

1 **MULTIPLE MYELOMA: FAMILY HISTORY AND MORTALITY IN SECOND**
2 **PRIMARY CANCERS**

3
4 Subhayan Chattopadhyay, Hongyao Yu, Amit Sud, Jan Sundquist, Asta Försti, Akseli Hemminki
5 and Kari Hemminki
6
7
8
9
10
11
12
13
14
15
16
17
18
19
20
21
22
23
24
25
26
27
28
29

30 Key words: myeloma, familial risk, lymphoma, leukemia, second cancer.
31

32 Running title: Familial second primary.
33
34

35 **CONFLICTS OF INTEREST**

36 A.H. is shareholder in Targovax ASA. A.H. is employee and shareholder in TILT
37 Biotherapeutics Ltd. Other authors declared no conflict of interest.
38
39

40 Since cancer survival rates in general are increasing, second primary cancers (SPCs) account for
41 an increasing proportion of the overall cancer burden. In some cancer registries they account for
42 more than 20% of new diagnoses (1). Contributing factors for SPCs may be multiple, including
43 iatrogenic adverse effects of chemotherapy or radiation, increased surveillance and the same
44 causes that influenced patients' first cancers, including family history and environmental causes
45 (2-4). Chemotherapy and radiation induce DNA damage which increases the risk of SPCs, and
46 therapy associated immunosuppression could also play a role. Treatment for MM involves
47 intense chemotherapy and concerns about SPCs have been raised, particularly relating to the
48 possible effects of lenalidomide and melphalan (5). The impact of family history was recently
49 shown in survivors of Hodgkin lymphoma with an excess of lung, colorectal and breast cancers
50 in survivors with a family history of these cancers (6). The potential importance of family history
51 is emphasized by the fact that about 50% of patients with first primary cancer have a first-degree
52 relative diagnosed with some cancer (7). This proportion was also high among patients
53 diagnosed with multiple myeloma (MM), 61% (7). The other cancers in family members were
54 diverse; including chronic lymphocytic leukemia and colorectal and prostate cancers (8, 9).

55
56 In the present study we use the Swedish Family-Cancer Database, with two goals, first to assess
57 the influence of family history on the risk of SPC, and second to estimate the influence of SPC
58 on mortality in MM in family members (7). A family history implies that the type of SPC (e.g.,
59 lung cancer) was the same cancer that was diagnosed in a parent or sibling (e.g. lung cancer).

61 **Methods**

62 In the Swedish Family-Cancer Database the second generation 'offspring' was defined as
63 individuals born after 1931 and their parents were defined as the parental generation. Another
64 truncation of data was caused by the start of cancer registration in Sweden in 1958. The study
65 included 25,787 MM diagnosed from 1958 to 2015; of these 5205 were diagnosed in the
66 offspring generation with a median age at diagnosis of 62 years. Among MM patients 360
67 (6.9%) were diagnosed with SPC after a median follow-up time of 4 years. Among these 360,
68 246 (68.3%) had a first-degree family history of any cancer.

69
70 Relative risks (RRs) were assessed with incidence rate ratios, estimated with RRs regressed over
71 a fixed effects generalized Poisson model. RRs for SPC were obtained by comparing incidence
72 rates for SPC X in MM patients with rates for first cancer X in the background population of the
73 database. Family history was defined among parents and siblings. Familial RRs were estimated
74 by comparing incidence rates between MM patients diagnosed with cancer X as SPC and having
75 a family history of cancer X against those diagnosed with first cancer X in the population; the
76 reference rate was the same as above. Sex, age group, calendar-period, socio-economic status
77 and residential areas were treated as potential confounders and were adjusted for in the
78 regression model. Follow-up commenced from diagnosis of MM and was terminated on SPC
79 diagnosis, emigration, death or end of follow-up period, i.e. 2015, whichever occurred first.

80 Confidence intervals were calculated for 5%, 1% and 0.1% level of significance (10). All
81 cancer-related deaths were stratified into MM, SPC and other causes, including cancers defined
82 in death certificates and non-neoplastic causes of death. Additive and multiplicative interactions
83 of family history and risk of SPC were tested as described (11).

84

85 The study was approved by the Ethical Committee of Lund University. Analyses are performed
86 in SAS v9.4; please contact the authors for codes.

87

88 **Results**

89

90 Among 5,205 MM patients, 360 (6.9%) were diagnosed with a SPC. Familial SPCs were
91 compared to non-familial SPCs in Table 1, which lists all SPCs with at least two cases having the
92 same (concordant) tumor in a parent or sibling. Ignoring the overlapping impact of more than
93 one cancer in family, prostate cancer was the major contributor to the family history (20%)
94 followed by colorectal (14%), breast (10%), bladder (5%), lung cancer and skin SCC (4% both).
95 In patients without a family history of cancer, the risk of SPC was increased for skin cancer
96 (squamous cell carcinoma, SCC, RR = 2.58) and leukemia (RR = 4.55). For patients with a
97 family history of cancer, even though case numbers were low, risks were significantly elevated
98 in a trend test for colorectal (RR/familial = 2.10 vs. RR/non-familial = 1.01), prostate
99 (RR/familial = 1.60 vs. RR/non-familial = 0.56) and skin SCC (RR/familial = SCC, 8.82 vs.
100 RR/non-familial = 2.58). The trend test was of borderline significance ($P = 0.061$) for lung
101 cancer (RR/familial = 5.40 vs. RR/non-familial = 1.13). The highest SPC risk was observed for
102 MM patients with a family history of leukemia (RR = 9.14, only 2 cases). Patients with SPC with
103 any familial cancer (N = 246) were 68.3% of all SPCs and the RR was 1.38 vs. 1.13 respectively
104 (trend test $P < 0.001$). We tested interactions of significant family risks and risk of SPC and
105 found a stronger than additive interaction for skin cancer ($P = 0.04$).

106

107 In order to check for possible skewed patient recruitment based on the multiple applied
108 conditions were plotted the patient accrual over the study period (Supplementary Figure 1). The
109 diagram shows MM patients with SPC and with or without family history (246 and 114 patients)
110 plotted by 5-year intervals of MM diagnosis. No skewing of case accrual was observed.

111

112 The total number of deaths by the end of 2015 was 2872 (55.2%) among 5205 patients; and the
113 total number of deaths among 360 patients with SPC was 228 (60.6%). The proportion was
114 equally high among 246 patients with familial SPC, of whom 146 (59.3%) had died.

115 Kolmogorov-Smirnov test on proportion difference found no evidence of statistical difference (P
116 > 0.05)

117

118 MM was the most common cause of death in patients without a SPC (83.4%, 2194/2629), with
119 16.6% of deaths due to other causes (data not shown). For MM patients with a SPC, the

120 distribution of causes of death is shown in Table 2. MM was the leading cause with 38.7% of
121 deaths, followed by SPC 35.8% and other causes (25.5%); among other causes the majority of
122 deaths (62.9%) were due to non-neoplastic causes. The mortality of SPC varied between second
123 cancer types. For second pancreatic cancer, all 7 patients died of this cancer; more than half of
124 MM patients died of SPC when it was lung or nervous system cancer or leukemia. Other causes
125 were important for CUP as SPC which is due to the practice of rarely describing CUP as a cause
126 of death (12). Among 82 deaths in patients with SPC without a cancer family history, majority
127 was due to MM (36.6%), closely followed by SPCs (34.2%). Kolmogorov-Smirnov test found no
128 significant difference in proportion contribution by the different causes of death in patients with
129 or without family history ($P > 0.05$).

130

131

132 **Discussion**

133

134 The novel aspect of this study was the demonstration of the impact of familial risk on SPCs in
135 MM patients. Accordingly, as many as 68.3% of SPCs were familial, i.e., a parent or sibling of
136 MM patients were diagnosed with any cancer, moderately higher compared to that of 59.9%
137 patients without an SPC. For three SPCs with significant risks, including colorectal, prostate and
138 skin cancers, the family members had exactly the same cancer as was the SPC. It is interesting
139 that in a recent study from this database the most consistent familial association between MM
140 and first primary cancers included colorectal and prostate cancer and leukemia (9). This may not
141 be coincidental and shared susceptibility may contribute to these findings. We showed also that
142 MM patients with SPC appeared to have moderately worse prognosis (60.6% dead) compared to
143 all MM patients (55.2% dead), while family history of SPCs did not increase mortality (59.3%
144 dead). The limitation of the study was a relatively small sample size in spite of nation-wide
145 coverage. The reason is that survival in MM, although improving, is still relatively poor whereby
146 the time-window for SPCs is narrow (13). Due to the small numbers we did not undertake formal
147 hazard ratio analysis for survival.

148

149 Therapy-related SPCs in MM have mainly been associated with acute myeloid leukemia which
150 has been increased also in a recent study on German and Swedish MM patients (5, 14). The
151 Swedish population of that study partially overlaps with the present one were a risk (RR 4.41) of
152 second leukemia was observed in patients lacking family history. Therapy-related side effects are
153 still considered relatively weak in MM but the situation may change when larger patient groups
154 achieve long survival times (5). Family history needs to be considered a possible confounder in
155 therapy related studies on SPCs.

156

157 In conclusion, 68.3% of MM patients with SPC in had a family history of any cancer.
158 Significantly increased associations were found for second colorectal, prostate and skin cancers
159 and family members diagnosed with these cancers. With continued therapeutic successes in MM

160 treatment SPCs will be receiving increasing attention whereby the contributing role of family
161 history deserves inquiry into its mechanistic underpinnings.

162

163 ACKNOWLEDGEMENTS

164 A.S. is the recipient of Guest scientist Fellowship of DKFZ. Supported by the Harald Huppert
165 Foundation, The German Federal Ministry of Education and Research (eMed, Cliommics
166 01ZX1309B), Deutsche Krebshilfe, Jane and Aatos Erkkö Foundation, University of Helsinki
167 and Helsinki University Central Hospital

168 AFFILIATIONS

169 Division of Molecular Genetic Epidemiology, German Cancer Research Center (DKFZ), Im
170 Neuenheimer Feld 580, D-69120, Heidelberg, Germany

171 Subhayan Chattopadhyay, Hongyao Yu, Asta Försti, Kari Hemminki

172

173 Faculty of Medicine, University of Heidelberg, Heidelberg, Germany

174 Subhayan Chattopadhyay, Hongyao Yu

175

176 Division of Genetics and Epidemiology, The Institute of Cancer Research, London, UK

177 Amit Sud

178

179 Center for Primary Health Care Research, Lund University, 20502 Malmö, Sweden

180 Jan Sundquist, Asta Försti, Kari Hemminki

181

182 Department of Family Medicine and Community Health, Department of Population Health

183 Science and Policy, Icahn School of Medicine at Mount Sinai, New York, USA

184 Jan Sundquist

185

186 Center for Community-based Healthcare Research and Education (CoHRE), Department of

187 Functional Pathology, School of Medicine, Shimane University, Japan

188 Jan Sundquist

189

190 Cancer Gene Therapy Group, Faculty of Medicine, University of Helsinki, Finland

191 Akseli Hemminki

192

193 Comprehensive Cancer Center, Helsinki University Hospital, Helsinki, Finland

194 Akseli Hemminki

195

196 CORRESPONDENCE

197 Subhayan Chattopadhyay

198 Division of Molecular Genetic Epidemiology,
199 German Cancer Research Center (DKFZ),
200 Im Neuenheimer Feld 580, Heidelberg-69120, Germany.
201 Telephone: +496221421800
202 Fax: +496221421810
203 Email: S.Chattopadhyay@dkfz.de

204 AUTHOR CONTRIBUTIONS

205 Design: KH
206 Acquisition of data: JS
207 Statistical analysis and interpretation: SC, HY, KH, AF, AS.
208 Manuscript writing: KH, SC, AH, AF.
209 Approval of the final text: All authors

210 REFERENCES

- 211
- 212 1. CentreforEpidemiology. Cancer incidence in Sweden 2012. Stockholm: The National
213 Board of Health and Welfare; 2013.
 - 214 2. Travis LB, Demark Wahnefried W, Allan JM, Wood ME, Ng AK. Aetiology, genetics
215 and prevention of secondary neoplasms in adult cancer survivors. *Nature reviews Clinical*
216 *oncology*. 2013;10(5):289-301.
 - 217 3. Schaapveld M, Aleman BM, van Eggermond AM, Janus CP, Krol AD, van der Maazen
218 RW, et al. Second Cancer Risk Up to 40 Years after Treatment for Hodgkin's Lymphoma. *N*
219 *Engl J Med*. 2015;373(26):2499-511.
 - 220 4. Wood ME, Vogel V, Ng A, Foxhall L, Goodwin P, Travis LB. Second malignant
221 neoplasms: assessment and strategies for risk reduction. *J Clin Oncol*. 2012;30(30):3734-45.
 - 222 5. Musto P, Anderson KC, Attal M, Richardson PG, Badros A, Hou J, et al. Second primary
223 malignancies in multiple myeloma: an overview and IMWG consensus. *Ann Oncol*.
224 2017;28(2):228-45.
 - 225 6. Sud A, Thomsen H, Sundquist K, Houlston RS, Hemminki K. Risk of second cancer in
226 Hodgkin lymphoma survivors and the influence of family history. *J Clin Oncol*. 2017;35:1584-
227 90.
 - 228 7. Frank C, Sundquist J, Yu H, Hemminki A, Hemminki K. Concordant and discordant
229 familial cancer: Familial risks, proportions and population impact *Int J Cancer*. 2017;140:1510-6.
 - 230 8. Kristinsson SY, Bjorkholm M, Goldin LR, Blimark C, Mellqvist UH, Wahlin A, et al.
231 Patterns of hematologic malignancies and solid tumors among 37,838 first-degree relatives of
232 13,896 patients with multiple myeloma in Sweden. *Int J Cancer*. 2009;125(9):2147-50.
 - 233 9. Frank C, Fallah M, T. C, Mai EK, Sundquist J, Forsti A, et al. Search for familial
234 clustering of multiple myeloma with any cancer Leukemia. 2016;30:627-32.
 - 235 10. Tibshirani R. Estimating Transformations for Regression via Additivity and Variance
236 Stabilization. *J Am Stat Associat*. 1988;83:394-405.
 - 237 11. Zhang H, Bermejo JL, Sundquist J, Hemminki K. Modification of second cancer risk
238 after malignant melanoma by parental history of cancer. *Br J Cancer*. 2008;99:536-8.

239 12. Hemminki K, Bevier M, Sundquist J, Hemminki A. Site-specific cancer deaths in cancer
240 of unknown primary diagnosed with lymph node metastasis may reveal hidden primaries. *Int J*
241 *Cancer*. 2013;132:944-50.

242 13. Kristinsson SY, Landgren O, Dickman PW, Derolf AR, Bjorkholm M. Patterns of
243 survival in multiple myeloma: a population-based study of patients diagnosed in Sweden from
244 1973 to 2003. *J Clin Oncol*. 2007;25:1993-9.

245 14. Chen T, Fallah M, Brenner H, Jansen L, Mai EK, Castro FA, et al. Risk of Second
246 Primary Cancers in Multiple Myeloma Survivors in German and Swedish Cancer Registries.
247 *Scientific reports*. 2016;6:22084.
248

Table 1. Relative risks of SPCs among all multiple myeloma patients stratified over family

Cancer	At least 1 FDR with cancer			No FDR with cancer			Total			Trend test P value
	N	RR	95% CI	N	RR	95% CI	N	RR	95% CI	
Colorectum	7	2.10	1.00 - 4.41	27	1.01	0.69 - 1.47	34	1.13	0.81 - 1.58	0.033
Lung	3	5.40	1.74 - 16.75	10	1.13	0.61 - 2.10	13	1.38	0.80 - 2.38	0.061
Breast	4	1.13	0.42 - 3.01	24	0.93	0.62 - 1.39	28	0.95	0.66 - 1.38	0.176
Prostate	20	1.60	1.03 - 2.48	38	0.56	0.41 - 0.77	58	0.72	0.56 - 0.93	0.006
Melanoma	2	5.04	1.26 - 20.14	18	1.46	0.92 - 2.32	20	1.57	1.01 - 2.44	0.087
Skin (SCC)	4	8.82	3.31 - 23.52	31	2.58	1.81 - 3.67	35	2.81	2.01 - 3.91	0.029
Leukemia	2	9.14	2.29 - 36.55	32	4.41	3.11 - 6.24	34	4.55	3.25 - 6.37	0.093
All	246	1.38	1.22 - 1.57	114	1.13	0.94 - 1.36	360	1.29	1.17 - 1.43	<0.001

Abbreviation:

FDR, first degree relative; N, frequency; RR, relative risk; CI, confidence interval;

SCC, squamous cell carcinoma;

Bold, italics and underline indicate 5%, 1% and 0.1% level of significance;

Table 2. Causes of death distribution of multiple myeloma patients diagnosed with SPC

	MM		^a SPC		Other causes	
	N	%	N	%	N	%
Cancer						
UAT	2	50.0	2	50.0	-	-
Stomach	-	-	4	100.0	-	-
Colorectum	8	33.3	11	45.8	5	20.9
Anus	-	-	1	100.0	-	-
Liver	2	33.3	3	50.0	1	16.7
Pancreas	-	-	7	100.0	-	-
Lung	3	13.6	15	68.2	4	18.2
Breast	6	42.9	1	7.1	7	50
Cervix	-	-	1	100.0	-	-
Ovary	1	50.0	1	50.0	-	-
Prostate	11	42.3	5	19.2	10	38.4
Kidney	3	37.5	3	37.5	2	25
Urinary bladder	5	41.7	3	25.0	4	33.3
Melanoma	7	58.3	3	25.0	2	16.7
Skin (SCC)	16	72.7	1	4.5	5	22.7
Nervous system	3	42.9	4	57.1	-	-
NHL	5	45.5	4	36.4	2	18.2
Hodgkin lymphoma	-	-	1	50.0	1	50
Leukemia	7	24.1	16	55.2	6	20.6
CUP	3	21.4	1	7.1	10	71.4
^b Total	94	38.7	87	35.8	62	25.5

Abbreviations:

MM multiple myeloma; SPC, second primary cancer; UAT, upper aerodigestive tract; SCC, squamous cell carcinoma; NHL, non-Hodgkin lymphoma; CUP, cancer of unknown primary;

^a Cases noted only when at least one death is observed due to second cancer.

^b Total includes all cancers without constraints.