1 Q1 Research Article

Subsequent Primary Malignancies in Patients with Nonmelanoma Skin Cancer in England: A National Record-

4 Linkage Study

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Abstract

Background: Conflicting evidence exists about whether people with a history of nonmelanoma skin cancer (NMSC) are at higher risk of subsequent primary malignant cancers than those without.

Methods: An all England record-linked hospital and mortality dataset spanning from 1999 to 2011 was used. We constructed two cohorts: one that comprised people with a history of NMSC (502,490 people), and a control cohort that comprised people without. We "followed up" these two cohorts electronically to determine observed and expected numbers of people with subsequent primary cancers in each, based on person-years at risk, and calculated standardized risk ratios (RR).

Results: Comparing the NMSC cohort with the non-NMSC cohort, the RR for all subsequent malignant cancers combined was 1.36 [95% confidence interval (CI), 1.35–1.37]. Significantly increased RRs (P < 0.05) were found for 26 of the 29 cancer types studied, in particular for salivary gland, melanoma, bone, and upper gastrointestinal tract cancers. The RRs were also particularly high when comparing younger people with and without NMSC.

Conclusions: NMSC is strongly associated with a broad spectrum of other primary cancers, particularly in younger age groups. The pattern suggests a genetic or early-acquired etiologic association.

Impact: These results represent what can be done using very large, linked, routinely collected administrative datasets; but such datasets lack detail. Further work to establish the mechanisms behind these associations is warranted. *Cancer Epidemiol Biomarkers Prev;* 1–9. ©2014 AACR.

Introduction

Nonmelanoma skin cancer (NMSC) is the commonest malignancy in white populations, and its incidence is increasing (1, 2). The two key known mechanisms of cutaneous carcinogenesis are UV-induced genetic damage and suppression of skin immune tumor surveillance responses (3). UV exposure is considered to account for an increased risk of subsequent NMSC and melanoma in individuals after a first NMSC (4, 5).

There is conflicting literature on whether individuals with NMSC are at increased risk of developing primary malignancies that are not known to be associated with UV radiation. Comparatively large individual studies investigating this question include two local UK regis-

Note: Supplementary data for this article are available at Cancer Epidemiology, Biomarkers and Prevention Online (http://cebp.aacrjournals.org/).

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43 try-based studies (6, 7), a nationwide study in Finland (8), and a US prospective cohort study of health profes-44 sionals (9), all of which showed significant increases in a 45range of cancers after NMSC. Another UK registry-46 based study, however, showed no such increased risks 47 and indeed significant risk reductions in certain cancers, 48 Q_{249} including of the breast and prostate (10). Of two worldwide meta-analyses that have investigated the literature 50on NMSC and other cancers, one found that the risk of 51cervical, colon, gastric, and rectal cancers is significantly 52reduced in people with NMSC, and concluded that solar 53UVB iance reduces the risk of many internal can-54cers, with the likely mechanism being the protective 55effects of increased production of vitamin D (11); the 56other meta-analysis concluded that NMSC is associated 57with a significantly increased risk of a broad spectrum 58of subsequent malignancies (5). Certain cancers, such as 59salivary, have consistently been found to be particularly 60 associated with NMSC (5, 11). Some evidence exists 61 that second primary cancer risks are higher in younger 62 individuals with NMSC than older (12), suggesting a 63 genetic or early-acquired etiology. 64

We aimed to contribute to the literature by using a linked dataset covering the whole of England during the period from 1999 to 2011, making this the largest single epidemiologic study so far to investigate whether a

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71history of NMSC is a significant risk factor for other 72subsequent primary cancers.

73**Materials and Methods**

Setting and dataset

The dataset contained Hospital Episode Statistics 7576 (administrative data routinely collected for each hospital 77 admission or episode of day-case care in all NHS hospitals) and data on mortality, obtained from national death 78 79registrations, from January 1, 1999, to December 31, 2011. 80 The hospital data were supplied by the English National 81 Information Centre for Health and Social Care and mortality data by the Office for National Statistics. Both 82 83 sources of data contain diagnosis codes using the Inter-84 national Classification of Diseases 10th Revision (ICD-85 10). Successive records for each individual were linked 86 together to create a single dataset for analysis. The record 87 linkage was carried out by the Oxford record-linkage 88 group (13).

89 Subjects

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90 A cohort was constructed comprising people with a record of hospital day-case care or inpatient admission for NMSC in which NMSC was the principal diagnosis on the record, by identifying the earliest episode of day-case care, or inpatient admission, for the condition in an NHS hospital during the study period (the "NMSC cohort"). NMSC was defined using code C44 in the 10th revision of 97 the International Classification of Diseases (ICD).

98A "reference cohort" was constructed (as a control 99 group) by identifying individuals with a record of hos-100 pital day-case care or inpatient admission for various other, mainly minor, medical and surgical conditions and 101 102injuries (see footnote in Table 1). We selected a wide range 103of different conditions, rather than relying on a narrow 104range (in case the latter are themselves atypical in their 105risk of subsequent disease). Any individual in the refer-106 ence cohort who was found to have a subsequent record of 107 NMSC contributed person-days, first, to the reference 108 cohort and were then placed into the NMSC cohort from 109the exact date of the first record of NMSC, in which they 110 then contributed person-days to the NMSC cohort.

111 For each outcome cancer studied (below), we excluded anyone who had a record of that outcome cancer before, or 112at the same time as, their record of the NMSC or reference 113 cohort condition, so as to establish the correct chronology 114 115of events for the investigation of NMSC as a potential risk 116factor for the subsequent outcome cancers.

117 Outcomes

The individuals in the NMSC cohort and reference 118 cohort were then "followed up" by searching the database 119for any subsequent NHS care for, or death from, other 120121primary malignant cancers besides NMSC ("outcome 122cancers"), during the study period. We searched, first, for 123any outcome cancer within the range ICD-10 C00-C41, 124C45–C75, and C81–C97, taking the earliest record of a 125cancer event within that range as the individual's outcome event. We then searched for specific outcor individually within that range (listed in the t ancers

Statistical analysis

Both cohorts were stratified for analysis, by age in 5year groups, sex, calendar year of first recorded admission, region of residence, and quintile of patients' index of multiple deprivation (IMD) score (a measure of socioeconomic status widely used in England). We calculated the observed incidence rate of individuals with each outcome cancer in each stratum of each of the two cohorts, based on person-days of "follow-up." "Date of entry" into each cohort was the date of earliest record of NMSC or any one of the reference cohort conditions. "Date of exit" was the date of the earliest subsequent record for the outcome cancer, death, or the end of data collection period (December 31, 2011).

The indirect method of standardization was used, with the NMSC and reference cohorts taken together as the standard population. For analysis of each outcome cancer, we applied the stratum-specific incidence rates that were observed in the standard population to the number of people in each stratum of the NMSC cohort and then, separately, to those in the reference cohort, to obtain the expected number of people with the outcome cancer in each stratum of the NMSC and reference cohort. Observed (O) and expected (E) numbers for each stratum were then summed to give total observed and expected numbers of people with the outcome cancer in each of the cohorts. We then compared the observed and expected numbers of people with the outcome cancer in the NMSC cohort with those in the reference cohort, using the formula: $(O^{\text{NMSC}}/E^{\text{NMSC}})$: $(O^{\text{REF}}/E^{\text{REF}})$, where the Os and Es are the observed and expected numbers in the NMSC and reference cohorts, respectively. The notation (O^{NMSC}) E^{NMSC}) gives the calculation of relative risk in the NMSC group, relative to the standard population; that of $(O^{\text{REF}}/E^{\text{REF}})$, gives a calculation of relative risk in the reference cohort, relative to the standard population; and we termed the result, $(O^{\text{NMSC}}/E^{\text{NMSC}}) : (O^{\text{REF}}/E^{\text{REF}})$, the "risk ratio (RR)." The RR, its confidence interval, and χ^2 statistics for its significance were calculated using standard statistical methods (14).

We used stratification, rather than matching, for the variables of age, sex, year, region, and deprivation to maximize the availability of the data, statistical power, and reduction of confounding. We did so because there is no merit in discarding data simply to have equal numbers (e.g., 1:1 or 2:1 matching) in each subgroup. To illustrate, in the age group 20 to 24 years there were 828 people in the NMSC cohort, 506,141 in the reference cohort, and 506,969 in the combined "standard" cohort. The stratum-specific rates within the 506,969 people in the standard cohort are then applied to the numbers in each substratum in the NMSC cohort and then the reference cohort. In this way, available data are maximized; statistical power is as high as it can be; there are adequate numbers to populate every 127

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Table 1. Age distribution of people with NMSC in the study, the percentage who were women, and the number and age distribution of people in the reference cohort^a

England (1999–2011)				
Age at admission to the NMSC cohort	Number in the NMSC cohort (% of total)	% of Women	Number in the eference cohort	
0–4	51 (<0.1)	63	589,344	
5–9	119 (<0.1)	60	610,791	
10–14	221 (<0.1)	53	492,236	
15–19	417 (0.1)	56	466,250	
20–24	813 (0.2)	60	506,114	
25–29	1,773 (0.4)	58	517,977	
30–34	3,618 (0.7)	55	498,316	
35–39	6,602 (1.3)	55	507,940	
40–44	11,041 (2.2)	54	487,475	
45–49	15,988 (3.2)	52	458,286	
50–54	23,497 (4.7)	49	469,656	
55–59	33,157 (6.6)	46	499,147	
60–64	47,141 (9.4)	43	514,123	
65–69	58,452 (11.6)	41	499,152	
70–74	70,951 (14.1)	41	501,378	
75–79	80,465 (16)	43	484,323	
80–84	72,375 (14.4)	47	369,574	
85–120	75,809 (15.1)	57	315,431	
All ages	502,490 (100)	46	8,787,513	

^aThe reference cohort consisted of people admitted with the following conditions coded under the Office of Population, Censuses, and Surveys code (OPCS) edition 4 for operations and the ICD revision 10 for diagnoses: adenoidectomy (OPCS4 E20), tonsillectomy (F34 + F36), appendectomy (H01–H03), total hip replacement (W37–W39), total knee replacement (W40–W42), cataract (ICD 10 H25), squint (H49-H51), otitis externa/media (H60-H67), varicose veins (I83), hemorrhoids (I84), upper respiratory tract infections (J00-J06), deflected septum, nasal polyp (J33 + J34.2), impacted tooth and other disorders of teeth (K00-K03), inguinal hernia (K40), in growing toenail and other diseases of nail (L60), sebaceous cyst (L72.1), bunion (M20.1), internal derangement of knee (M23), superficial injury and contusion (S00, S10, S20, S30, S40, S50, S60, S70, S80, and S90), dislocations, sprains and strains (S03, S13, S23, S33, S43, S53, S63, S73, S83, and S93), head injury (S06), and selected limb fractures (S42, S52, S62, S82, and S92).

cell in the potential confounder data; and the resultant 185 186 RRs are adjusted, as much as all available data allow, in respect of age, sex, year, region, and deprivation. 187

188Q5 We carried out subanalyses by the age group and time 189interval, splitting results by age at first admission for the NMSC or reference cohort condition (grouped as <24, 25–44, 45–59, and 60+) by time interval between the 190191 192event of first admission for NMSC (or the reference cohort 193condition) and the event of the subsequent outcome 194cancer (<1, 1–4, and 5+ years).

195Results

196Table 1 shows the number, age, and sex distribution of 197people at entry to the NMSC cohort, and the number and 198age distribution of people in the reference cohort. In total, 199there were 502,490 people in the NMSC cohort and there 200were 8,787,513 people in the reference cohort. The mean period of follow-up in the NMSC cohort was 5.1 years; the 201202mean period of follow-up in the reference cohort was 6.0 203years. Of note, 462 people who started in the reference cohort later crossed over into the NMSC cohort. The RR, 204

comparing the NMSC cohort with the reference cohort, for 206 all outcome cancers combined in people across all ages 207 and all time intervals, was 1.36 (95% CI, 1.35-1.37). When 208we excluded melanoma from the combined outcome 209cancers range, the RR decreased but remained high at 2101.27 (95% CI, 1.26-1.28). We found increased risks in 26 of 21129 individual outcome cancers (P < 0.05). The highest RRs 212were found for salivary gland (5.78 and 5.29-6.32), mel-213anoma (5.54 and 5.37-5.71), bone (2.93 and 2.66-3.23), and 214 upper gastrointestinal tract malignancies (2.36 and 2.25-2152.48). Results for all individual outcome cancers are 216shown in Table 2. NMSC was not found to be protective 217against any cancer, but cervical, uterine, and testicular 218cancers were not significantly elevated (P > 0.05). Any 219differences between males and females were small (Sup-220plementary Table S1). 221

Results by age

Table 3 and Table 4 show RRs for people ages <25, 25–	223
44, 45–59, and 60+ years at the time of entry to their cohort.	224
Generally, RRs for all outcome cancers decreased with	225

Table 2. Analysis by type of cancer: observed numbers of people with NMSC who had a subsequent record of primary malignant cancer, RRs and 95% CIs

	Cancer following admission for NMSC			
Site (ICD-10 code)	0	RR (95% CI)		
Any malignant primary cancer excluding NMSC ^a	67,148	1.36 (1.35–1.37)		
Any malignant primary cancer excluding all skin ^b	62,377	1.27 (1.26–1.28)		
Bladder (C67)	5,372	1.13 (1.10–1.16)		
Bone (C40–C41)	554	2.93 (2.66-3.23)		
Brain (C71)	781	1.07 (1.00–1.16)		
Breast (C50)	6,154	1.24 (1.21–1.28)		
Cervix (C53)	193	0.97 (0.83-1.12)		
Colon (C18–C19)	7,259	1.16 (1.13–1.19)		
GI-upper (C00-C06.8, C09-C10, and C12-C14)	1,974	2.36 (2.25-2.48)		
Kidney (C64–C65)	1,909	1.18 (1.13–1.24)		
Larynx (C32)	649	1.32 (1.21–1.43)		
Leukemia-lymphoid (C91)	2,096	2.01 (1.92-2.11)		
Leukemia-myeloid (C92)	1,173	1.28 (1.21–1.37)		
Liver (C22)	1,095	1.10 (1.03–1.17)		
Lung (C33–C34)	10,942	1.31 (1.28–1.33)		
Lymphoma—Hodgkins (C81)	298	1.68 (1.48–1.89)		
Lymphoma-non-Hodgkin (C82-C85)	3,632	1.63 (1.58–1.69)		
Melanoma-malignant (C43)	6,693	5.54 (5.37-5.71)		
Multiple myeloma (C90)	1,225	1.13 (1.07–1.20)		
Nasopharynx (C11)	98	1.48 (1.19–1.82)		
Nervous system—other (C70)	62	1.91 (1.43-2.51)		
Esophageal (C15)	2,274	1.10 (1.06–1.15)		
Ovary (C56)	898	1.08 (1.01–1.16)		
Pancreas (C25)	1,975	1.05 (1.00–1.10)		
Prostate (C61)	11,730	1.12 (1.10–1.14)		
Rectum (C20–C21)	3,529	1.43 (1.38–1.48)		
Salivary gland (C07–C08)	881	5.78 (5.29-6.32)		
Stomach (C16)	2,205	1.08 (1.04–1.13)		
Testis (C62)	69	1.28 (0.99–1.62)		
Thyroid (C73)	226	1.25 (1.08–1.43)		
Uterus—body of (C54)	830	1.05 (0.98–1.13)		
NOTE: All these ICD codes are specifically for primary maligna ^a C00–C43, C45–C75, and C81–C97.	ncy.			

^bC00–C41, C45–C75, and C81–C97.

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 increasing age but remained elevated throughout. Risk

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 ratios for all outcome cancers combined at ages <25, 25–44,</td>

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 45–59, and 60+, respectively, were 22.6 (95% CI, 18.0–

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 28.2), 3.52 (95% CI, 3.30–3.75), 1.74 (95% CI, 1.70–1.79), and

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 1.32 (95% CI, 1.30–1.33).

23:Q6 Given that only 0.3% of the people in our NMSC cohort 234were under 25 years (Table 1)—a reflection of the fact that 235cancer incidence rates are very low in this age group 236generally-the observed and expected numbers of people in this age group with specific outcome cancers were small (usually less than 3), resulting in the statistical power 237238239and wider confidence interval s. We report, in Table 3, RRs 240 for specific outcome cancers in which the observed or 241expected values in this age group were 3 or more in the 242NMSC cohort. The RR for melanoma and salivary gland cancers were particularly high at 94.4 (95% CI, 65.3–133.0) and 93.4 (95% CI, 18.4–295.1), respectively.

Short- (<1 year), medium- (1–4 years), and long-term (5+ years) associations

Table 5 shows results by time intervals between entry to the NMSC or reference cohort and the first cancer outcome event for each individual cancer. For most outcome cancers, the RRs were highest within the first year and, for most outcome cancers, the RRs decreased but remained significantly high for first cancer events at 1 to 4 years, and for first events after 5+ years. The RRs for several specific outcome cancers—of the bladder, brain, breast, colon, liver, lung, pancreas, prostate, and stomach—remained consistently elevated across all time intervals; some

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Table 3. People younger than 25 years with NMSC who had a subsequent record of another cancer: observed numbers. RRs and 95% CIs Ages 0-24 Site (ICD-10 code) Ο RR (95% CI) Any malignant primary cancer excluding NMSC^a 82 22.64 (17.97-28.16) Any malignant primary cancer excluding all skin^b 48 14.45 (10.63-19.19) Bone (C40-C41) 53.43 (24.15-103.05) 9 Brain (C71) 4 20.18 (5.46-52.27) Leukemia-lymphoid (C91) 5 26.75 (8.65-62.79) Melanoma-malignant (C43) 37 94.41 (65.25-133.01) Salivary gland (C07-C08) 93.44 (18.43-295.10) 3 ^aC00-C43, C45-C75, and C81-C97. ^bC00-C41, C45-C75, and C81-C97.

260 increased with increasing time (brain, colon, and261 prostate).

262 Discussion

263 Principal findings

A previous record of NMSC was associated with an increased subsequent risk of a broad spectrum of other primary malignant cancers in this national cohort study. Younger people with NMSC, in particular, are at much higher risk of subsequent primary malignant cancer incidence than people without NMSC, although cancer incidence is, overall, rarer in younger people.

271 Strengths, weaknesses, and potential biases

With 502,490 NMSC cases, this is the largest study on
this topic so far, with high statistical power and precision
in analyses split by cancer site, age, and time interval.

The main shortcoming is that we were unable to dis-275276tinguish squamous cell carcinomas (SCC) from basal cell carcinomas (BCC) because these cancers are coded togeth-277278er in the ICD. It might be that one type of NMSC is more 279strongly associated with increased risks of subsequent 280primaries; however, only subtle differences have been noted in studies that do differentiate SCCs and BCCs 281(5, 9). T-cell lymphomas should be coded separately, but 282might occasionally be miscoded as NMSC; however, 283given their rarity compared with SCCs and BCCs, any 284effect on our results is likely to be very small. 285

We have not adjusted for multiple comparisons of cancer sites, but because almost all of the RRs go in the same direction of excess risk, it is highly improbable that these are chance findings. The height of some of the RRs, and the ubiquity of elevated risk across so many cancer sites, is such that the associations are very unlikely to be attributable to artefacts of data collection or study design.

In our stratified analysis, we accounted for age, sex, calendar year of first recorded admission, region of residence, and IMD score, but were unable to adjust for other potential confounders like body mass index, smoking, and UV exposure. However, smoking-related cancers,

299 such as lung, had a RR of less than the average elevation of cancer site risk, and, after excluding melanoma from the 300 analyses, the RRs for cancer overall were still significantly 301 high, suggesting that factors other than smoking and UV 302 exposure are at play. Studies that have been able to control 303 for these potentially confounding variables have not 304 found substantial difference between the age-adjusted 305 and multivariable-adjusted risks, demonstrating that the 306 observed association is unlikely to be explained by such 307 factors (9). Furthermore, the effects of acquired risks, such 308 as smoking and other behavioral factors, are cumulative, 309 and one would expect an increasing relative risk with 310 increasing age if they were major factors behind our 311 associations. We found the opposite (higher relative risks 312 at young ages), suggesting a genetic cause or early envi-313 ronmental exposure might explain our results rather than 314 a later acquired cause of association. 315

It was not possible for us to directly ascertain the 316 number of NMSC cases that were missed by our dataset. 317 Most dermatologic excisions in ambulatory care should be 318 captured under hospital day-case admissions (15), but 319 some will be classified as outpatient sand some will be 320 treated wholly in primary care. The people admitted as a 321 day case or inpatient may be at the more severe end of the 322 diseases spectrum. It seems unlikely, however, that the 323 profile of RRs seen in our study is attributable to missing 324 cases of NMSC that are treated in primary care. Further-325 more, the decision to treat a person with NMSC as a day 326 case or inpatient, rather than within primary care or as an 327 outpatient, is perhaps more likely to relate to the prefer-328 ences of the treating clinician than the characteristics of the 329 NMSC. 330

We do not have data on the cohorts' migration into or out of the study area. However, migration bias is unlikely to be an issue because there is no apparent reason why the migration pattern of one cohort would be significantly different from the other.

We examined associations at different time intervals between the first known record of NMSC and the first subsequent record of the outcome cancer. This was partly 331

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Table 4. People ages 25–44, 45–64, and 65+ with NMSC who had a subsequent record of another cancer:observed numbers, RRs, and 95% CIs

	Ages 25–44		Ages 45–59		Ages 60+	
Site (ICD-10 code)	0	RR (95% CI)	0	RR (95% CI)	0	RR (95% CI)
Any malignant primary cancer excluding NMSC ^a	1,001	3.52 (3.30–3.75)	5,787	1.74 (1.70–1.79)	60,278	1.32 (1.30–1.33)
Any malignant primary cancer excluding all skin ^b	674	2.47 (2.29–2.67)	4,951	1.52 (1.47–1.56)	56,704	1.25 (1.24–1.26)
Bladder (C67)	20	2.30 (1.39–3.58)	213	1.12 (0.97–1.29)	5,139	1.13 (1.10–1.16)
Bone (C40–C41)	32	12.19 (8.13–17.73)	84	4.38 (3.42–5.56)	429	2.66 (2.37–2.98)
Brain (C71)	24	2.03 (1.29–3.04)	92	1.36 (1.09–1.69)	661	1.02 (0.93–1.10)
Breast (C50)	174	1.79 (1.53–2.08)	853	1.25 (1.17–1.34)	5,127	1.23 (1.20–1.27)
Cervix (C53)	12	1.31 (0.67–2.30)	23	0.99 (0.62–1.50)	157	0.94 (0.79–1.10)
Colon (C18–C19)	21	1.20 (0.74–1.84)	395	1.30 (1.17–1.44)	6,841	1.15 (1.12–1.18)
GI-upper (C00-C06.8, C09-C10, and C12-C14)	52	4.42 (3.28-5.86)	354	2.62 (2.34–2.93)	1,567	2.34 (2.21–2.48)
Kidney (C64–C65)	14	1.59 (0.86–2.68)	148	1.32 (1.11–1.56)	1,747	1.17 (1.11–1.23)
Larynx (C32)	2	0.81 (0.10–2.99)	85	1.57 (1.24–1.96)	562	1.30 (1.18–1.42)
Leukemia-lymphoid (C91)	10	2.97 (1.41–5.55)	90	1.73 (1.38–2.15)	1,991	2.06 (1.96–2.17)
Leukemia-myeloid (C92)	10	2.05 (0.98–3.82)	59	1.31 (0.99–1.70)	1,103	1.28 (1.20–1.37)
Liver (C22)	5	1.53 (0.49–3.62)	81	1.32 (1.04–1.65)	1,009	1.08 (1.01–1.16)
Lung (C33—C34)	34	1.95 (1.35–2.75)	787	1.74 (1.61–1.87)	10,121	1.28 (1.26–1.31)
Lymphoma—Hodgkins (C81)	10	1.87 (0.89–3.47)	52	2.16 (1.59–2.87)	236	1.65 (1.42–1.90)
Lymphoma—non-Hodgkin (C82–C85)	28	1.80 (1.19–2.61)	318	1.90 (1.69–2.14)	3,285	1.63 (1.57–1.69)
Melanoma-malignant (C43)	377	20.63 (18.36–23.13)	1,040	9.40 (8.72–10.13)	5,239	5.08 (4.90-5.27)
Multiple myeloma (C90)	4	1.33 (0.36–3.46)	65	1.14 (0.87–1.46)	1,156	1.13 (1.06–1.21)
Nasopharynx (C11)	1	0.62 (0.02–3.52)	28	2.28 (1.49–3.38)	69	1.33 (1.01–1.72)
Nervous system—other (C70)	3	7.85 (1.53–25.33)	5	1.67 (0.52–4.13)	53	1.84 (1.33–2.51)
Esophageal (C15)	12	1.98 (1.01–3.49)	157	1.19 (1.01–1.40)	2,105	1.10 (1.05–1.15)
Ovary (C56)	16	1.54 (0.88–2.53)	92	1.06 (0.85–1.31)	788	1.08 (1.00–1.16)
Pancreas (C25)	9	2.24 (1.01–4.32)	103	1.07 (0.87–1.31)	1,862	1.05 (1.00–1.10)
Prostate (C61)	9	1.72 (0.78–3.29)	493	1.21 (1.10–1.32)	11,228	1.12 (1.09–1.14)
Rectum (C20–C21)	105	10.68 (8.60–13.15)	423	2.59 (2.33–2.87)	2,999	1.29 (1.24–1.35)
Salivary gland (C07–C08)	19	12.34 (7.20–20.05)	70	5.98 (4.51–7.84)	789	6.33 (5.71–7.02)
Stomach (C16)	13	2.70 (1.43–4.68)	89	1.08 (0.86–1.34)	2,102	1.08 (1.03–1.13)
Testis (C62)	7	1.00 (0.40–2.07)	18	1.58 (0.92–2.54)	43	1.26 (0.89–1.76)
Thyroid (C73)	10	1.30 (0.62–2.40)	36	1.44 (0.99–2.02)	179	1.22 (1.03–1.42)
Uterus—body of (C54)	10	1.58 (0.75–2.95)	105	1.06 (0.86–1.29)	715	1.05 (0.97–1.13)
^a C00–C43, C45–C75, and C81–C97.						

^bC00–C41, C45–C75, and C81–C97.

341 to judge any effect of surveillance bias. This would arise if, 342 for example, care for NMSC led to the prompt identifi-343 cation of another cancer. For all primary outcome cancers combined, the RR decreased after the first year, but 344 345remained significantly high. Increased surveillance might 346 partially account for the increased rates of melanoma, but is unlikely to account for the increased rates of other 347 nonvisible internal cancers, particularly in which the 348349 increased risk continues beyond 5 years after the diagno-350sis of NMSC, as was the case with most outcome cancers. 351

We cannot rule out the possibility of some misclassification of secondary cancer sites as primary in the source data, although we selected ICD codes for primary cancers only. We are unable to link our records to histologic samples. If misclassification of secondary cancers were an explanation for the pattern of our results, the effects of it would have to be not only considerable, but also longterm because the RRs for most cancers remain significantly increased after 5 years. The incidence of metastasis in people with BCC is extremely rare, with reported incidences ranging from 0.0028% to 0.5% (16).

Treatment and mortality bias is likely to be negligible, as almost all NMSCs are treated curatively by excision without the use of potentially carcinogenic radiotherapy or systemic chemotherapy (17). Despite the high incidence of NMSC, mortality rates from NMSC are low at 0.91 per 100,000 persons per year (18).

Comparison with other studies

A meta-analysis that combined the results of three cohort studies (5) and accounted for individual level risk factors, like smoking, showed an overall summary

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Table 5. People who were admitted with NMSC and had a subsequent record of another cancer, by time interval between admissions (<1 year, 1–4 years, and 5 or more years): observed numbers, RRs, and 95% Cls

	<1 y	ear time interval	1–4 yea	ars time interval	5+ yea	ars time interval
Cancer site (ICD-10 code)	0	RR (95% CI)	0	RR (95% CI)	0	RR (95% CI)
Any malignant primary cancer excluding NMSC ^a	16,485	1.64 (1.61–1.67)	33,304	1.30 (1.29–1.32)	17,359	1.27 (1.25–1.29)
Any malignant primary cancer excluding all skin ^b	14,104	1.41 (1.38–1.43)	31,667	1.25 (1.23–1.26)	16,606	1.22 (1.20–1.24)
Bladder (C67)	1,087	1.08 (1.01–1.16)	2,792	1.14 (1.10–1.19)	1,493	1.14 (1.08–1.20)
Bone (C40–C41)	264	5.37 (4.59–6.27)	211	2.37 (2.03–2.76)	79	1.66 (1.30-2.10)
Brain (C71)	128	0.85 (0.70–1.02)	371	1.05 (0.94–1.17)	282	1.25 (1.11–1.42)
Breast (C50)	1,228	1.31 (1.23–1.39)	3,075	1.20 (1.15–1.24)	1,851	1.29 (1.23–1.35)
Cervix (C53)	38	1.08 (0.76–1.51)	110	1.05 (0.86–1.28)	45	0.75 (0.55-1.01)
Colon (C18–C19)	1,376	1.07 (1.01–1.13)	3,817	1.18 (1.14–1.22)	2,066	1.19 (1.14–1.25)
GI-upper (C00-C06.8, C09-C10, and C12-C14)	713	3.45 (3.15–3.76)	827	2.05 (1.90-2.21)	434	1.98 (1.78–2.19)
Kidney (C64–C65)	366	1.22 (1.08–1.36)	976	1.17 (1.09–1.25)	567	1.18 (1.08–1.29)
Larynx (C32)	161	1.40 (1.17–1.66)	321	1.29 (1.15–1.46)	167	1.31 (1.11–1.54)
Leukemia—lymphoid (C91)	556	2.58 (2.33–2.85)	1,083	2.09 (1.95–2.24)	457	1.53 (1.38–1.69)
Leukemia-myeloid (C92)	234	1.28 (1.11–1.47)	620	1.31 (1.20–1.43)	319	1.24 (1.10–1.39)
Liver (C22)	194	1.04 (0.88–1.21)	565	1.11 (1.02–1.22)	336	1.12 (1.00–1.25)
Lung (C33–C34)	2,166	1.30 (1.24–1.37)	5,723	1.32 (1.29–1.36)	3,053	1.29 (1.24–1.34)
Lymphoma—Hodgkins (C81)	74	1.84 (1.42–2.35)	159	1.71 (1.44–2.02)	65	1.49 (1.14–1.93)
Lymphoma—non-Hodgkin (C82–C85)	878	1.85 (1.71–1.99)	1,842	1.64 (1.56–1.73)	912	1.48 (1.38–1.58)
Melanoma-malignant (C43)	2,912	12.42 (11.69–13.20)	2,553	4.26 (4.06–4.47)	1,228	3.72 (3.49–3.97)
Multiple myeloma (C90)	234	1.03 (0.89–1.18)	659	1.19 (1.09–1.29)	332	1.11 (0.99–1.24)
Nasopharynx (C11)	38	1.51 (1.05–2.12)	39	1.34 (0.93–1.87)	21	1.73 (1.04–2.74)
Nervous system—other (C70)	12	1.85 (0.90–3.51)	35	2.08 (1.39–3.02)	15	1.66 (0.90-2.86)
Esophageal (C15)	464	1.20 (1.08–1.33)	1,186	1.08 (1.01–1.14)	624	1.10 (1.01–1.19)
Ovary (C56)	152	0.98 (0.82–1.16)	474	1.08 (0.98–1.19)	272	1.15 (1.01–1.30)
Pancreas (C25)	348	1.02 (0.91–1.14)	1,027	1.03 (0.96–1.10)	600	1.11 (1.02–1.21)
Prostate (C61)	2,212	1.03 (0.98–1.08)	6,080	1.14 (1.11–1.17)	3,438	1.15 (1.11–1.19)
Rectum (C20–C21)	1,194	2.35 (2.20–2.51)	1,537	1.19 (1.13–1.26)	798	1.20 (1.11–1.29)
Salivary gland (C07–C08)	290	8.74 (7.27–10.52)	443	5.99 (5.26-6.81)	148	3.71 (3.05-4.49)
Stomach (C16)	434	1.08 (0.97–1.20)	1,149	1.06 (1.00–1.13)	622	1.12 (1.03–1.22)
Testis (C62)	22	1.55 (0.96–2.39)	32	1.23 (0.83–1.75)	15	1.10 (0.61–1.82)
Thyroid (C73)	66	1.90 (1.44–2.48)	114	1.19 (0.97–1.45)	46	0.92 (0.67-1.24)
Uterus—body of (C54)	140	0.96 (0.80–1.15)	456	1.11 (1.00–1.22)	234	1.01 (0.88–1.15)
^a C00–C43, C45–C75, and C81–C97.						

^bC00–C41, C45–C75, and C81–C9.

relative risk of cancer after NMSC of 1.49 (95% CI, 1.12–
1.98), similar to our RR of 1.36 (95% CI, 1.35–1.37), notwithstanding differences in methodology. Our results
also corroborate a particularly high risk of melanoma and
salivary gland cancer, found by others (5, 11).
Risks of breast, colorectal, and prostate cancers have

Risks of breast, colorectal, and prostate cancers have
been reported as low in people with NMSC (11, 19, 20). It
has been suggested that increased sunlight exposure and
vitamin D levels play a protective role in their development. We did not find low risks for these cancers.

385Other subgroups known to have high risk of cancer are386transplant patients. However, although SCCs are up to 65-387fold more prevalent in transplant patients than matched388controls (21), the percentage of people with NMSC in our389study who have had a transplant is likely to be very small.

The risk and pattern of cancers in people who have391undergone transplantation, in whom the excess cancer392risk particularly affects the kidney, liver, and non-Hodg-393kin lymphoma (22), would not in itself account for our394results. Findings of a recent study also showed that, even395among transplant recipients, SCC was a marker of396increased risk for other cancers (23).397

Interpretation and implications

Mechanisms for these associations remain elusive. It is399plausible that UV-induced oxidative damage resulting in400gene mutation, immunosuppression, and inflammation401may also act more systemically to increase the risk of402cancer in predisposed individuals in other sites, such as403in immunosuppressed transplant recipients (24, 25) and404

407 those with cancer-predisposing syndromes (26, 27). 408 Recent studies suggest that genetic predisposition to 409 reduced DNA repair capacity may be an underlying 410 susceptibility factor for NMSC and other cancers (28-411 31). We considered that the occurrence of NMSC in people 412in our study ages <25 might be associated with certain 413congenital skin disorders, in particular xeroderma pig-414mentosum. We, therefore, conducted a post-hoc analysis to 415ascertain the number of people in this age group of the 416 NMSC cohort who also had a record of xeroderma pigmentosum either before or after their record of NMSC. We 417 found a total of 6, 5 of whom had a record of xeroderma 418 419 pigmentosum before the record of NMSC. In a cohort of 420 1,621 people with NMSC ages <25, the occurrence of 6 421cases of xeroderma pigmentosum is unlikely to be of 422 significant impact.

423 Our findings should be regarded as supporting the 424 hypothesis of a raised risk of other cancers after diagnosis 425of NMSC, but not as definitive. The results represent what 426can be done using very large, linked, routinely collected 427administrative datasets; but such datasets lack detail. Alternative designs of similar scope, which could incor-428429porate data on genetic profiling and biomarkers, would be 430substantial undertakings, but the benefits of more precise 431characterization of those with NMSC who are at risk of 432other specific malignancies would be considerable.

433 For those cancers in which screening, or other app-434roaches to cancer prevention, has proven benefit, a next step would be to define guidelines that translate these 435436results into clinical practice and lifestyle advice, especially 437for younger individuals with NMSC. Guidelines on 438NMSC do not refer to surveillance for any specific cancer 439types apart from other skin cancers (32, 33); our results 440 suggest surveillance for other cancers might be warranted 441 if supported by further clinical research and cost-benefit 442 analyses. 443

Further work to elucidate why people with NMSC, particularly the young, are at increased risk of other

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- Fransen M, Karahalios A, Sharma N, English DR, Giles G, Sinclair RD, 1. Non-melanoma skin cancer in Australia, Med J Aust 2012;197;565-8.
- Dipegen TL, Mahler V. The epidemiology of skin cancer. Br J Dermatol 2. 2002;146 Suppl 61:1-6.
- З. Halliday G. Inflammation, gene mutation and photoimmunosuppression in response to UVR-induced oxidative damage contributes to photocarcinogenesis. Mut Res 2005;571:107-20.
- Marcil I, Stern RS. Risk of developing a subsequent nonmelanoma skin 4. cancer in patients with a history of nonmelanoma skin cancer: a critical review of the literature and meta-analysis. Arch Dermatol 2000;136: 1524-30.
- 5. Wheless J, Black J, Alberg AJ. Nonmelanoma skin cancer and the risk of second primary cancers: a systematic review. Cancer Epidemiol Biomarkers Prev 2010:19:1686-95
- Maitra SK, Gallo H, Rowland-Payne C, Robinson D, Møller H. Second 6. primary cancers in patients with squamous cell carcinoma of the skin. Br J Cancer 2005;92:570-1.
- 7. Cantwell MM, Murray LJ, Catney D, Donnelly D, Autier P, Boniol M, et al. Second primary cancers in patients with skin cancer: a population-based study in Northern Ireland, Br J Cancer 2009:100:174-7.

malignancies could be an important step to a more fun- damental understanding of carcinogenesis.	$\begin{array}{c} 446 \\ 447 \end{array}$
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Ethical Approval

Ethical approval for analysis of the record-linkage study data were obtained from the Central and South Bristol Multi-Centre Research Ethics Committee (04/O2006/176)

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- Milán T. Pukkala E. Verkasalo PK. Kaprio J. Jansén CT. Koskenvuo M. 8. et al. Subsequent primary cancers after basal-cell carcinoma: a nationwide study in Finland from 1953 to 1995. Int J Cancer 2000;87:283-8.
- Song F, Qureshi AA, Giovannucci EL, Fuchs CS, Chen WY, Stampfer MJ. et al. Risk of a second primary cancer after non-melanoma skin cancer in white men and women: a prospective cohort study. PLoS Med 2013:10:e1001433.
- 10. Bower CPR, Lear J, Bygrave S, Etherington D, Harvey I, Archer CB. Basal cell carcinoma and risk of subsequent malignancies: a cancer registrybased study in southwest England. J Am Acad Dermatol 2000;42:988-91.
- 11. Grant WB. A meta-analysis of second cancers after a diagnosis of nonmelanoma skin cancer: additional evidence that solar ultraviolet-B irradiance reduces the risk of internal cancers. J Steroid Biochem Mol Biol 2007:103:668-74.
- 12. Chen J, Ruczinski I, Jorgensen TJ, Yenokyan G, Yao Y, Alani R, et al. Nonmelanoma skin cancer and risk for subsequent malignancy. J Natl Cancer Inst 2008;100:1215-22
- 13. Goldacre M, Kurina L, Yeates D, Seagroatt V, Gill L. Use of large medical databases to study associations between diseases. QJM 2000:93:669-75.

- Breslow NE, Day NE. Statistical methods in cancer research. Volume II-the design and analysis of cohort studies. IARC Sci Publ 1987;82: 1–406.
- Milkeljevic J, Johnston C, Adamson PJ, Wright A, Bishop JA, Batman P, et al. How complete has skin cancer registration been in the UK? A study from Yorkshire. Eur J Cancer Prev 2003;12: 125–33.
- Lo JS, Snow SN, Reizner GT, Mohs FE, Larson PO, Hruza GJ. Metastatic basal cell carcinoma: report of twelve cases with a review of the literature. J Am Acad Dermatol 1991;24:715–9.
- Neville JA, Welch E, Leffell DJ. Management of nonmelanoma skin cancer in 2007. Nat Clin Pract Oncol 2007;4:462–9.
- Lewis KG, Weinstock MA. Non melanoma skin cancer mortality (1988– 2000): the Rhode Island follow–back study. Arch Dermatol 2004;140: 837–42.
- De Vries E, Soerjomataram I, Houterman S, Louwman MW, Coeberg JW. Decreased risk of prostate cancer after skin cancer diagnosis: a protective role of ultraviolet radiation? Am J Epidemiol 2007;165: 966–72.
- **20.** Gorham ED, Garland CF, Garland FC, Grant WB, Mohr SB, Lipkin M, et al. Optimal vitamin D status for colorectal cancer prevention: a quantitative meta analysis. Am J Prev Med 2007;32:210–6.
- Jensen P, Hansen S, Møller B, Leivestad T, Pfeffer P, Geiran O, et al. Skin cancer in kidney and heart transplant recipients and different long-term immunosuppressive therapy regimens. J Am Acad Dermatol 1999;40:177–86.
- Engels EA, Pfeiffer RM, Fraumeni JF Jr, Kasiske BL, Israni AK, Snyder JJ, et al. Spectrum of cancer risk among US solid organ transplant recipients. JAMA 2011;306:1891–901.
- 23. Wisgerhof HC, Wolterbeek R, de Fijter JW, Willemze R, Bouwes Bavinck JN. Kidney transplant recipients with cutaneous squamous

cell carcinoma have an increased risk of internal malignancy. J Invest Dermatol 2012;132:2176-83. 559

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588

- Glover MT, Deeks JJ, Raftery MJ, Cunningham J, Leigh IM. Immunosuppression and risk of non-melanoma skin cancer in renal transplant recipients. Lancet 1997:249:398.
- Berg D, Otley CC. Skin cancer in organ transplant recipients: epidemiology, pathogenesis, and management. J Am Acad Dermatol 2002;47:1–17.
- **26.** Nikolaou V, Stratigos AJ, Tsao H. Heridatary nonmelanoma skin cancer. Semin Cutan Med Surg 2012;31:204–10.
- Ponti G, Pellacani G, Seidenari S, Pollio A, Muscatello U, Tomasi A. Cancer-associated genodermatoses: skin neoplasms as clues to hereditary tumor syndromes. Crit Rev Oncol Hematol 2013;85:239–56.
- Brewster AM, Alberg AJ, Strickland PT, Hoffman SC, Helzlsouer K. XPD polymorphism and risk of subsequent cancer in individuals with nonmelanoma skin cancer. Cancer Epidemiol Biomarkers Prev 2005; 13:1271–5.
- Li C, Wang LE, Wei Q. DNA repair phenotype and cancer susceptibility-a mini review. Int J Cancer 2009;124:999–1007.
- **30.** Ruczinski I, Jorgensen TJ, Shugart YY, Schaad YB, Kessing B, Hoffman-Bolton J, et al. A population-based study of DNA repair gene variants in relation to nonmelanoma skin cancer as a marker of a cancer-prone phenotype. Carcinogenesis 2012;33:1692–8.
- Kitagishi Y, Kobayashi M, Matsuda S. Defective DNA repair systems and the development of breast and prostate cancer (review). Int J Oncol 2013;42:29–34.
- **32.** Drake LA, Ceilley RI, Cornelison RL, Dobes WA, Dorner W, Goltz RW, et al. Guidelines of care for cutaenous squamous cell carcinoma. J Am Acad Dematol 1993;28:628–31.
- Alam M, Ratner D. Cutaneous squamous–cell carcinoma. N Engl J Med 2001;344:975–83.

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