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Clinicopathologic features of incident and subsequent tumors in patients with multiple primary cutaneous melanomas

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Abstract

Background—0.6–12.7% of patients with primary cutaneous melanoma will develop additional melanomas. Pathologic features of tumors in patients with multiple primary cutaneous melanomas have not been well described. In this large international multi-center case-control study, we compared the clinicopathologic features of a subsequent melanoma with the preceding (usually the first) melanoma in patients with multiple primary cutaneous melanomas, and with those of melanomas in patients with single primary cutaneous melanomas.

Methods—Multiple primary melanoma (cases) and single primary invasive melanoma (controls) patients from the Genes, Environment and Melanoma (GEM) study were included if their tumors were available for pathologic review and confirmed as melanoma. Clinicopathologic

characteristics of invasive subsequent and first melanomas in cases and invasive single melanomas in controls were compared.

Results—473 pairs comprising a subsequent and a first melanoma and 1989 single melanomas were reviewed. Forward stepwise regression modeling in 395 pairs with complete data showed that, compared to first melanomas, subsequent melanomas were: more commonly contiguous with a dysplastic nevus; more prevalent on the head/neck and legs than other sites; and thinner. Compared with single primary melanomas, subsequent melanomas were also more likely to be: associated with a contiguous dysplastic nevus; more prevalent on the head/neck and legs; and thinner. The same differences were observed when subsequent melanomas were compared with single melanomas. First melanomas were more likely than single melanomas to have associated solar elastosis and no observed mitoses.

Conclusions—Thinner subsequent than first melanomas suggest earlier diagnosis, perhaps due to closer clinical scrutiny. The association of subsequent melanomas with dysplastic nevi is consistent with the latter being risk factors or risk markers for melanoma.

Keywords

Diagnosis; Melanoma; Multiple primary melanoma; Pathology; Risk factors

INTRODUCTION

In patients with primary cutaneous melanoma (hereafter referred to as melanoma), the reported incidence of a second or higher order melanoma ranges between 0.6% and 12.7%.¹⁻³ In patients with one melanoma, a family history of melanoma confers an increased risk for a subsequent melanoma.^{2, 4-12} Many studies based on data from single centers, and a few based on population data, have shown that patients with non-familial melanoma also have an increased risk of developing additional melanomas.^{1-5, 7-10, 13-33} The relative risk of additional melanomas is highest in the first year after diagnosis of the first melanoma and decreases progressively with time, but is increased for at least 20 years.^{1, 25}

Most studies of multiple melanomas have detailed the clinical and genetic characteristics of affected patients, but pathologic features of tumors in patients with MPMs have been incompletely described. To date, the pathologic features assessed have been limited to thickness and Clark level,^{8, 34} associated dysplastic nevus,^{2, 17} and, rarely, ulceration^{2, 21} and regression.¹⁶ A detailed comparison of the pathologic differences between tumors in patients with multiple primary melanomas has not been reported.

In the Genes, Environment and Melanoma (GEM) Study, a large international multi-institution case-control study, we sought to examine in detail the clinicopathologic features of melanomas in patients with multiple primary melanomas (GEM cases) and to contrast features of the subsequent melanoma with those of the preceding, usually the first, melanoma, in these patients. Furthermore, we compared the clinicopathologic features of both the subsequent and first melanoma in cases with features of the melanoma in patients with single primary melanomas (GEM controls). The aim of the study was to determine whether there were significant differences in clinicopathologic features between these groups.

METHODS

Patients and selection criteria

The Genes, Environment and Melanoma (GEM) Study³⁵ has investigated genetic and environmental risk factors in cutaneous melanoma patients with newly incident multiple and single primary tumors identified in eight population-based cancer registries in Australia, Canada, Italy and the United States and one hospital center estimated to cover around 50% of the melanomas diagnosed in the state of Michigan. The GEM study is designed as a case control study in which the population at risk is survivors of a first primary melanoma. Controls were diagnosed with a first invasive primary melanoma in 2000 and cases had a newly incident second or higher order invasive or in situ melanoma in 2000–2003 and, in four centers, in 1998 and 1999 also. We used incident sampling to identify both GEM cases and GEM controls and also ascertained the preceding (usually the first) melanoma in GEM cases in local cancer registry records (average 47 months between the first and the subsequent, case-defining melanoma). Further details of the study design and its rationale have been published.³⁵

The Study was approved by the appropriate institutional review committees and met the guidelines of the responsible governmental agencies. For the analyses in this report, GEM Study participants with single primary melanomas (controls) or multiple primary melanomas (cases) were included if their tumors were confirmed as invasive melanoma and were available for pathologic review by an experienced dermatopathologist. Patients whose tumors were entirely in situ (some subsequent melanomas in GEM cases) were excluded.

In cases, clinicopathologic characteristics of the newly incident subsequent melanoma and their preceding, usually first, melanoma were compared; we refer to these melanomas as ‘subsequent’ and ‘first’ for convenience. The order of diagnosis of the multiple melanomas could not be determined for some cases because diagnosis month and year only were obtained for the study; thus multiple melanomas with the same month of diagnosis were treated as simultaneously diagnosed and excluded from this comparison. There was a shorter time between diagnosis of the subsequent and first melanoma in cases in this analysis (mean 47 months, interquartile range (IQR) 10–72) than in all GEM cases (mean 72 months, IQR 14–110); this is probably explained by the greater difficulty of gaining access to pathology sections or slides suitable for review of first melanomas diagnosed longer ago. In separate analyses, clinicopathologic features of the subsequent and first melanomas in cases were compared with the single melanomas in controls. The clinicopathologic characteristics assessed in each melanoma are listed in Tables 1 and 2.

Statistical methods

Our primary analyses compared the clinicopathologic characteristics of subsequent and first melanomas in individual cases by estimating prevalence ratios (subsequent compared with first melanoma) using Cox regression models with equal survival times given to each subject and robust variance. This method gives unbiased effect estimates and confidence intervals and is superior to the use of logistic regression, in which odds ratio estimates of prevalence ratios are non-conservatively biased.³⁶ Analyses were clustered on individual cases to account for within case correlation and adjusted for age at diagnosis, sex and study center. The variables ‘site of melanoma’ and ‘presence of mitoses’ were included in the multivariable models instead of the variables ‘location on exposed sites’ and ‘mitotic rate’ due to their smaller p-values in the univariate analysis. A p-value of <0.2 in the single variable analyses was used as a cut-off for entry into a forward stepwise multiple regression model of all clinicopathologic characteristics. The analyses comparing each of subsequent and first melanomas with single melanomas also estimated prevalence ratios in models

adjusted for age, sex and study center but did not cluster on individual patients; a forward stepwise regression approach was again used. Only patients with valid values for all pathology variables were included in the stepwise analyses.

RESULTS

There were 3,676 patients in the GEM study population, of whom 2,470 had a single primary melanoma and 1,210 had multiple primary melanomas. Following pathology review, 1,214 (33.0%) patients were excluded because pathology slides were not available for review, primary tumor was missing from the slides available or the review diagnosis was missing (866, 23.6%); or slide review classified the index tumor as 'not melanoma' (67, 1.8%) or 'in situ melanoma' (281, 7.6%). Totals of 473 cases with invasive multiple (subsequent and first) melanomas and 1989 controls with single melanomas were available for the analysis; 78 of the former were excluded, however, because they had the same month of diagnosis and were considered to have been diagnosed simultaneously. The age and sex distribution of the included patients (Table 1) was similar to all GEM participants. In the main GEM study, cases were generally older than GEM controls and the female/male ratio fell with age, reflecting the higher rate of increase in melanoma incidence with age in men than in women.³⁵ More than half (58%) of the 473 patients with multiple melanomas in this report had their most recent melanoma diagnosed within 3 years of the preceding melanoma, compared with 52% of all 1,210 GEM multiple melanoma patients.

There were a number of sizeable differences in clinicopathologic characteristics between the subsequent and first melanomas (Table 2). The subsequent melanomas were more often located on the head and neck than MPM1 (23% vs. 16%), though the trunk was the dominant site for both subsequent and first tumors; 50.2% of the subsequent tumors in males and 24.6% in females were on the trunk (Table 1). The subsequent melanomas were more often lentigo maligna melanoma histologic subtype (23% vs. 14%), though 57% of pairs were of the same histologic type, they were thinner (77% vs. 68% ≤ 1.0 mm thick), more often Clark level II (60% vs. 48%) or were lacking vertical growth (57% vs. 46%). A higher proportion of subsequent than first melanomas was associated with remnants of a contiguous dysplastic nevus (24% vs. 15%) and more often had an associated in situ component (92% vs. 89%), marked solar elastosis (24% vs. 20%), mitoses rated as absent (66% vs. 63%), and occurred on a usually exposed site (29% vs. 26%). Adjusting for age at diagnosis did not produce materially different results from those in Table 2. Generally speaking, differences between MPM2 and single primary melanoma were similar to but greater than those between MPM2 and MPM1 (Table 2).

There were no material differences between subsequent melanomas diagnosed within 1–3 months of the first melanoma (synchronous melanomas) and those diagnosed 3 months or more after it (metachronous melanomas). However, when we compared melanoma pairs with <3 years and ≥ 3 years between tumours, the subsequent melanomas diagnosed in the later interval were more likely than those diagnosed in the earlier interval to be of nodular and unclassified type (20% vs 8%), >1mm thick (31% vs 16%), Clark level IV or V (46% vs 34%) and to have mitoses (39% vs 28%) ($p < 0.05$ in each case; results not shown).

Apart from the higher prevalence of subsequent melanomas on the head and neck, the site distributions of subsequent and first melanomas were similar. There was a weak degree of concordance within pairs by body site (46%; Kappa 0.22, 95% CI: 0.21–0.35; $p < 0.001$).

Only a co-existing nevus, location on the head and neck or legs and less deep invasion (whether assessed by Breslow thickness, Clark level or presence of vertical growth) were identified as independently more likely to be present in subsequent melanomas than in first

melanomas in the forward stepwise regression model (which included adjustment for age at diagnosis, sex and center, Table 3). The model fit was marginally poorer when either Clark level or Breslow thickness was substituted for vertical growth (results not shown). There were no differences between the directions of the associations of these variables with MPM2 in the multivariable model (Table 3) and those in the single variable analyses (Table 2).

Only two characteristics appeared important in a forward stepwise regression model (adjusted for age, sex and center) comparing first melanomas in multiple melanoma patients (cases) with single melanomas (controls): mitoses were present less often (prevalence ratio (PR) 0.79, 95% CI 0.65–0.96) in first melanomas than in single melanomas and mild/moderate (PR 1.43, 95% CI 1.11–1.84) and marked solar elastosis (PR 1.41, 95% CI 1.04–1.93) was present more often (results not shown). In a similar analysis comparing subsequent melanomas with single melanomas, having a contiguous dysplastic nevus (PR 1.45, 95% CI 1.15–1.83), less frequent vertical growth (PR 0.79, 95% CI 0.63–0.99) and location on the head and neck (reference category, PR 1.00) or legs (PR 1.05, 95% CI 0.76–1.46) were associated more often with subsequent melanomas than single melanomas (results not shown). These results are similar to those for the comparison of subsequent and first melanomas (Table 3).

DISCUSSION

We examined many histopathologic and some clinical characteristics of multiple melanomas and found that subsequent melanomas in patients with multiple melanomas were more likely than their first melanomas to have evidence of a contiguous dysplastic nevus, to have invaded less, and to be located on the head and neck or legs.

In studying patients with multiple melanomas, it is important to ensure that the subsequent melanomas are independent primary tumors and not cutaneous metastases from an antecedent primary melanoma. Classifying cutaneous melanomas as primary or secondary solely on histologic grounds may be challenging.³⁷ More reliable classification is based on correlation of several clinical and pathologic features, such as location, the presence of an associated precursor/in-situ lesion, lymphatic invasion and dense lymphocytic inflammation, although both primary and metastatic melanomas may share some of these characteristics. In a recent study, Orlov and colleagues³⁸ compared the somatic mutational profiles of pairs of melanomas designated as independent primary tumors on the basis of their clinical and pathologic characteristics. They found no significant evidence of clonal origin of the two primaries in 17 of the 19 patients examined by molecular profiling using a set of highly polymorphic genetic markers. These results suggest that most second melanomas designated clinically and pathologically as independent primary tumors are indeed independent occurrences of the disease, supporting the validity of the criteria used by experienced clinicians and pathologists in distinguishing new primaries from metastases.

Pathologic features of tumors in patients with multiple melanomas have been incompletely reported to date. Most commonly, subsequent melanomas have been reported to have invaded less than preceding melanomas, both in terms of Breslow thickness^{20–22} and Clark level.^{8, 27} Studies that included in situ melanomas reported that a greater proportion of subsequent melanomas were in situ.^{2, 3, 20, 21, 29, 39} Although melanomas that were exclusively in situ were excluded from the present analysis, invasive subsequent melanomas were more commonly associated with an in situ component (92%) than preceding melanomas (89%) and single melanomas (88%). The high prevalence of an associated in situ component supports the proposition that the subsequent melanomas in patients with multiple melanomas in this study were primary at the site of diagnosis.

We observed a weak but significant concordance between the sites of occurrence of multiple melanoma pairs. While a significant correlation between their sites has not been reported in most studies,^{8, 10, 14, 16, 20, 22, 40} several studies^{19, 34, 39} did find site concordance ranging between 52% and 56%. Some degree of site concordance between melanomas in the same patient would be expected because the same patterns of sun exposure and sun protection underlie the occurrence of both lesions. The comparatively weak site concordance and the fact that subsequent melanomas are diagnosed synchronously (within three months) with the first melanoma in up to 60% of patients with multiple melanomas (27% of the 395 pairs in the present study – results not shown)²¹ highlight the need for careful and complete skin examination when assessing patients with melanoma.^{39, 41, 42} Moreover, the increased risk of metachronous melanomas, the long intervals (>20 years¹) within which they may be diagnosed, and the probable benefits of early diagnosis of additional melanomas and metastatic disease suggest that patients with one melanoma may benefit from regular follow up.^{27, 43} This suggestion is strengthened by our findings that subsequent melanomas diagnosed ≥ 3 years after a first, when perhaps clinical follow-up has become less intense or ceased, were thicker, more likely to be of nodular type and to have mitoses, than subsequent melanomas diagnosed within 3 years of the first.

In addition to the greater site concordance of multiple melanomas, subsequent melanomas were more likely than first melanomas to be on the head and neck or legs than on the upper limbs or trunk. This too might be expected, at least for the head and neck, because of the head and neck's generally greater exposure to the sun and greater risk of melanoma per unit of surface area than other body sites, particularly in older people.^{44, 45} This possible association of multiple melanomas with higher sun exposure is supported by the stronger association of solar elastosis with first melanomas than single melanomas in the present study, observed associations of high sun exposure⁴⁴ and lack of sunscreen use⁴ with an increased risk of multiple melanomas, and the finding that solar exposure at any age was associated with increased risk of developing multiple primary melanomas in the GEM study.⁴⁶

Superficial spreading melanoma, lentigo maligna melanoma and nodular melanoma were, in decreasing order, the commonest histologic subtypes in each of subsequent, first and single primary melanomas in our study. Scheibner et al¹⁶ found the commonest subtypes to be superficial spreading melanoma and nodular melanoma, and that each of subsequent and first melanomas were of the same histologic type in 74% of cases, which compares with 54% in the present study. The prominence of lentigo maligna melanoma in our study probably reflects refined diagnostic criteria and better clinical recognition of this subtype of melanoma, as well as the common occurrence in Australian populations (42% of melanomas in this analysis were from the Australian GEM center – data not shown) of high solar UV exposure, which is well known to be associated with lentigo maligna melanoma.⁴⁷

Ulceration has previously been shown to be less common^{2, 21} and regression to be more common¹⁶ in subsequent than in first melanomas. Consistently, ulceration was less common in subsequent melanomas in our study, which is in keeping with the lesser thickness of these tumors and a known correlation of ulceration with tumor thickness,⁴⁸ and regression was more common, but both could have been chance differences. The difference between our findings for regression and those of Scheibner et al¹⁶ may be due to the fact that there is considerable interobserver variation in, and poor reproducibility of, the histologic assessment of regression.^{49, 50}

Similarly to our study, dysplastic nevi have been found to occur more frequently in patients with multiple (38–63%) than single primary melanomas^{17, 22} and in the general population.⁵¹ These observations suggest that dysplastic nevi are markers of risk for

additional melanomas. They are in keeping with results of previous studies, which have shown that the presence of clinically and histologically diagnosed dysplastic nevi,^{5, 33} a family history of dysplastic nevi,⁴ and classical atypical mole syndrome²⁰ are associated with increased risk of multiple primary melanomas. Dysplastic nevi are also risk markers for the development of melanoma in melanoma-prone families.^{52–54} However, it is generally easier to detect a nevus remnant in thin melanomas (e.g. subsequent melanomas in patients with multiple melanomas) than in thick melanomas (e.g. first melanomas), as in the latter any residual nevus may have been overgrown by the invasive melanoma.

Differences in pathology between the paired melanomas in patients with multiple tumors, notably the reduced thickness and vertical growth in subsequent melanomas, are likely to reflect closer clinical surveillance and earlier diagnosis. Other differences, such as the more common occurrence of dysplastic nevi in association with subsequent melanomas, and the stronger association of subsequent melanomas with the most exposed body site (head and neck) are consistent with dysplastic nevi and sun exposure being risk factors or risk markers for and, in the case of dysplastic nevi, possible precursors to additional melanomas in patients with a cutaneous melanoma.

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Age, sex and anatomic site of tumors in patients with single primary melanomas and multiple primary melanomas included in the analysis

Table 1

Age	Single primary melanomas N=1989		First melanoma* N=395		Subsequent melanoma* N=395	
	Males	Females	Males	Females	Males	Females
<50 years	265 (25.5%)	458 (47.8%)	33 (12.0%)	39 (32.8%)	23 (8.3%)	23 (19.3%)
50–69 years	482 (46.8%)	314 (32.8%)	141 (51.1%)	55 (46.2%)	113 (40.9%)	54 (45.4%)
≥70 years	286 (27.7%)	186 (19.4%)	102 (37.0%)	25 (21.0%)	140 (50.7%)	42 (35.3%)
<i>Anatomic site</i>						
Head & neck	194 (18.9%)	109 (11.5%)	49 (17.8%)	15 (12.7%)	72 (26.2%)	20 (16.9%)
Trunk	596 (58.0%)	290 (30.5%)	160 (58.2%)	27 (22.9%)	138 (50.2%)	29 (24.6%)
Arms	147 (14.3%)	222 (23.3%)	37 (13.5%)	32 (27.1%)	32 (11.6%)	26 (22.0%)
Legs	90 (8.8%)	330 (34.7%)	29 (10.5%)	44 (37.3%)	33 (12.0%)	43 (36.4%)

* First and subsequent melanomas are primary invasive cutaneous melanomas in patients with multiple primary melanomas.

Table 2
Comparison of clinicopathologic features of subsequent and first melanomas in patients with multiple primary melanomas and single primary melanomas[†]

Pathology variable	Subsequent melanomas* N=395		First melanomas* N=395		Single primary melanomas N=1,989	
	No.	(%)	No.	(%)	No.	(%)
Site of melanoma						
Head & neck	92	(23)	64	(16)	303	(15)
Trunk	167	(42)	188	(48)	886	(45)
Arms	58	(15)	69	(17)	369	(19)
Legs	77	(19)	73	(18)	420	(21)
Missing	1		1		11	
				p=0.02 [§]		p<0.001 [‡]
Site of melanoma and exposure						
Usually exposed (head and neck)	108	(29)	95	(26)	442	(23)
Not usually exposed (trunk, limbs)	264	(71)	277	(74)	1,494	(77)
Missing	23		23		53	
				p=0.22		p=0.009
Histologic subtype						
Superficial spreading melanoma	246	(63)	266	(68)	1,411	(72)
Nodular melanoma	31	(8)	36	(9)	185	(9)
Lentigo maligna melanoma	91	(23)	55	(14)	196	(10)
Melanoma, not otherwise specified	19	(5)	24	(6)	135	(7)
Other	4	(1)	10	(3)	46	(2)
missing	4		4		16	
				p=0.002		p<0.001
Association with in situ melanoma						
In-situ & invasive	362	(92)	351	(89)	1,754	(88)
Invasive only	33	(8)	44	(11)	235	(12)
				p=0.21		p=0.05
Breslow thickness (mm)						

Pathology variable	Subsequent melanomas* N=395		First melanomas* N=395		Single primary melanomas N=1,989	
	No.	(%)	No.	(%)	No.	(%)
0.01–1.00	301	(77)	265	(68)	1,339	(68)
1.01–2.00	52	(13)	70	(18)	377	(19)
2.01–4.00	28	(7)	45	(11)	182	(9)
>4.00	11	(3)	12	(3)	79	(4)
<i>missing</i>	3		3		12	
				p=0.03		p=0.007
Clark level						
II	236	(60)	188	(48)	814	(41)
III	79	(20)	103	(26)	548	(28)
IV & V	78	(20)	102	(26)	608	(31)
<i>Missing</i>	2		2		19	
				p=0.003		p<0.001
Vertical growth						
Absent	223	(57)	180	(46)	706	(36)
Present	171	(43)	214	(54)	1,264	(64)
<i>Missing</i>	1		1		19	
				p=0.002		p<0.001
Ulceration						
Absent	358	(93)	344	(89)	1,782	(91)
Present	28	(7)	42	(11)	179	(9)
<i>Missing</i>	9		9		28	
				p=0.09		p=0.24
Presence of mitoses						
Absent	260	(66)	245	(63)	1,109	(56)
Present	132	(34)	147	(38)	877	(44)
<i>Missing</i>	3		3		3	
				p=0.25		p<0.001
Mitotic rate (per mm ²)						
Absent	255	(70)	233	(64)	1,109	(59)

Pathology variable	Subsequent melanomas* N=395		First melanomas* N=395		Single primary melanomas N=1,989		p-value
	No.	(%)	No.	(%)	No.	(%)	
1	27	(7)	35	(10)	296	(16)	
2	18	(5)	21	(6)	136	(7)	
3-4	18	(5)	24	(7)	125	(7)	
5-9	25	(7)	28	(8)	105	(6)	
10+	20	(6)	22	(6)	111	(6)	
Missing	32		32		107		p<0.001
Tumor-infiltrating lymphocytes							
Non-brisk	236	(60)	245	(63)	1,254	(64)	
Brisk	74	(19)	71	(18)	273	(14)	
Absent	82	(21)	76	(19)	430	(22)	
Missing	3		3		32		p=0.04
Presence of regression							
Absent	263	(67)	278	(71)	1,322	(67)	
Present	129	(33)	114	(29)	639	(33)	
Missing	3		3		28		p=0.18
Co-existing nevus							
Not identified	257	(65)	271	(69)	1,430	(73)	
Dysplastic	96	(24)	60	(15)	226	(12)	
Common acquired	34	(9)	59	(15)	267	(14)	
Congenital & other	7	(2)	4	(1)	38	(2)	
Missing	1		1		28		p<0.001
Satellites							
Absent	314	(81)	326	(84)	1,400	(71)	
Present	6	(2)	4	(1)	13	(1)	
N/A (biopsy only)	70	(18)	60	(15)	546	(28)	

Pathology variable	Subsequent melanomas* N=395		First melanomas* N=395		Single primary melanomas N=1,989		p-value
	No.	(%)	No.	(%)	No.	(%)	
<i>Missing</i>	5		5		30		p<0.001
Pigmentation							
Absent	24	(6)	29	(7)	179	(9)	
Present	366	(94)	361	(93)	1,796	(91)	
<i>Missing</i>	5		5		14		
Solar elastosis							p=0.45
Absent	69	(19)	78	(21)	663	(35)	
Mild/moderate	208	(57)	213	(58)	938	(49)	
Marked	88	(24)	74	(20)	310	(16)	
<i>Missing</i>	30		30		78		
Total	395	(100)	395	(100)	1,989	(100)	p<0.001

[†] Missing and N/A categories were excluded from the percentages.

* Subsequent and first melanomas are primary invasive cutaneous melanomas in patients with multiple primary melanomas.

[‡] P-values for comparison of single primary melanomas and subsequent melanomas.

[§] P-values for comparison of first and subsequent melanomas in patients with multiple melanomas.

Table 3

Results of a stepwise multiple regression analysis to identify variables that independently distinguished subsequent from first melanomas in patients with multiple primary melanomas

Pathology variables	Tumor		Prevalence ratios [†] (95% CI) for MPM2	p-value
	First melanoma*	Subsequent melanoma*		
Co-existing nevus				<0.001
Not identified	226	227	1.0	
Dysplastic	54	83	1.29 (1.10–1.53)	
Common acquired	54	22	0.61 (0.43–0.87)	
Congenital & other	2	4	1.78 (0.97–3.29)	
Vertical growth				0.01
Absent	157	191	1.0	
Present	179	145	0.82 (0.70–0.95)	
Site of melanoma				0.04
Head & neck	56	78	1.0	
Trunk	156	143	0.81 (0.68–0.96)	
Arms	61	48	0.78 (0.60–1.02)	
Legs	63	67	0.98 (0.81–1.19)	

* First and subsequent melanomas are primary invasive cutaneous melanomas in patients with multiple primary melanomas.

[†] Prevalence ratio for subsequent melanoma compared with first melanoma in patients with multiple primary melanomas, clustered on patient and adjusted for age at diagnosis of lesion, sex, GEM center and all other variables in table. Highest p-value for entry 0.2; 336 cases with complete data for all variables included in the analysis.