 (Bsml) Breast | | | | | | | |
| Mishra, 2013 | USA | 6 | 232 | TT vs TC/CC | | 10.4% | 1.49 (0.53 , 3.33 |
| Pande, 2013 | USA | 6 | 1029 | TT vs CC | H HH H | 31.8% | 1.05 (0.92 , 1.21 |
| Subtotal (Breast) | | | | | \diamond | | 1.08 (0.85 , 1.32 |
| Prostate Pao, 2013 | Taiwan | 8 | 601 | TT vs CC | H | 2.3% | 1.00 (0.14 , 7.21 |
| Williams, 2004 | USA | 7 | 728 | TT vs CC | | 13.1% | 0.60 (0.31 , 1.18 |
| Holt, 2010 | USA | 8 | 1294 | TT vs CC | | 18.4% | 1.10 (0.60 , 1.80 |
| Subtotal (Prostate) | UUA | 0 | 1204 | 11 13 00 | \diamond | 10.476 | 0.93 (0.54 , 1.32 |
| Lung | | | | | | | |
| Liu, 2011 | China | 6 | 568 | TT vs CC | ⊢ ∎−−−1 | 24.1% | 1.64 (1.16 , 2.31) |
| SUBTOTAL | | | 4,452 | | • | | 1.19 (0.88 , 1.50) |
| Rs2228570 (Fokl) | | | | | | | |
| Breast Mishra, 2013 | | G | 232 | | ⊢ ∎i | 01 70/ | 0.57 (0.20 . 0.00 |
| Prostate | USA | 6 | 232 | TT vs TC/CC | | 21.7% | 0.57 (0.30 , 0.90 |
| Pao, 2013 | Taiwan | 8 | 601 | TT vs CC | ⊢ ⊷ ⊣ | 22.6% | 0.99 (0.74 , 1.33 |
| Holt, 2010 | USA | 8 | 1292 | TT vs CC | | 19.9% | 1.01 (0.60 , 2.00 |
| Subtotal (Prostate) | 00/1 | 0 | 1202 | | \diamond | 1010/0 | 0.99 (0.72 , 1.26 |
| Head and Neck | | | | | | | |
| Hama, 2011 | Japan | 6 | 204 | TT vs TC/CC | | ▶ 15.5% | 3.03 (1.62 , 5.67 |
| Skin Halsall, 2014 | | F | 171 | | | 00.49/ | 0.77 (0.24 . 2.50 |
| SUBTOTAL | UK | 5 | 171 2,500 | TT vs TC/CC | | 20.4% | 0.77 (0.24 , 2.50) 1.28 (0.49 , 2.08) |
| Rs731236 (Taql) | | | , | | | | |
| | | | | | | | |
| Breast Mishra, 2013 | USA | 6 | 232 | GG/AG <i>vs</i> AA | ⊢ | 40.7% | 1.30 (0.70 , 2.40) |
| | | | 202 | | | | |
| Prostate Holt, 2010 | USA | 8 | 1292 | GG <i>vs</i> AA | ⊢ ∎ | 48.8% | 0.80 (0.50 , 1.40) |
| | 00/1 | 0 | | | | 1010/0 | 0.00 (0.00 , 11.0) |
| Skin Halsall, 2014 | UK | 5 | 171 | GG <i>vs</i> AA/AG | | 10.5% | 0.60 (0.22 , 2.00) |
| 11013011, 2014 | OIX | 5 | | | | 10.576 | |
| SUBTOTAL | | | 1,695 | | • | | 0.98 (0.59 , 1.37) |
| | | | | | | | |
| Rs4516035 (EcoRV) | | | | | | | |
| Breast | | | | | | | |
| Pande, 2013 | USA | 6 | 1029 | GG <i>vs</i> AA | H | 53% | 0.95 (0.82 , 1.08) |
| Skin | | | | | | | |
| Halsall, 2014 | UK | 5 | 171 | GG/AG vs AA | • | ▶ 47% | 2.90 (1.10 , 7.50) |
| SUBTOTAL | | | 1,200 | | | - | 1.87 (-0.04 , 3.77 |
| | | | | | | | |
| Rs2282679 (GC) | | | | | | | |
| Prostate | | | | | | | |
| Trummer, 2015 | Austria | 6 | 702 | TG <i>vs</i> TT | i − | 60.1% | 1.26 (0.93 , 1.70) |
| Colorectal | | | | | | | |
| Szkandera, 2013 | Austria | 5 | 264 | GG vs TT/TG | ⊢ ••−−1 | 39.9% | 1.17 (0.81 , 1.69 |
| SUBTOTAL | | | 966 | | | | 1 22 (0 00 1 46 |
| GODIVIAL | | | 900 | | | | 1.22 (0.99 , 1.46) |
| | | | | | | | |
| | | | | | | | |
| | | | | | 0.00 1.50 3.00 | 4.50 | |
| | | | | | Hazard ratio | | |

Figure 6. Cancer progression and vitamin D receptor polymorphisms and other vitamin D-related genetic variants: adjusted meta-analysis. l^2 for Apal: 0, Bsml prostate: 0.52, Bsml breast: 0.1, Bsml All: 0.61, Fokl Prostate: 0, Fokl All: 0.90, Taql all: 0, Rs4516035: 0.94, Rs22382679: 0.

Second, various assays were used for 25OHD measurement in the different studies, while 25OHD was also sampled at variable timepoints, including pre-diagnosis, before treatment and after treatment, which may impact the results. Also, in disease progression studies, different outcome definitions were used for example, disease-free survival, local or distant recurrence.

In the present study, results for all cancers combined are given, in addition to site-specific findings, we yet fully acknowledge that cancer is a heterogeneous disease. However, numerous studies have shown involvement of vitamin D on key hallmarks of cancer, many of which are common to all cancers; preclinical studies demonstrate effects on cell cycle arrest, cell adhesion, differentiation, proliferation, tumour angiogenesis, and apoptosis in human cancer cell lines (Simboli-Campbell et al, 1997; Chen et al, 2000; Krishnan et al, 2003; Deeb et al, 2007; Kizildag et al, 2010; Hsu et al, 2011; Ting et al, 2012), while reduction in cancer proliferation has been shown in carcinogen-exposed rats (Mokady et al, 2000) and cancer phenotypes are more commonly observed in vitamin D receptor (VDR) knockout mice (Zheng et al, 2012). Nevertheless, the heterogeneity in pooled results between different cancer types and the small number of studies for certain cancers limits the strength of the current study in demonstrating an association between circulating 25-hydroxyvitamin D and total cancer survival.

Next, in reporting the impact of genetic variation on outcome, we acknowledge that ethnic differences in VDR variation exist, which might interfere with the findings from genetic studies, as ethnicity is directly linked to the skin type and vitamin D synthesis. Meanwhile, VDR variants may interact with circulating 25OHD to impact outcome, yet only a small number of studies examined these putative gene–environment interactions. Finally, we observed some evidence of heterogeneity and publication bias overall; however, findings from sensitivity analysis were highly consistent and supportive of main findings.

Despite these limitations, the present work includes a novel meta-analysis, investigating the association between vitamin D-related genetic variation and cancer outcome, in addition to a \sim 50% larger meta-analysis of circulating 25OHD and cancer outcome compared to a previous review (Li *et al*, 2014). Moreover, stringent quality assessment of original studies and corresponding sensitivity analysis were conducted and strikingly inconsistent 25OHD category definitions were addressed in stratified analysis.

In conclusion, the consistent evidence across the studies presented in the current review demonstrates a clear and strong association between low baseline vitamin D levels and poorer cancer survival. The associations between vitamin D-related genetic variants and cancer survival support an interpretation that vitamin D may play an important role in influencing cancer outcome. However, a causal link cannot be conclusively established from observational studies; hence, well-designed and adequately powered RCTs are needed to evaluate the clinical application of vitamin D in augmenting standard follow-up and adjuvant chemotherapy regimens. Understanding the mechanism of action of genetic factors promises to provide further insight into biological determinants of response to treatment and could help inform prognosis.

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CONFLICT OF INTEREST

The authors declare no conflict of interest.

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