



OPEN ACCESS

Original research

# Cancer risk and tumour spectrum in 172 patients with a germline *SUFU* pathogenic variation: a collaborative study of the SIOPE Host Genome Working Group

Léa Guerrini-Rousseau <sup>1,2</sup>, Julien Masliah-Planchon,<sup>3</sup> Sebastian M Waszak,<sup>4,5</sup> Pia Alhopuro,<sup>6</sup> Patrick R Benusiglio,<sup>7</sup> Franck Bourdeaut,<sup>3</sup> Ines B Brecht,<sup>8</sup> Giada Del Baldo,<sup>9</sup> Sandeep Kumar Dhanda <sup>10</sup>, Maria Luisa Garré,<sup>11</sup> Corrie E M Gidding,<sup>12</sup> Steffen Hirsch,<sup>13,14</sup> Pauline Hoarau,<sup>1</sup> Mette Jorgensen,<sup>15</sup> Christian Kratz,<sup>16</sup> Lucie Lafay-Cousin,<sup>17</sup> Angela Mastronuzzi,<sup>18</sup> Lorenza Pastorino,<sup>19,20</sup> Stefan M Pfister,<sup>14,21,22</sup> Christopher Schroeder,<sup>23</sup> Miriam Jane Smith <sup>24</sup>, Pia Vahteristo,<sup>6,25</sup> Roseline Vibert <sup>26</sup>, Catheline Vilain,<sup>27,28</sup> Nicolas Waespe <sup>29,30</sup>, Ingrid M Winship,<sup>31</sup> D Gareth Evans,<sup>32</sup> Laurence Brugieres<sup>2,33</sup>

► Additional supplemental material is published online only. To view, please visit the journal online (<http://dx.doi.org/10.1136/jmedgenet-2021-108385>).

For numbered affiliations see end of article.

## Correspondence to

Dr Léa Guerrini-Rousseau, Department of Children and Adolescents Oncology, Gustave Roussy, Villejuif, Île-de-France, France; [lea.guerrini-rousseau@gustaveroussy.fr](mailto:lea.guerrini-rousseau@gustaveroussy.fr)

Received 10 January 2022

Accepted 23 April 2022

Published Online First 29 June 2022



© Author(s) (or their employer(s)) 2022. Re-use permitted under CC BY. Published by BMJ.

**To cite:** Guerrini-Rousseau L, Masliah-Planchon J, Waszak SM, *et al*. *J Med Genet* 2022;**59**:1123–1132.

## ABSTRACT

**Background** Little is known about risks associated with germline *SUFU* pathogenic variants (PVs) known as a cancer predisposition syndrome.

**Methods** To study tumour risks, we have analysed data of a large cohort of 45 unpublished patients with a germline *SUFU* PV completed with 127 previously published patients. To reduce the ascertainment bias due to index patient selection, the risk of tumours was evaluated in relatives with *SUFU* PV (89 patients) using the Nelson-Aalen estimator.

**Results** Overall, 117/172 (68%) *SUFU* PV carriers developed at least one tumour: medulloblastoma (MB) (86 patients), basal cell carcinoma (BCC) (25 patients), meningioma (20 patients) and gonadal tumours (11 patients). Thirty-three of them (28%) had multiple tumours. Median age at diagnosis of MB, gonadal tumour, first BCC and first meningioma were 1.5, 14, 40 and 44 years, respectively. Follow-up data were available for 160 patients (137 remained alive and 23 died). The cumulative incidence of tumours in relatives was 14.4% (95% CI 6.8 to 21.4), 18.2% (95% CI 9.7 to 25.9) and 44.1% (95% CI 29.7 to 55.5) at the age of 5, 20 and 50 years, respectively. The cumulative risk of an MB, gonadal tumour, BCC and meningioma at age 50 years was: 13.3% (95% CI 6 to 20.1), 4.6% (95% CI 0 to 9.7), 28.5% (95% CI 13.4 to 40.9) and 5.2% (95% CI 0 to 12), respectively. Sixty-four different PVs were reported across the entire *SUFU* gene and inherited in 73% of cases in which inheritance could be evaluated.

**Conclusion** Germline *SUFU* PV carriers have a life-long increased risk of tumours with a spectrum dominated by MB before the age of 5, gonadal tumours during adolescence and BCC and meningioma in adulthood, justifying fine-tuned surveillance programmes.

## INTRODUCTION

Gorlin syndrome (GS) (MIM 109400), or nevoid basal cell carcinoma syndrome (NBCCS), is an autosomal dominantly inherited syndrome characterised by developmental anomalies including

## WHAT IS ALREADY KNOWN ON THIS TOPIC

- ⇒ Germline *SUFU* pathogenic variant (PV) was described for the first time associated with the occurrence of medulloblastoma by Michael Taylor *et al* in 2002.
- ⇒ Before our study, germline *SUFU* PVs were known to be associated with a cancer predisposition syndrome predisposing to SHH-medulloblastoma during the first 3 years of life as well as cancers associated with Gorlin syndrome.
- ⇒ During the last years, >100 patients with a germline *SUFU* PV have been reported, but most often, these publications are case reports in which the *SUFU* PV was identified after the occurrence of cancer.
- ⇒ Due to the rarity of this clinical situation, little was known about tumour risks and outcome of patients in this condition.

## WHAT THIS STUDY ADDS

- ⇒ Germline *SUFU* PV carriers have a life-long increased risk of tumours.
- ⇒ Data from this large series allow describing the oncological spectrum of *SUFU* PVs, dominated by medulloblastoma before the age of 5 years, gonadal tumors during adolescence and basal cell carcinoma and meningioma in adulthood.
- ⇒ We also aimed to evaluate cancer risk, but as most index patients have been identified after the occurrence of a malignancy, we analysed the cumulative risk of tumour in relatives only, after exclusion of the index cases in order to reduce bias.
- ⇒ We were able to confirm that the tumour penetrance (any type of tumour) is high, although incomplete reaching 44% at 50 years.

**HOW THIS STUDY MIGHT AFFECT RESEARCH, PRACTICE AND/OR POLICY**

- ⇒ Thanks to this large international cooperation, we could describe the spectrum of tumours, the cumulative risk of cancer as well as the period of onset of each tumour type during life associated with a germline *SUFU* PV.
- ⇒ These information allow designing guidelines for PV carriers follow-up based on comprehensive data.

macrocephaly, frontal bossing, hypertelorism and has been described as a cancer predisposition syndrome.<sup>1,2</sup> The tumour spectrum includes malignant tumours, mostly basal cell carcinomas (BCC) and medulloblastomas (MB), and benign tumours such as keratocystic odontogenic tumours, meningiomas, ovarian or cardiac fibromas.<sup>13–5</sup> Most individuals affected by GS have a heterozygous germline pathogenic variant (PV) in Sonic Hedgehog pathway genes: Patched 1 (*PTCH1*)<sup>6,7</sup> or Suppressor of fused (*SUFU*).<sup>8,9</sup> A GS-like clinical presentation has been recently described in children with heterozygous germline *GPR161* variants.<sup>10</sup> The role of Patched 2 (*PTCH2*)<sup>11,12</sup> in the pathogenesis of GS has also been suggested and questioned.<sup>13</sup>

Defining the incidence and spectrum of tumours in *SUFU*-associated GS is complicated due to the paucity of information collected so far. The association of germline *SUFU* PVs and nodular desmoplastic MB was described for the first time in 2002 by Taylor *et al.*<sup>14</sup> Since then, most information we have on germline *SUFU* mutation carriers comes from patients identified after the occurrence of a tumour, mainly MB. A few *SUFU* mutation carriers have also been identified after the occurrence of a meningioma<sup>15,16</sup> and cutaneous cancers.<sup>17,18</sup> Additional information comes from cohorts of patients presenting the clinical characteristics of GS in whom 5% are identified with a *SUFU* PV.<sup>19</sup> In a large series of 1022 patients with MB analysed for germline variants,<sup>20</sup> 6% of the patients were found to carry a germline PV in a known cancer predisposition gene, including 11 patients (1.1%) with a germline *SUFU* PV and 9 patients (0.9%) with a germline *PTCH1* PV, all with SHH-activated MB (SHH-MB).<sup>21,22</sup> The prevalence of germline *SUFU* or *PTCH1* PVs in SHH-MB below the age of 3 years was 21%.<sup>20</sup>

In recent years, >100 patients with a germline *SUFU* PV have been reported,<sup>8,9,14,19,20,23–36</sup> mainly as case reports; but data quantifying tumour risk and outcome of these patients remain scarce. The recent creation of the Host Genome Working Group (HGWG) in the European branch of the International Society of Pediatric Oncology (SIOPE) aimed at improving care for patients with paediatric cancer predisposition syndromes. The SIOPE-HGWG allowed us to set up a large international collaboration to increase the knowledge on this predisposition syndrome. The objective of this study was to describe the tumour spectrum and cancer risks specifically associated with germline *SUFU* PVs in order to provide recommendations to affected patients and their family members based on a comprehensive cohort.

**PATIENTS AND METHODS****Inclusion criteria**

This study includes only individuals with a germline PV in the *SUFU* gene referred to as *SUFU* PV carriers. We analysed the literature to collect all patients with a *SUFU* PV in articles published before 1 January 2021. We contacted the authors of these publications to obtain follow-up information. In

addition, through the SIOPE-HGWG, we also collected data of unpublished patients from eight different countries.

**Data collection**

For all *SUFU* PV carriers, we collected data on tumours identified so far and vital status at the last follow-up. In each family, the first patient in whom the PV was identified was defined as the index patient, whether she/he had a tumour or not. All their family members, in whom the *SUFU* PV was identified after its identification in the index patient, were qualified as relatives. Data from French patients, currently collected in the French ‘Observatory of Genetic Cancer Predisposition Syndromes in Children and Adolescents’ (Observatoire des syndromes de prédisposition génétique au cancer des enfants et des adolescents, PREDCAP, IRB00003888) were merged with data obtained from each national group.

**Statistics**

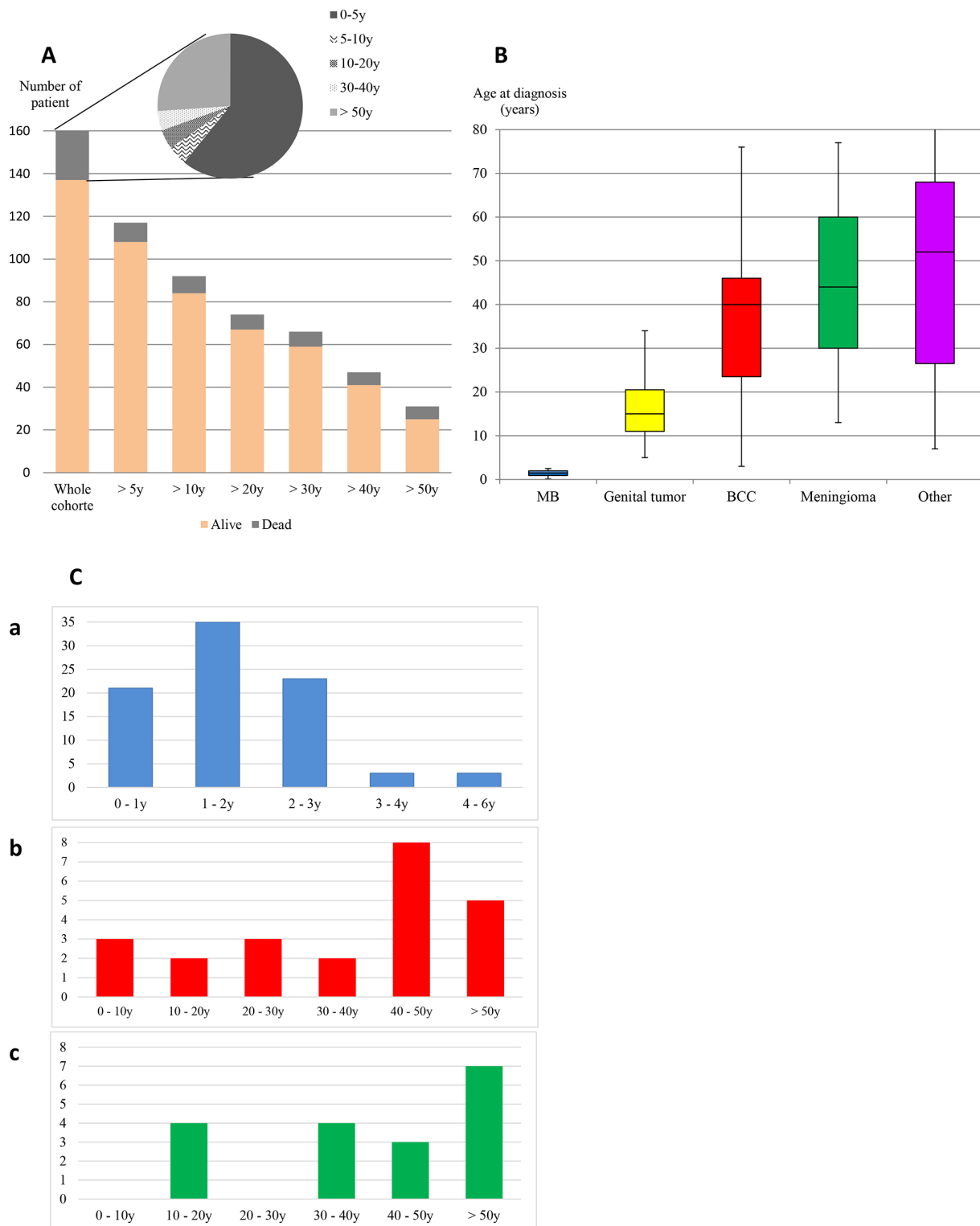
Baseline values (ie, at diagnosis) were expressed as medians and ranges for continuous variables, and as numbers and percentages for categorical variables, and compared using the  $\chi^2$  test. Overall survival (OS) rates were calculated using the Kaplan-Meier method. Overall survival time after MB was estimated from the date of diagnosis of the MB to death, whatever the cause, or the date of the last follow-up. The 95% CI values for OS rates were estimated with the Rothman method. The Nelson and Aalen estimator<sup>37</sup> was used to model the cumulative incidence curves in relatives carrying the *SUFU* PV. To study genotype-phenotype correlation, the impact of the type of the *SUFU* PV on the risk of MB was estimated using the  $\chi^2$  test, and p values <0.05 were considered statistically significant.

**RESULTS****General characteristics**

Overall, we identified 172 *SUFU* PV carriers (83 index patients and 89 relatives) from 83 families, including 127 individuals previously reported.<sup>8,9,14,19,20,23–36,38,39</sup> In most cases, the *SUFU* PV had been identified in patients with MB either through systematic screening for *SUFU* PV (in 74 index patients, 89%) or because of a familial history of MB (5 families). Another tumour type was the presenting feature in only seven patients: BCC (one patient),<sup>28</sup> BCC and meningioma (three patients),<sup>30,32,38</sup> multiple meningiomas (one patient), a bilateral ovarian stromal tumour (one patient) and a pancreatic carcinoma (one patient).<sup>33</sup> In two patients, the *SUFU* PV was identified in the exploration of a GS phenotype associated with developmental delay but without tumour.

Follow-up data were available for 160 patients. The median age at last follow-up for the whole cohort was 19.5 years (range 0.1–91). At least one tumour has been reported in 117 individuals (68%) while 55 (32%) individuals (including two index patients) were defined as healthy PV carriers, that is, without tumour until their last follow-up (median 38.5 years, range 2–91). Healthy carriers were significantly older at last follow-up than affected patients (median 10 years, range 0.1–85) ( $p=0.00015$ ).

The age distribution (patient exposed to the risk) and status at the last follow-up are illustrated in [figure 1A](#) for 160 patients with available data. At the last follow-up, 137 patients were alive with a median age of 25.5 years (range 0.1–91), and 23 patients died at a median age of 3.5 years (range 0.1–85). Overall, 1–5 malignant or benign



**Figure 1** Distribution of age of *SUFU* pathogenic variant carriers. (A) Number of patient exposed to the risk according to the age, (B) patient's age at diagnosis of tumour onset depending on the type of tumour, (C) patient's age at diagnosis of tumour onset, for medulloblastoma (MB) (C (a)), for the first basal cell carcinoma (BCC) (C (b)) and for the first meningioma (C (c)).

tumours were diagnosed in 117 patients, 81/83 (97.5%) index patients and 36/89 (44.5%) relatives. The distribution of tumour types, number of patients and age at diagnosis are described in [table 1](#), [figure 1B](#) and online supplemental figure 1. At least one second tumour was diagnosed in 33 patients (28%).

### Medulloblastoma

Overall, 86 patients (74 index patients and 12 relatives) were diagnosed with an MB in 76 distinct families. Multiple cases were diagnosed in six families (online supplemental table 1 and online supplemental figure 2). The MB was discovered during surveillance after presymptomatic screening for only one patient in this cohort.

**Table 1** Description of the distribution of the different tumour types in the cohort

Tumour type	Number of patients	Age at diagnosis (years) (median/range)
Medulloblastoma	86	1.5 (0.1–5)
BCC*	25	40 (3–76)
Meningioma*	20	44 (13–77)
Gonadal tumours	11	14 (5–34)
▶ Ovarian fibroma (n=4, including one patient with two successive asynchronous fibromas)		
▶ Bilateral ovarian stromal tumour (n=3, including one patient with successive asynchronous tumours)		
▶ Fibrosarcoma (n=1, ovarian; n=1, testicular)		
▶ Bilateral immature teratoma (n=1) <sup>27</sup>		
▶ Bilateral leiomyosarcoma (n=1) <sup>39</sup>		
<b>Other tumours</b>		
Sarcoma	2	46, 47, 47, 53
Carcinoma (n=10)		
▶ Breast carcinoma	3	37, 71, 77
▶ Thyroid carcinoma	2	7, 20
▶ Squamous cell carcinoma	2	>50
▶ Pancreatic carcinoma <sup>33</sup>	1	NA
▶ Bladder carcinoma	1	73
▶ Renal cell carcinoma	1	79
Skin hamartoma	3	2, 9, 55
Acute myeloid leukaemia	1	8
Pilocytic astrocytoma	1	20
Myeloma	1	79
Benign ileum myoma	1	42
Thoracic macrocystic lymphangioma	1	At birth

\*Age at diagnosis of first BCC/meningioma. BCC, basal cell carcinoma; NA, not available.

MB occurred as a first tumour in all these patients, with a median age at diagnosis of 1.5 years (range 0.1–5.8). MB was diagnosed before the age of 3 in all patients 81/86 (94.2%) but five (5.8%) aged respectively 3.3, 3.4, 3.8, 4 and 5.8 years (figure 1C (a)). Molecular subgrouping and/or histopathological subtype was available for 73/86 patients. All 35 patients with MB with molecular subgrouping belonged to the SHH subgroup. For 39 additional patients, histological subgrouping was concordant with SHH subgroup in 36/39 patients: nodular desmoplastic in 22 or extensive nodularity in 13. In three cases diagnosed before 2010 and with no histological review, the local pathologist reported a classic histology.

The median age at last follow-up was 6 years (range 0.1–37) in 74 patients with MB with follow-up data. Overall, 58/74 (78%) children were alive (median follow-up since diagnosis: 6.0 years, range 0.2–36) with 23 patients older than 10 years at the last follow-up. Sixteen patients died due to progression of the MB in 15 (median time since diagnosis: 1.2 years, range 0–4.3) and because of acute myeloid leukaemia as a second malignancy at age 7.8 years in one. The 5-year OS was 76% (95%CI 64% to 85%) (online supplemental figure 3).

A second tumour has been reported in 17 (out 71) patients after the occurrence of the MB, including 13/17 (76%) patients aged 10 years or more at the last follow-up. Among them, nine have received radiotherapy as part of the MB treatment (table 2).

### Basal cell carcinomas

BCC was reported in 25 patients (11 index patients and 14 relatives). The median age at diagnosis of the first BCC was 40 years (range 3–76) (figure 1C (b)). Only five patients were diagnosed with a BCC before the age of 20 years, all of them as a second malignancy. The number of BCCs is available for 18 patients (median age at last follow-up was 40.5 years). Half of them (aged 9–52 years at diagnosis of first BCC and 31–79 years at last follow-up) developed >20 BCC.

BCCs occurred as the first tumour in 16 patients, all occurring after the age of 20 years (median age of 43 years, range 22–76). Most patients (11/16) were identified through a systematic screening for *SUFU* PV in GS cohorts and have already been reported.<sup>8 19 28 30 32</sup> All patients but eight developed at least one other tumour before (11 tumours in 9 patients, including 7 MBs, 2 sarcomas, 1 ovarian tumour and 1 meningioma) or after BCC diagnosis (18 tumours in 15 patients). Among the 23 MB survivors aged 10 years or more at the last follow-up, 6 patients (26%) developed a BCC with a median time between the diagnosis of MB and BCC of 16.8 years (range 7–26). Of those with available data, 5/5 patients were treated with radiotherapy (table 2).

### Meningiomas

Twenty patients from 13 different families have been reported with a meningioma (13 index patients and 7 relatives); all were intracranial. They were described as a first, second, third and fourth tumour for four, nine, six and one patient(s). Meningiomas have been reported in members of a large family with several cases,<sup>27</sup> in patients treated for MB or in their relatives, or in patients with clinical features suggestive of GS. Seven patients developed multiple meningiomas, including the five familial cases reported in 2012.<sup>27</sup> The median age at diagnosis of the first meningioma was 44 years (range 13–77) (figure 1C (c)). Eight patients had a meningioma before the age of 35 years (median age of 24.5 years, range 13–35), occurring after radiotherapy for an MB in the 6/7 patients with data on MB treatment (table 2). Only one patient developed a meningioma 12 years after the occurrence of an MB treated without radiotherapy.

### Gonadal and other tumours

Eleven patients (seven index patients and four relatives) were diagnosed with a gonadal tumour, which occurred as a first tumour in four cases. Ten tumours were classified as a sexual cord or stromal tumours: five ovarian fibromas, four ovarian stromal tumours and one tumour diagnosed as an ovarian fibrosarcoma (sexual cord tumour with clear evidence of a fibrothecoma with malignant features). One bilateral immature teratoma<sup>27</sup> and one bilateral leiomyosarcoma<sup>39</sup> were previously reported. One relative was reported with a testicular fibrosarcoma at the age of 10 years. Overall, bilateral ovarian tumours were observed in six patients. These gonadal tumours mostly occurred at paediatric age, with a median age at diagnosis of 14 years (range 5–34). The four ovarian stromal tumours, which occurred in three girls, were bilateral in all the cases, synchronous for two of them and subsequent for the last patient. In cases where treatment data were available, surgery was performed in all, associated with chemotherapy in two.

Several other tumours were reported, as shown in table 1. No cardiac fibromas were observed.

### Risk of tumours in relatives

*SUFU* PV inheritance could be tested in 41 of 83 families. The PV was de novo in 11 patients (27%) and inherited in 30/41

**Table 2** Patients with secondary tumours after diagnosis of MB

Study	Status and age range at last follow-up	Age range at MB onset (histological type if available)	RT	Secondary malignancies	Other tumour(s)	Number of BCC
Smith <i>et al</i> <sup>8</sup> Evans <i>et al</i> <sup>19</sup>	Alive (30s)	Infant (desmoplastic)	Yes	BCC	Meningioma Pilocytic astrocytoma	65
Smith <i>et al</i> <sup>8</sup> Evans <i>et al</i> <sup>19</sup>	Alive (30s)	Infant (desmoplastic)	Yes	BCC	Meningioma	3
Smith <i>et al</i> <sup>8</sup> Evans <i>et al</i> <sup>19</sup>	Alive (20s)	Infant (desmoplastic)	Yes	BCC	Unilateral ovarian fibroma	11
Taylor <i>et al</i> <sup>14</sup> Ng <i>et al</i> <sup>23</sup>	Alive (20s)	NA (desmoplastic)	Yes	Meningioma		None
Kijima <i>et al</i> <sup>26</sup>	Alive (30s)	Infant	NA	BCC	Meningioma	NA
Mann <i>et al</i> <sup>28</sup>	Dead (childhood)	Infant (desmoplastic)	NA	BCC infundibulocystic		NA
Mann <i>et al</i> <sup>28</sup>	Dead (infant)	Infant	NA	Skin hamartoma		None
Guerrini-Rousseau <i>et al</i> <sup>31</sup>	Alive (10s)	Infant (MBEN)	No	Stromal ovarian tumour	Meningioma	None
Guerrini-Rousseau <i>et al</i> <sup>31</sup>	Dead (childhood)	Infant (MBEN)	Yes (PF only)	AML		None
Guerrini-Rousseau <i>et al</i> <sup>31</sup>	Alive (10s)	Infant (MBEN)	No	Thyroid carcinoma		None
Guerrini-Rousseau <i>et al</i> <sup>31</sup>	Alive (30s)	Infant (classic)	Yes (CSI)	BCC	Meningioma	>20
Guerrini-Rousseau <i>et al</i> <sup>31</sup>	Alive (10s)	Infant (desmoplastic)	No	Hamartoma		None
Ogden <i>et al</i> <sup>39</sup>	Alive (30s)	Infant	Yes (CSI)	Bilateral ovarian leiomyosarcoma	BCC Meningioma	>100
Present report	Alive (20s)	Infant	No	Bilateral ovarian stromal tumour		None
Present report	Alive (childhood)	Infant (desmoplastic)	No	Ovarian fibroma	Controlateral ovarian fibroma	None
Present report	Alive (10s)	Infant	Yes (CSI)	Meningioma	Meningioma	None
Present report	Alive (20s)	Infant (MBEN)	Yes (CSI)	Meningioma	Ovarian fibrosarcoma Thyroid carcinoma	None

Age range at last follow-up (2.2–37 years), age range at MB onset (0.5–2.5 years), age range at diagnosis of the other tumours (7–35 years)  
 AML, acute myeloid leukaemia; BCC, basal cell carcinoma; CSI, craniospinal radiotherapy; MB, medulloblastoma; MBEN, medulloblastoma with extensive nodularity; NA, not available; PF, posterior fossa; RT, radiotherapy.

families (73%). In these 30 families, 119 mutation carriers have been identified: 30 index and 89 relatives. We could analyse the occurrence of tumours in 89 relatives in whom the median age at last follow-up was 40.5 years (range 0.1–91). A total of 53 individuals (60%) were alive without a tumour at a median age of 37 years (range 1–91), and 36 patients (40%) developed a total of 55 tumours, some of them after the age of 50 years. The median age at the occurrence of the first tumour was 34 years. The cumulative incidence of any tumour at the age of 5, 20 and 50 years was, respectively, 14.4% (95% CI 6.8 to 21.4), 18.2% (95% CI 9.7 to 25.9) and 44.1% (95% CI 29.7 to 55.5) (figure 2A). The cumulative incidence of MB was 13.3% (95% CI 6 to 20.1) at 5 years and remained stable afterwards. The cumulative incidence of gonadal tumours was 2.8% (95% CI 0 to 6.6) and 4.6% (95% CI 0 to 9.7) at 20 and 50 years, respectively. The cumulative incidence of BCC and meningioma at 50 years was 28.5% (95% CI 13.4 to 40.9) and 5.2% (95% CI 0 to 12) (figure 2B).

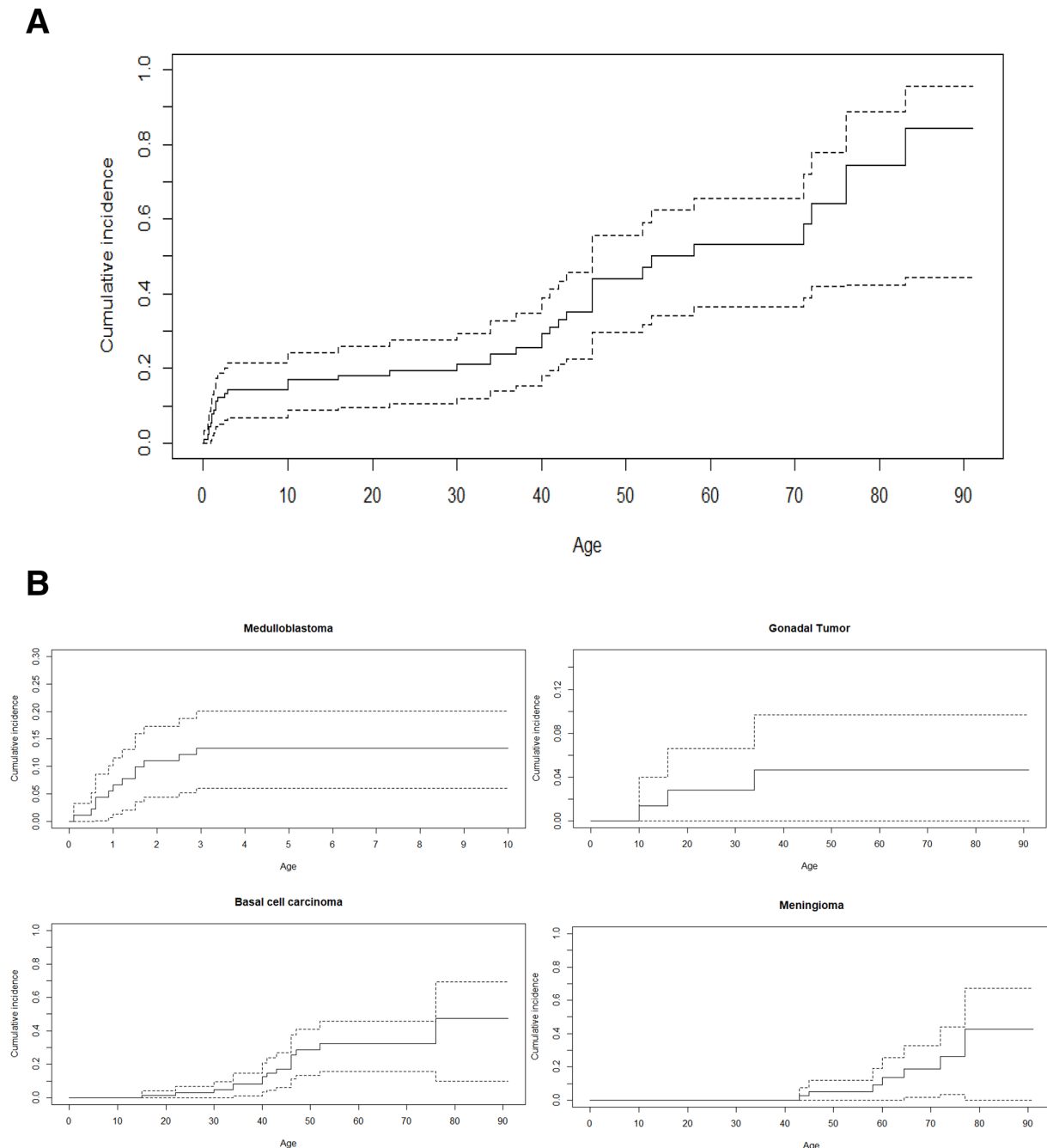
### Germline *SUFU* PV analysis

Overall, 64 different germline *SUFU* PVs were identified in 83 families, reported across the entire *SUFU* gene (figure 3). Nine different variants were identified in 28 families (online supplemental table 3). We identified 24 frameshift variants (37%), 9

nonsense variants (14%), 5 missense variants (8%), 19 splice site variants (30%) and 7 different structural variations involving either the complete *SUFU* gene, several exons or just one exon (11%). Nucleotide position 71 in the first exon (3 deletions of the cytosine and 5 duplications at that position) was found to be mutated in 8 families and the c.1022+1G>A variant which has been shown to result in the skipping of exon 8<sup>14</sup> was the splice PV the most frequently reported (in 7 families). Three patients with a severe intellectual disability have large structural variants that could be associated with a contiguous gene syndrome. There was not a significantly higher risk of MB according to the type of variation, the expected protein effects (structural variations, nonsense and frameshift PVs vs missense and splice PVs) ( $p=0.8728$ ,  $\chi^2$  test) or the involvement of the PV in the DNA binding domain of the *SUFU* gene ( $p=0.0827$ ,  $\chi^2$  test) (online supplemental table 4).

### DISCUSSION

Analysing together data from 48 unpublished patients and all cases previously reported allowed us to constitute a large series of germline *SUFU* PV carriers in which we could analyse tumour occurrence. In such a rare situation where presymptomatic testing is not proposed as a routine procedure, a precise evaluation of tumour risk cannot be performed. As most index patients in this



**Figure 2** Estimated cumulative incidence curves of tumours, among relatives carrying a *SUFU* pathogenic variant, according to the Nelson-Aalen estimator, with 95% CIs represented as dotted lines. (A) For all tumours. (B) For each main tumour type: medulloblastoma, basal cell carcinoma, meningioma and genital tumour.

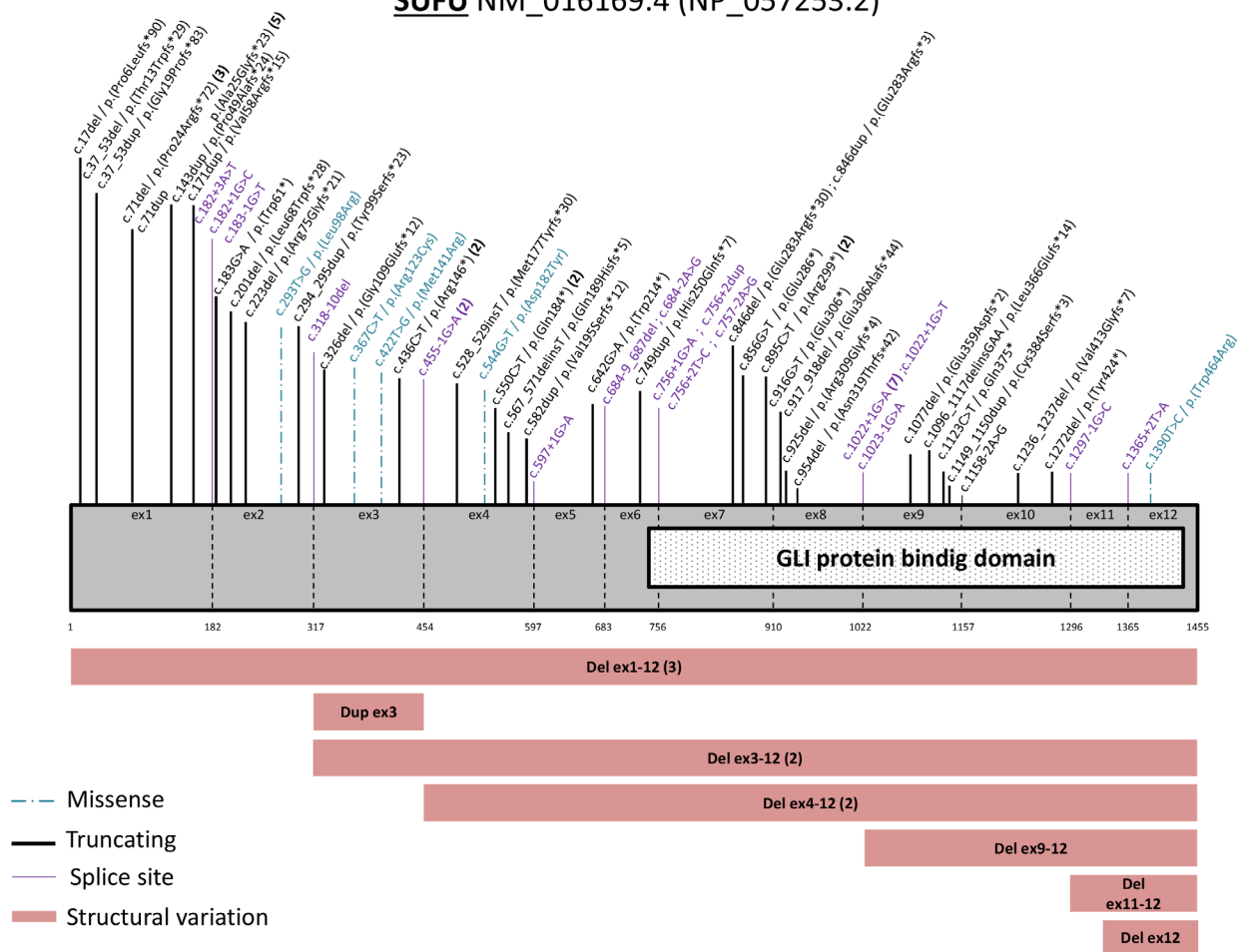
series have been identified after the occurrence of a malignancy, the evaluation of tumour risk on the entire population overestimates the risk. In order to reduce this ascertainment bias, we analysed the risk of tumour in relatives only after excluding the index cases. Nevertheless, a residual ascertainment bias related to the selection of relatives from families characterised by the presence of an affected case cannot be excluded. However, data from this series allow describing the oncological spectrum and risk associated with germline *SUFU* PV and the period of onset of each tumour type during life, which is the most important information for designing guidelines for PV carriers follow-up.

We confirmed that the tumour penetrance is high, although incomplete reaching 44% at 50 years. It cannot be compared

with *PTCH1* PV carriers since such an evaluation of cancer incidence in relatives is not available in this population. The overall tumour risk in the entire population, including index patients, reaches 68%. This is in the range of the risk described in *PTCH1*-associated GS, where tumour risk has been estimated around 55%–60%.<sup>19</sup> In addition, we could confirm that the spectrum of *SUFU*-associated GS differs from *PTCH1*-associated GS as previously suggested in smaller studies.<sup>19 31</sup>

The most frequent tumour is MB. It affects 89% of index patients. This rate is clearly higher than in patients carrying a germline *PTCH1* PV in whom MB incidence has been reported to be 2%.<sup>19</sup> The cumulative incidence of MB in relatives reaches 13.3%. This risk may be slightly overestimated since several

## SUFU NM\_016169.4 (NP\_057253.2)



**Figure 3** Representation of all the pathogenic or likely pathogenic variants described on the *SUFU* gene.

families were tested after the occurrence of MB in siblings but is lower than previously estimated.<sup>19,25</sup> As previously reported, all MBs were classified in the SHH subgroup<sup>20,34</sup> and most of them with desmoplastic/nodular histology.<sup>25</sup> We confirmed that the occurrence of MB was mainly limited to the first 3 years of life, with only 5/86 patients (5.8%) occurring after age 3 years but before 6 years.

Gonadal tumours mostly occur in children and teenagers with a median age at diagnosis of 14 years (range 5–34). In the Manchester cohort, systematic ultrasound in individuals with GS led to the detection of an ovarian tumour in 3/7 females (43%) with germline *SUFU* PV, compared with 4/68 females (5.9%) with *PTCH1* PV.<sup>19</sup> In the present series, the cumulative incidence of ovarian tumours has been estimated at around 10% at 50 years in females (4.6% in the entire population). The occurrence of malignant stromal tumours, which have not been reported yet in *PTCH1*-related GS where only fibromas have been reported, has to be underlined.

The cumulative incidence of meningioma at 50 years is about 5%. Nevertheless, meningioma seems to be more frequent in *SUFU* PV carriers (11%) than in the Manchester cohort of 126 patients with GS associated with *PTCH1* germline PVs in whom the incidence of meningiomas was <2%.<sup>19</sup> In contrast, the risk of BCC for *SUFU* PV carriers is clearly lower than for those with a *PTCH1* PV with only 11/31 (35%) *SUFU* patients >50 years affected with BCC. The cumulative incidence is 28.5% at 50 years of age in this series compared with 76.5%–80% at 50 years

in *PTCH1* PV carriers.<sup>3</sup> The occurrence of BCC or meningioma in germline *SUFU* PV carriers seems to be similar with a double peak occurrence of onset. As a first oncological event, apart from any previous treatment by radiotherapy or chemotherapy, the onset of these tumours seems to occur in a few patients (about 5%), mainly in adults around 40 years. In children treated for MB and exposed to chemotherapy and/or radiotherapy, the risk of BCC and meningioma is higher, and the onset is earlier (before the age of 30 years). It is noteworthy that neither odontogenic keratocysts nor cardiac fibromas occurred in the present series, with both tumours being hallmarks of *PTCH1*-associated GS.<sup>19</sup>

The risk of multiple tumours in *SUFU* PV carriers is clearly high, affecting 28% (33/117) of patients who developed a first tumour. The risk of second neoplasms, especially BCC and meningioma, after treatment of an MB can only be assessed for the first years following treatment since the follow-up is still short in most patients. However, since 12/23 (52%) patients aged >10 years at last follow-up after MB diagnosis have developed at least one secondary malignancy, this risk is clearly much higher than in an unselected series of MB in which a rate of secondary primary tumours of 3.1% at 10 years has been reported.<sup>40</sup> This high incidence warrants specific guidelines for the follow-up of these patients.<sup>41</sup> It is noteworthy that one-third of meningioma reported in this study occurred in patients previously treated with cranial radiotherapy for an MB. The 5-year OS of patients treated for an MB was 76% and is in the range of survival rates described in a large series of young children

either with nodular desmoplastic MB (5-year OS=89% and 81% for M0 and M+ patients, respectively)<sup>42</sup> or SHH-MB in infants (5-year OS=62%).<sup>43</sup> With the relatively short follow-up of the patients with MB in this cohort, most of the secondary tumours observed were not life-threatening. Except for one case, death was always linked to MB progression. Given the incidence of germline mutations in young patients with MB<sup>20</sup> and the consequences of the presence of a germline variant on care and follow-up, genetic testing for *SUFU* and *PTCH1* is of paramount importance in all children with SHH-MB before the age of 5 years. The best therapeutic strategy aiming to keep this high survival rate while sparing patients of the risk of second malignancy has to be evaluated. Upfront radiation sparing approaches could be justified given the expected high cure rate of SHH-MB in infants even in the absence of radiation.<sup>42</sup> The high risk of a secondary tumour in *SUFU* PV carriers treated for MB warrants early detection of BCC and meningioma, even in the absence of irradiation. Presymptomatic testing should be offered to the relatives during family genetic counselling to allow appropriate tumour surveillance.<sup>41</sup>

Because of this new insight into the *SUFU*-related cancer spectrum and risks requiring specific management, we think the clinical condition associated with *SUFU* PVs should be described as a specific syndrome requiring specific management. In contrast, the term 'GS' should be restricted to clinical manifestation associated with *PTCH1* PVs, acknowledging that these syndromes overlap. Testing for a germline *SUFU* mutations should be proposed in all patients presenting with a tumour belonging to *SUFU* spectrum (SHH-MB, BCC, meningioma, ovarian stromal or fibrous tumour) and for whom genetic predisposition is suspected because of young age at diagnosis, a familial history of cancer of multiple tumours. Recommendations for cancer surveillance in GS already published,<sup>4 44</sup> were recently adapted by the European Host Genome Working Group to the genetic background (*PTCH1* or *SUFU* PV) with the support of data presented here<sup>41</sup> (table 3). As the risk of MB is 13.3% at the age of 5 years, early postnatal testing for a *SUFU* PV can be offered because the result of the analysis will have a major impact on the surveillance of children with *SUFU* PV in the first years of life.

This large study also described molecular data of germline *SUFU* PVs and highlighted a certain level of recurrence for some

variants. Apart from structural variants that could be associated with a contiguous gene syndrome, no significant genotype-phenotype associations could be identified.

In conclusion, germline *SUFU* PV carriers have a life-long increased risk of tumours with a spectrum dominated by medulloblastoma before the age of 5 years, gonadal tumours during adolescence and BCC and meningioma in adulthood, justifying fine-tuned surveillance programmes, and the identification of healthy mutation carriers among relatives.

#### Author affiliations

<sup>1</sup>Department of Children and Adolescents Oncology, Gustave Roussy, Villejuif, France

<sup>2</sup>Team "Genomics and Oncogenesis of pediatric Brain Tumors"—Paris Saclay University, INSERM U981, VILLEJUIF, France

<sup>3</sup>INSERM U830, Laboratory of Translational Research in Pediatric Oncology, SIREDO Pediatric Oncology Center, Institute Curie, Paris, France

<sup>4</sup>Centre for Molecular Medicine Norway (NCMM), Nordic EMBL Partnership, University of Oslo and Oslo University Hospital, Oslo, Norway

<sup>5</sup>Department of Pediatric Research, Oslo University Hospital, Oslo, Norway

<sup>6</sup>Department of Medical and Clinical Genetics, University of Helsinki, Helsinki, Finland

<sup>7</sup>Département de Génétique et Institut Universitaire de Cancérologie, Sorbonne University Faculty of Médecine Pitié-Salpêtrière Campus, Paris, France

<sup>8</sup>Department of Pediatric Oncology and Hematology, University Hospitals Tübingen, Tübingen, Germany

<sup>9</sup>Department of Hematology/Oncology, Cell Therapy, Gene Therapy and Hemopoietic Transplant, IRCCS, Bambino Gesù Pediatric Hospital, Roma, Italy

<sup>10</sup>Department of Oncology, St Jude Children's Research Hospital, Memphis, Tennessee, USA

<sup>11</sup>Neuro-Oncology Unit, Department of Neurochirurgia, IRCCS Istituto Giannina Gaslini, Genova, Italy

<sup>12</sup>Neuro-Oncology Department, Princess Maxima Center for Pediatric Oncology, Utrecht, The Netherlands

<sup>13</sup>Institute of Human Genetics, University Hospital Heidelberg, Heidelberg, Germany

<sup>14</sup>Hopp Children's Cancer Center Heidelberg (KITZ), Heidelberg Health Center, Heidelberg, Germany

<sup>15</sup>Oncology, Great Ormond Street Hospital For Children NHS Foundation Trust, London, UK

<sup>16</sup>Paediatric Haematology and Oncology, Hannover Medical School, Hannover, Germany

<sup>17</sup>Section of Pediatric Hematology Oncology and Bone Marrow Transplantation, Alberta Children's Hospital and Cumming School of Medicine, University of Calgary, Calgary, Alberta, Canada

<sup>18</sup>Pediatric Hematology/Oncology and Stem Cells Transplantation, Bambino Gesù Pediatric Hospital, Roma, Italy

<sup>19</sup>Department of Oncology, Biology and Genetics, University of Genoa, Genoa, Italy

<sup>20</sup>Genetics of Rare Cancers, IRCCS Ospedale Policlinico San Martino, Genoa, Italy

<sup>21</sup>Division of Pediatric Neurooncology, DKFZ, Heidelberg, Germany

<sup>22</sup>Department of Pediatric Oncology, Hematology and Immunology, University Hospital Heidelberg, Heidelberg, Germany

<sup>23</sup>Institute of Medical Genetics and Applied Genomics, University of Tübingen Institute of Human Genetics, Tübingen, Germany

<sup>24</sup>Division of Evolution, Infection and Genomics, The University of Manchester, Manchester, UK

<sup>25</sup>Department of Medical and Clinical Genetics, Applied Tumor Genomics Research Program, University of Helsinki, Helsinki, Finland

<sup>26</sup>Department of Genetics, PSL Research University, Institute Curie, Paris, France

<sup>27</sup>Department of Genetics, Hôpital Universitaire des Enfants Reine Fabiola, ULB Center of Human Genetics, Université Libre de Bruxelles, Bruxelles, Belgium

<sup>28</sup>Department of Genetics, Hôpital Erasme, ULB Center of Human Genetics, Université Libre de Bruxelles, Bruxelles, Belgium

<sup>29</sup>CANSEARCH Research Platform, Department of pediatric oncology and hematology, University of Geneva, Geneva, Switzerland

<sup>30</sup>Childhood Cancer Research Group, Institute of Social and Preventive Medicine, University of Bern, Bern, Switzerland

<sup>31</sup>Department of Medicine, The Royal Melbourne Hospital, Parkville, Victoria, Australia

<sup>32</sup>Manchester Centre for Genomic Medicine, St Mary's Hospital, Manchester Academic Health Science Centre, School of Biological Sciences, Division of Evolution, Infection and Genomics, The University of Manchester, Manchester, UK

<sup>33</sup>Department of Children and Adolescents Oncology, Gustave Roussy Institute, Villejuif, France

**Twitter** Sandeep Kumar Dhanda @drsandeepdhanda and Roseline Vibert @RoselineVibert

**Table 3** Screening recommendations for patients with *SUFU* germline pathogenic variations

Tumours	Methods for screening	Surveillance recommendations in carriers of germline <i>SUFU</i> variants (periodicity, age to start and to end)
BCC	Dermatological examination	Annually ▶ beginning at age 20 years ▶ earlier if previous radiotherapy
Medulloblastoma	Brain MRI without contrast agent except for the first MRI	Every 3–4 months during the first 3 years then every 6 months until 5 years
Meningioma	Brain MRI	Every 3–5 years ▶ beginning at age 30 years for patients with no previous MB ▶ after healing of the MB in other patients
Ovarian tumour	Pelvic ultrasound	Every 3 years, beginning at age 5 years
Cardiac fibroma	Echocardiogram	At the time of diagnosis of GS, ideally in the first 6 months of life

BCC, basal cell carcinoma; GS, Gorlin syndrome; MB, medulloblastoma.



**Acknowledgements** This project was additionally supported by 'la Fondation Gustave Roussy', the Italian Association for Cancer Research (AIRC), the PedBrain Tumor Project contributing to the International Cancer Genome Consortium (ICGC), funded by the German Cancer Aid (109252), the German Federal Ministry of Education and Research (BMBF) (01KU1201A, 01KU1201C) and the BMBF grants BioTop (01EK1502A, 01EK1502B), ICGC-DE-Mining (01KU1505F), MedSys (0315416C) and NGFNplus (01GS0883).

**Contributors** LG-R, SMW, FB, PRB, IBB, GDB, SKD, MLG, CEMG, SH, MJ, CK, LL-C, AM, LP, SMP, MJS, PV, CV, NW, IMW, DGE and LB provided clinical and molecular data. LG-R and LB reviewed literature and compiled clinical data. JM-P and SKD analysed the genetic data. LG-R and LB contributed to the study design. RV, LG-R and LB provided the statistical analysis with N and A estimation method. LG-R, PH and LB are in charge of the PREDCAP database. All authors contributed to the article writing, reviewing and editing.

**Funding** LB and LG-R have been supported by la Fondation Gustave Roussy campaign: Guérir Le Cancer de l'Enfant au 21<sup>ème</sup> siècle. DGE and MJS are supported by the National Institute for Health Research (NIHR) BRC Manchester (Grant Reference Number 1215-200074). CPK and SMP have been supported by the Deutsche Kinderkrebsstiftung (DKS2019.13).

**Competing interests** None declared.

**Patient consent for publication** Not applicable.

**Ethics approval** This study was approved by an Ethics Committee(s) or Institutional Board(s): CEEI Inserm/IRB00003888. Written informed consent was obtained for the genetic analysis according to local good clinical practice guidelines. Consent for data collection was obtained from the parents or guardians, according to each national ethics requirements.

**Provenance and peer review** Not commissioned; externally peer reviewed.

**Data availability statement** All data relevant to the study are included in the article or uploaded as supplementary information.

**Supplemental material** This content has been supplied by the author(s). It has not been vetted by BMJ Publishing Group Limited (BMJ) and may not have been peer-reviewed. Any opinions or recommendations discussed are solely those of the author(s) and are not endorsed by BMJ. BMJ disclaims all liability and responsibility arising from any reliance placed on the content. Where the content includes any translated material, BMJ does not warrant the accuracy and reliability of the translations (including but not limited to local regulations, clinical guidelines, terminology, drug names and drug dosages), and is not responsible for any error and/or omissions arising from translation and adaptation or otherwise.

**Open access** This is an open access article distributed in accordance with the Creative Commons Attribution 4.0 Unported (CC BY 4.0) license, which permits others to copy, redistribute, remix, transform and build upon this work for any purpose, provided the original work is properly cited, a link to the licence is given, and indication of whether changes were made. See: <https://creativecommons.org/licenses/by/4.0/>.

#### ORCID iDs

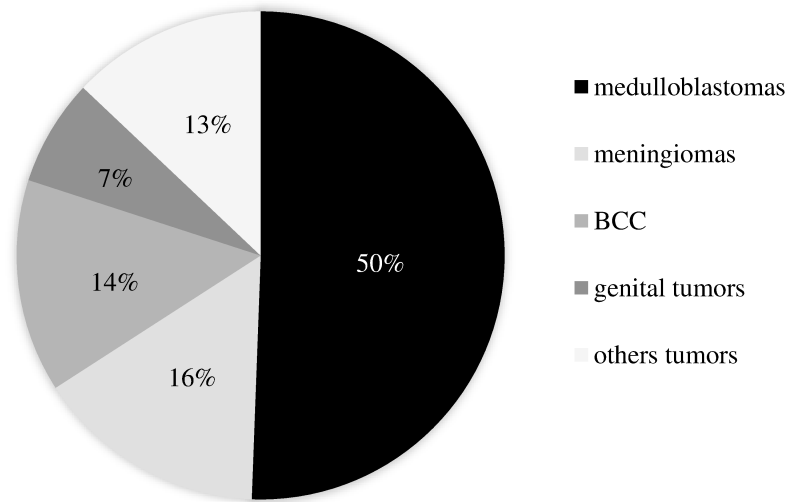
Léa Guerrini-Rousseau <http://orcid.org/0000-0003-0050-5407>  
Sandeep Kumar Dhanda <http://orcid.org/0000-0003-1381-7434>  
Miriam Jane Smith <http://orcid.org/0000-0002-3184-0817>  
Roseline Vibert <http://orcid.org/0000-0001-5202-0212>  
Nicolas Waespe <http://orcid.org/0000-0002-2271-8959>

#### REFERENCES

- 1 Kimonis VE, Goldstein AM, Pastakia B, Yang ML, Kase R, DiGiovanna JJ, Bale AE, Bale SJ. Clinical manifestations in 105 persons with nevoid basal cell carcinoma syndrome. *Am J Med Genet* 1997;69:299–308.
- 2 Shanley S, McCormack C. Diagnosis and management of hereditary basal cell skin cancer. *Recent Results Cancer Res* 2016;205:191–212.
- 3 Jones EA, Sajid MI, Shenton A, Evans DG. Basal cell carcinomas in Gorlin syndrome: a review of 202 patients. *J Skin Cancer* 2011;2011:217378.
- 4 Bree AF, Shah MR, for the BCNS Colloquium Group. Consensus statement from the first international Colloquium on basal cell nevus syndrome (BCNS). *Am J Med Genet A* 2011;155:2091–7.
- 5 Evans DG, Fardon PA. Nevoid Basal Cell Carcinoma Syndrome. In: Adam MP, Ardinger HH, Pagon RA, Wallace SE, Bean LJ, Mirzaz G, eds. Seattle (WA): GeneReviews® University of Washington, Seattle, 1993.
- 6 Johnson RL, Rothman AL, Xie J, Goodrich LV, Bare JW, Bonifas JM, Quinn AG, Myers RM, Cox DR, Epstein EH, Scott MP. Human homolog of *patched*, a candidate gene for the basal cell nevus syndrome. *Science* 1996;272:1668–71.
- 7 Hahn H, Wicking C, Zaphiropoulos PG, Gailani MR, Shanley S, Chidambaram A, Vorechovsky I, Holmberg E, Uden AB, Gillies S, Negus K, Smyth I, Pressman C, Leffell DJ, Gerrard B, Goldstein AM, Dean M, Toftgard R, Chenevix-Trench G, Wainwright B, Bale AE. Mutations of the human homolog of *Drosophila patched* in the nevoid basal cell carcinoma syndrome. *Cell* 1996;85:841–51.
- 8 Smith MJ, Beetz C, Williams SG, Bhaskar SS, O'Sullivan J, Anderson B, Daly SB, Urquhart JE, Bholah Z, Oudit D, Cheesman E, Kelsey A, McCabe MG, Newman WG, Evans DGR. Germline Mutations in *SUFU* Cause Gorlin Syndrome—Associated Childhood Medulloblastoma and Redefine the Risk Associated With *PTCH1* Mutations. *JCO* 2014;32:4155–61.
- 9 Pastorino L, Ghiorzo P, Nasti S, Battistuzzi L, Cusano R, Marzocchi C, Garrè ML, Clementi M, Scarrà GB. Identification of a *SUFU* germline mutation in a family with Gorlin syndrome. *Am J Med Genet A* 2009;149A:1539–43.
- 10 Begemann M, Waszak SM, Robinson GW, Jäger N, Sharma T, Knopp C, Kraft F, Moser O, Mynarek M, Guerrini-Rousseau L, Brugières L, Varlet P, Pietsch T, Bowers DC, Chintagumpala M, Sahn F, Korbel JO, Rutkowski S, Eggermann T, Gajjar A, Northcott P, Elbracht M, Pfister SM, Kontny U, Kurth I. Germline *GPR161* Mutations Predispose to Pediatric Medulloblastoma. *J Clin Oncol* 2020;38:43–50.
- 11 Fan Z, Li J, Du J, Zhang H, Shen Y, Wang C-Y, Wang S. A missense mutation in *PTCH2* underlies dominantly inherited NBCCS in a Chinese family. *J Med Genet* 2008;45:303–8.
- 12 Fujii K, Ohashi H, Suzuki M, Hatsuse H, Shiohama T, Uchikawa H, Miyashita T. Frameshift mutation in the *PTCH2* gene can cause nevoid basal cell carcinoma syndrome. *Fam Cancer* 2013;12:611–4.
- 13 Smith MJ, Evans DG. *Ptch2* is not a strong candidate gene for Gorlin syndrome predisposition. *Fam Cancer* 2021;32.
- 14 Taylor MD, Liu L, Raffel C, Hui C-chung, Mainprize TG, Zhang X, Agatep R, Chiappa S, Gao L, Lowrance A, Hao A, Goldstein AM, Stavrou T, Scherer SW, Dura WT, Wainwright B, Squire JA, Rutka JT, Hogg D. Mutations in *Sufu* predispose to medulloblastoma. *Nat Genet* 2002;31:306–10.
- 15 Smith MJ. Germline and somatic mutations in meningiomas. *Cancer Genet* 2015;208:107–14.
- 16 Pathmanaban ON, Sadler KV, Kamaly-Asl ID, King AT, Rutherford SA, Hammerbeck-Ward C, McCabe MG, Kilday J-P, Beetz C, Poplawski NK, Evans DG, Smith MJ. Association of genetic predisposition with solitary schwannoma or meningioma in children and young adults. *JAMA Neurol* 2017;74:1123–9. 01.
- 17 Gorlin RJ, Goltz RW. Multiple nevoid basal-cell epithelioma, jaw cysts and bifid rib. A syndrome. *N Engl J Med* 1960;262:908–12.
- 18 Fogel AL, Sarin KY, Teng JMC. Genetic diseases associated with an increased risk of skin cancer development in childhood. *Curr Opin Pediatr* 2017;29:426–33.
- 19 Evans DG, Oudit D, Smith MJ, Rutkowski D, Allan E, Newman WG, Lear JT. First evidence of genotype-phenotype correlations in Gorlin syndrome. *J Med Genet* 2017;54:530–6.
- 20 Waszak SM, Northcott PA, Buchhalter I, Robinson GW, Sutter C, Groebner S, Grund KB, Brugières L, Jones DTW, Pajtler KW, Morrissy AS, Kool M, Sturm D, Chavez L, Ernst A, Brabetz S, Hain M, Zichner T, Segura-Wang M, Weisenfeldt J, Rausch T, Mardin BR, Zhou X, Baciu C, Lawrenz C, Chan JA, Varlet P, Guerrini-Rousseau L, Fults DW, Grajkowska W, Hauser P, Jabado N, Ra Y-S, Zitterbart K, Shringarpure SS, De La Vega FM, Bustamante CD, Ng H-K, Perry A, MacDonald TJ, Hernáiz Driever P, Bendel AE, Bowers DC, McCowage G, Chintagumpala MM, Cohn R, Hassall T, Fleischhack G, Eggen T, Wesenberg F, Feychting M, Laner B, Schüz J, Johansen C, Andersen TV, Rööslil M, Kuehni CE, Grotzer M, Kjaerheim K, Monoranu CM, Archer TC, Duke E, Pomeroy SL, Shelagh R, Frank S, Sumerauer D, Scheuren W, Ryzhova MV, Milde T, Kratz CP, Samuel D, Zhang J, Solomon DA, Marra M, Eils R, Bartram CR, von Hoff K, Rutkowski S, Ramaswamy V, Gilbertson RJ, Korshunov A, Taylor MD, Lichter P, Malkin D, Gajjar A, Korbel JO, Pfister SM. Spectrum and prevalence of genetic predisposition in medulloblastoma: a retrospective genetic study and prospective validation in a clinical trial cohort. *Lancet Oncol* 2018;19:785–98.
- 21 Kool M, Jones DTW, Jäger N, Northcott PA, Pugh TJ, Hovestadt V, Piro RM, Esparza LA, Markant SL, Remke M, Milde T, Bourdeaut F, Ryzhova M, Sturm D, Pfaff E, Stark S, Hutter S, Seker-Cin H, Johann P, Bender S, Schmidt C, Rausch T, Shih D, Reimand J, Sieber L, Wittmann A, Linke L, Witt H, Weber UD, Zaparka M, König R, Beroukhim R, Berghold G, van Sluis P, Volckmann R, Koster J, Versteeg R, Schmidt S, Wolf S, Lawrenz C, Bartholomae CC, von Kalle C, Unterberg A, Herold-Mende C, Hofer S, Kulozik AE, von Deimling A, Scheuren W, Felsberg J, Reifemberger G, Hasselblatt M, Crawford JR, Grant GA, Jabado N, Perry A, Cowdrey C, Croul S, Zadeh G, Korbel JO, Doz F, Delattre O, Bader GD, McCabe MG, Collins VP, Kieran MW, Cho Y-J, Pomeroy SL, Witt O, Brors B, Taylor MD, Schüller U, Korshunov A, Eils R, Wechsler-Reya RJ, Lichter P, Pfister SM, ICGC PedBrain Tumor Project. Genome sequencing of Shh medulloblastoma predicts genotype-related response to smoothened inhibition. *Cancer Cell* 2014;25:393–405.
- 22 Cavalli FMG, Remke M, Rampasek L, Peacock J, Shih DJH, Luu B, Garzia L, Torchia J, Nor C, Morrissy AS, Agnihotri S, Thompson YY, Kuzan-Fischer CM, Faraoo H, Isaev K, Daniels C, Cho BK, Kim SK, Wang KC, Lee JY, Grajkowska WA, Perek-Polnik M, Vasiljevic A, Faure-Conter C, Jouveta A, Giannini C, Nageswara Rao AA, Li KKW NHK, Eberhart CG, Pollack IF, Hamilton RL, Gillespie GY, Olson JM, Leary S, Weiss WA, Lach B, Chambless LB, Thompson RC, Cooper MK, Vibhakar R, Hauser P, van Veelen MC, Kros JM, French PJ, YS R, Kumabe T, López-Aguilar E, Zitterbart K, Sterba J, Finocchiaro G, Massimino M, Van Meir EG, Osuka S, Shofuda T, Klekner A, Zollo M, Leonard JR, Rubin JB, Jabado N, Albrecht S, Mora J, Van Meter TE, Jung S, Moore AS, Hallahan AR, Chan JA, Tirapelli DPC, Carlotti CG, Fouladi M, Pimentel J, Faria CC, Saad AG,

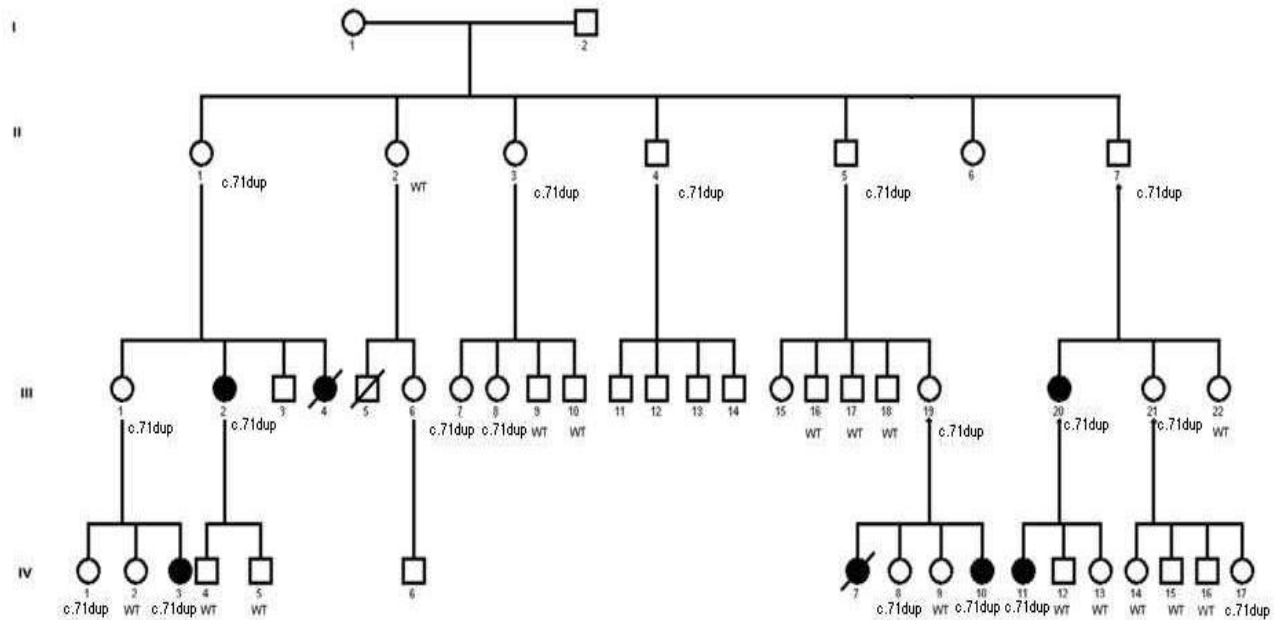
- Massimi L, Liou LM, Wheeler H, Nakamura H, Elbabaa SK, Perezpeña-Diazconti M, Chico Ponce de León F, Robinson S, Zapotocky M, Lassaletta a, Huang a, Hawkins Ce, Tabori U, Bouffet E, Bartels U, Dirks Pb, Rutka JT, Bader Gd, Reimand J, Goldenberg a, Ramaswamy V, Taylor MD. Intertumoral heterogeneity within medulloblastoma subgroups. *Cancer Cell* 2017;31:737–54.
- 23 Ng D, Stavrou T, Liu L, Taylor MD, Gold B, Dean M, Kelley MJ, Dubovsky EC, Vezina G, Nicholson HS, Byrne J, Rutka JT, Hogg D, Reaman GH, Goldstein AM. Retrospective family study of childhood medulloblastoma. *Am J Med Genet A* 2005;134:399–403.
- 24 Slade I, Murray A, Hanks S, Kumar A, Walker L, Hargrave D, Douglas J, Stiller C, Izatt L, Rahman N. Heterogeneity of familial medulloblastoma and contribution of germline PTCH1 and Sufu mutations to sporadic medulloblastoma. *Fam Cancer* 2011;10:337–42.
- 25 Brugières L, Remenieras A, Pierron G, Varlet P, Forget S, Byrde V, Bombled J, Puget S, Caron O, Dufour C, Delattre O, Bressac-de Paillerets B, Grill J. High frequency of germline Sufu mutations in children with desmoplastic/nodular medulloblastoma younger than 3 years of age. *J Clin Oncol* 2012;30:2087–93.
- 26 Kijima C, Miyashita T, Suzuki M, Oka H, Fujii K. Two cases of nevoid basal cell carcinoma syndrome associated with meningioma caused by a PTCH1 or Sufu germline mutation. *Fam Cancer* 2012;11:565–70.
- 27 Aavikko M, Li S-P, Saarinen S, Alhopuro P, Kaasinen E, Morgunova E, Li Y, Vesanen K, Smith MJ, Evans DGR, Pöyhönen M, Kiuru A, Auvinen A, Aaltonen LA, Taipale J, Vahteristo P. Loss of Sufu function in familial multiple meningioma. *Am J Hum Genet* 2012;91:520–6.
- 28 Mann K, Magee J, Guillaud-Bataille M, Blondel C, Bressac-de Paillerets B, Yeatman J, Winship I. Multiple skin hamartomata: a possible novel clinical presentation of Sufu neoplasia syndrome. *Fam Cancer* 2015;14:151–5.
- 29 Šoukalová J, Vejmelková K, Cermanová T, Kašiková K, Mikulášková A, Janyšková H, Melichárková K, Pavelka Z, Ježová M, Pospíšilová Š, Kuglík P, Valášková I, Gaillyová R, Štěrba J, Zitterbart K. [Identification of a Family with SUFU Germline Deletion Based on a Case of Desmoplastic Medulloblastoma in an Infant]. *Klin Onkol* 2016;29 Suppl 1:S83–8.
- 30 Schulman JM, Oh DH, Sanborn JZ, Pincus L, McCalmont TH, Cho RJ. Multiple Hereditary Infundibulocystic Basal Cell Carcinoma Syndrome Associated With a Germline *SUFU* Mutation. *JAMA Dermatol* 2016;152:323–7.
- 31 Guerrini-Rousseau L, Dufour C, Varlet P, Masliah-Planchon J, Bourdeaut F, Guillaud-Bataille M, Abbas B, Bertozzi A-I, Fouyssac F, Huybrechts S, Puget S, Bressac-De Paillerets B, Caron O, Sevenet N, Dimaria M, Villebasse S, Delattre O, Valteau-Couanet D, Grill J, Brugières L. Germline *SUFU* mutation carriers and medulloblastoma: clinical characteristics, cancer risk, and prognosis. *Neuro Oncol* 2018;20:1122–32.
- 32 Askaner G, Lei U, Bertelsen B, Venzo A, Wadt K. Novel *SUFU* Frameshift Variant Leading to Meningioma in Three Generations in a Family with Gorlin Syndrome. *Case Rep Genet* 2019;2019:9650184.
- 33 Skaro M, Nanda N, Gauthier C, Felsenstein M, Jiang Z, Qiu M, Shindo K, Yu J, Hutchings D, Javed AA, Beckman R, He J, Wolfgang CL, Thompson E, Hruban RH, Klein AP, Goggins M, Wood LD, Roberts NJ. Prevalence of germline mutations associated with cancer risk in patients with intraductal papillary mucinous neoplasms. *Gastroenterology* 2019;156:1905–13.
- 34 Lafay-Cousin L, Bouffet E, Strother D, Rudneva V, Hawkins C, Eberhart C, Horbinski C, Heier L, Souweidane M, Williams-Hughes C, Onar-Thomas A, Billups CA, Fouladi M, Northcott P, Robinson G, Gajjar A. Phase II Study of Nonmetastatic Desmoplastic Medulloblastoma in Children Younger Than 4 Years of Age: A Report of the Children's Oncology Group (ACNS1221). *JCO* 2020;38:223–31.
- 35 Wang Y, Wu J, Li W, Li J, Liu R, Yang B, Li C, Jiang T. Retrospective investigation of hereditary syndromes in patients with medulloblastoma in a single institution. *Childs Nerv Syst* 2021;37:411–7.
- 36 Robinson GW, Orr BA, Wu G, Gururangan S, Lin T, Qaddoumi I, Packer RJ, Goldman S, Prados MD, Desjardins A, Chintagumpala M, Takebe N, Kaste SC, Rusch M, Allen SJ, Onar-Thomas A, Stewart CF, Fouladi M, Boyett JM, Gilbertson RJ, Curran T, Ellison DW, Gajjar A. Vismodegib exerts targeted efficacy against recurrent sonic Hedgehog-Subgroup medulloblastoma: results from phase II pediatric brain tumor Consortium studies PBTC-025B and PBTC-032. *J Clin Oncol* 2015;33:2646–54.
- 37 Hobbs BP. On nonparametric hazard estimation. *J Biom Biostat* 2015;6:232.
- 38 Huq AJ, Walsh M, Rajagopalan B, Finlay M, Trainer AH, Bonnet F, Sevenet N, Winship IM. Mutations in Sufu and PTCH1 genes may cause different cutaneous cancer predisposition syndromes: similar, but not the same. *Fam Cancer* 2018;17:601–6.
- 39 Ogden T, Higgins S, Elbaum D, Wysong A. The relevance of a suppressor of fused (Sufu) mutation in the diagnosis and treatment of Gorlin syndrome. *JAAD Case Rep* 2018;4:196–9.
- 40 Nantavithya C, Paulino AC, Liao K, McGovern SL, Grosshans DR, McAleer MF, Woodhouse KD, Khatua S, Chintagumpala MM, Majd NK, Yeboa DN. Development of second primary tumors and outcomes in medulloblastoma by treatment modality: a surveillance, epidemiology, and end results analysis. *Pediatr Blood Cancer* 2020;67:e28373.
- 41 Guerrini-Rousseau L, Smith MJ, Kratz CP, Doergeloh B, Hirsch S, Hopman SMJ, Jorgensen M, Kuhlen M, Michaeli O, Milde T, Ridola V, Russo A, Salvador H, Waespe N, Claret B, Brugières L, Evans DG. Current recommendations for cancer surveillance in Gorlin syndrome: a report from the SIOPE host genome Working Group (SIOPE HGWG). *Fam Cancer* 2021;20:317–25.
- 42 Dhall G, O'Neil SH, Ji L, Haley K, Whitaker AM, Nelson MD, Gilles F, Gardner SL, Allen JC, Cornelius AS, Pradhan K, Garvin JH, Olshefski RS, Hukin J, Comito M, Goldman S, Atlas MP, Walter AW, Sands S, Spoto R, Finlay JL. Excellent outcome of young children with nodular desmoplastic medulloblastoma treated on "Head Start" III: a multi-institutional, prospective clinical trial. *Neuro Oncol* 2020;22:1862–72.
- 43 Schwalbe EC, Lindsey JC, Nakjang S, Crosier S, Smith AJ, Hicks D, Rafiee G, Hill RM, Iliasova A, Stone T, Pizer B, Michalski A, Joshi A, Wharton SB, Jacques TS, Bailey S, Williamson D, Clifford SC. Novel molecular subgroups for clinical classification and outcome prediction in childhood medulloblastoma: a cohort study. *Lancet Oncol* 2017;18:958–71.
- 44 Foulkes WD, Kamihara J, Evans DGR, Brugières L, Bourdeaut F, Molenaar JJ, Walsh MF, Brodeur GM, Diller L. Cancer surveillance in Gorlin syndrome and rhabdoid tumor predisposition syndrome. *Clin Cancer Res* 2017;23:e62–7.

# Figure S1



**Figure S1.** Distribution of tumor types in the cohort

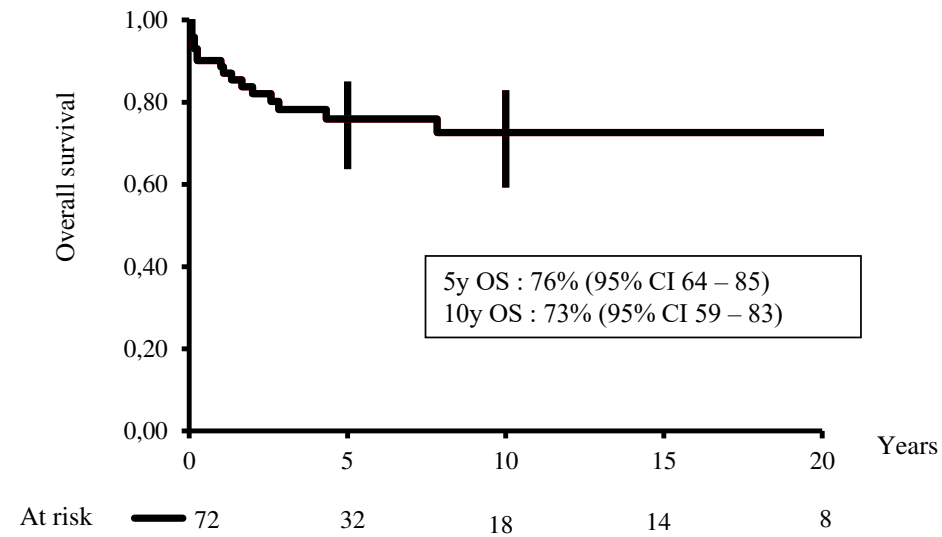
# Figure S2



**Figure S2.** Genealogical table of one family with an history of 5 children (III-2, III-20, IV-3, IV-10 and IV-11) with a documented germline *SUFU* PV affected with a proven medulloblastoma (Family 3)

Legend: Girls affected with medulloblastoma (probable or proven) are shown with filled circles.

# Figure S3



**Figure S3.** Overall survival after a medulloblastoma of children with germline *SUFU* PV (n=72).

# Table S1

Family	c. mutation type	Index patient or relative	MB subgroup	Age at diagnosis (years)
Family 1	c.37_53dup	Index patient	MB,SHH	0.4
		Relative (sibling)	MB (NA)	NA
Family 2*	c.756+1G>A	Relative (sibling)	ND MB	0.6
		Relative (sibling)	MB (NA)	0.5
Family 3	c.71dup	Index patient	MBEN	0.9
		Relative (cousin)	ND MB	2.9
		Relative (cousin)	ND MB	1.7
		Relative (aunt)	MB (NA)	0.6
		Relative (cousin)	MB (NA)	1.2
Family 4	c.71del	Index patient	MBEN	0.3
		Relative (sibling)	MBEN	0.1
Family 5	c.567_571delinsT	Index patient	MBEN	2.2
		Relative (sibling)	Classic MB	2.5
		Relative (sibling)	Classic MB	1.5
Family 6	c.201delC	Index patient	MB (NA)	2.8
		Relative (cousin)	MB (NA)	0.9

**Table S1.** Family history with multiple children with a germline *SUFU* PV affected with medulloblastoma in the same family

\* Index patient affected with BCC (28)

# Table S2

**Table S2.** Description of the various *SUFU* PVs identified in the cohort (NM\_016169.4)

publication	mutation c	mutation p	Mutation type	Recurrence in n families
(9,24,31, 34) (5 families) and present report (2 families)	c.1022+1G>A	p.?	Splicing	7 families
(31,35) (2 families) and present report (3 families)	c.71dup	p.(Ala25Glyfs*23)	Frameshift	5 families
(20,31) (3 families)	c.71del	p.(Pro24Argfs*72)	Frameshift	3 families
(14,29) and present report	c.1-? 1455+?del	del exon 1-12	Structural variation	3 families
(8,19, 31)	c.318-? 1455+?del	del exon 3-12	Structural variation	2 families
(8,19,26)	c.550C>T	p.(Gln184*)	Nonsense	2 families
(20) and present report	c.436C>T	p.(Arg146*)	Nonsense	2 families
(20) and present report	c.455-1G>A	p.?	Splicing	2 families
(20) and present report	c.895C>T	p.(Arg299*)	Nonsense	2 families
(8,19)	c.544G>T	p.(Asp182Tyr)	Missense	1 family
(20)	c.1077del	p.(Glu359Aspfs*2)	Frameshift	1 family
(20)	c.37_53del	p.(Thr13Trpfs*29)	Frameshift	1 family
(20)	c.37_53dup	p.(Gly19Profs*83)	Frameshift	1 family
(20)	c.642G>A	p.(Trp214*)	Nonsense	1 family
(20)	c.684-2A>G	p.?	Splicing	1 family
(20)	c.749dup	p.(His250Glnfs*7)	Frameshift	1 family
(20)	c.925del	p.(Arg309Glyfs*4)	Frameshift	1 family
(14,23)	c.143dup	p.(Pro49Alafs*24)	Frameshift	1 family
(14,23)	c.183-1G>T	p.?	Splicing	1 family
(24)	c.846dup	p.(Glu283Argfs*3)	Frameshift	1 family
(27)	c.367C>T	p.(Arg123Cys)	Missense	1 family
(28)	c.756+1G>A	p.?	Splicing	1 family
(30)	c.757-2A>G	p.?	Splicing	1 family
(31)	c.1023-? 1455+?del	del exon 9-12	Structural variation	1 family
(31)	c.1096_1117delinsGAA	p.(Leu366Glufs*14)	Frameshift	1 family
(31)	c.1123C>T	p.(Gln375*)	Nonsense	1 family
(31)	c.1149_1150dup	p.(Cys384Serfs*3)	Frameshift	1 family
(31)	c.1297-1G>C	p.?	Splicing	1 family
(31)	c.182+3A>T	p.?	Splicing	1 family
(31)	c.294_295dup	p.(Tyr99Serfs*23)	Frameshift	1 family
(31)	c.318-? 454+?dup	dup exon 3	Structural variation	1 family
(31)	c.318-10del	p.?	Splicing	1 family
(31)	c.422T>G	p.(Met141Arg)	Missense	1 family
(31)	c.567_571delinsT	p.(Gln189Hisfs*5)	Frameshift	1 family
(32)	c.954del	p.(Asn319Thrfs*42)	Frameshift	1 family
(33)	c.223del	p.(Arg75Glyfs*21)	Frameshift	1 family
(34)	c.1023-1G>A	p.?	Splicing	1 family
(34)	c.171dup	p.(Val58Argfs*15)	Frameshift	1 family
(34)	c.528_529insT	p.(Met177Tyrfs*30)	Frameshift	1 family
(34)	c.684-9_687del	p.?	Splicing	1 family
(34)	c.916G>T	p.(Glu306*)	Nonsense	1 family
(36)	c.1272del	p.(Tyr424*)	Nonsense	1 family
(38)	c.1365+2T>A	p.Ile433_Glu455del	Splicing	1 family
(39)	c.597+1dupG	p.?	Splicing	1 family
Present report	c.1022+1G>T	p.?	Splicing	1 family
Present report	c.1158-2A>G	p.?	Splicing	1 family
Present report	c.1236_1237del	p.(Val413Glyfs*7)	Frameshift	1 family
Present report	c.1297-? 1455+?del	del exon 11-12	Structural variation	1 family
Present report	c.1366-? 1455+?del	del exon 12	Structural variation	1 family
Present report	c.1390T>C	p.(Trp464Arg)	Missense	1 family
Present report	c.17del	p.(Pro6fs*89)	Frameshift	1 family
Present report	c.182+1G>C	p.?	Splicing	1 family
Present report	c.183G>A	p.(Trp61*)	Nonsense	1 family
Present report	c.201del	p.(Leu68Trpfs*28)	Frameshift	1 family
Present report	c.293T>G	p.(Leu98Arg)	Missense	1 family
Present report	c.326del	p.(Gly109Glufs*12)	Frameshift	1 family
Present report	c.455-? 1455+?del	del exon 4-12	Structural variation	1 family
Present report	c.582dup	p.(Val195Serfs*12)	Frameshift	1 family
Present report	c.597+1G>A	p.?	Splicing	1 family
Present report	c.756+2dup	p.?	Splicing	1 family
Present report	c.756+2T>C	p.?	Splicing	1 family
Present report	c.846del	p.(Glu283Argfs*30)	Frameshift	1 family
Present report	c.856G>T	p.(Glu286*)	Nonsense	1 family
Present report	c.917_918del	p.(Glu306Alafs*44)	Frameshift	1 family

# Table S3

	<b>p</b>
Frameshift (n=84) versus other (n=88)	0.169796
Missense (n=14) versus other (n=158)	0.163241
Nonsense (n=14) versus other (n=158)	0.050943
Splice (n=46) versus other (n=126)	0.605297
Structural variation (n=14) versus other (n=158)	0.402844
Structural variation, nonsense or frameshift PVs (n=112) versus missense or splicing PVs (n=60)	0.872893

**Table S3.** Risk of medulloblastoma according to the *SUFU* PV. Chi-square test depending on the type of variation or the expected protein effects of the *SUFU* gene

p-values <0.05 were considered statistically significant

PVs : pathogenic variants