



Nervous system (NS) Tumors in Cancer Predisposition Syndromes

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Abstract

Genetic syndromes which develop one or more nervous system (NS) tumors as one of the manifestations can be grouped under the umbrella term of NS tumor predisposition syndromes. Understanding the underlying pathological pathways at the molecular level has led us to many radical discoveries, in understanding the mechanisms of tumorigenesis, tumor progression, interactions with the tumor microenvironment, and development of targeted therapies. Currently, at least 7–10% of all pediatric cancers are now recognized to occur in the setting of genetic predisposition to cancer or cancer predisposition syndromes. Specifically, the cancer predisposition rate in pediatric patients with NS tumors has been reported to be as high as 15%, though it can approach 50% in certain tumor types (i.e., choroid plexus carcinoma associated with Li Fraumeni Syndrome). Cancer predisposition syndromes are caused by pathogenic variation in genes that primarily function as tumor suppressors and proto-oncogenes. These variants are found in the germline or constitutional DNA. Mosaicism, however, can affect only certain tissues, resulting in varied manifestations. Increased understanding of the genetic underpinnings of cancer predisposition syndromes and the ability of clinical laboratories to offer molecular genetic testing allows for improvement in the identification of these patients. The identification of a cancer predisposition syndrome in a CNS tumor patient allows for changes to medical management to be made, including the initiation of cancer surveillance protocols. Finally, the identification of at-risk biologic relatives becomes feasible through cascade (genetic) testing. These fundamental discoveries have also broadened the horizon of novel therapeutic possibilities and have helped to be better predictors of prognosis and survival. The treatment paradigm of specific NS tumors may also vary based on the patient's cancer predisposition syndrome and may be used to guide therapy (i.e., immune checkpoint inhibitors in constitutional mismatch repair deficiency [CMMRD] predisposition syndrome) [8]. Early diagnosis of these cancer predisposition syndromes is therefore critical, in both unaffected and affected patients. Genetic counselors are uniquely trained master's level healthcare providers with a focus on the identification of hereditary disorders, including hereditary cancer, or cancer predisposition syndromes. Genetic counseling, defined as “the process of helping people understand and adapt to the medical, psychological and familial implications of genetic contributions to disease” plays a vital role in the adaptation to a genetic diagnosis and the overall management of these diseases. Cancer predisposition syndromes that increase risks for NS tumor development in childhood include classic neurocutaneous disorders like neurofibromatosis type 1 and type 2 (NF1, NF2) and tuberous sclerosis complex (TSC) type 1 and 2 (TSC1, TSC2). Li Fraumeni Syndrome, Constitutional Mismatch Repair Deficiency, Gorlin syndrome (Nevoid Basal Cell Carcinoma), Rhabdoid Tumor Predisposition syndrome, and Von Hippel-Lindau disease. Ataxia Telangiectasia will also be discussed given the profound neurological manifestations of this syndrome. In addition, there are other cancer predisposition syndromes like Cowden/PTEN Hamartoma Tumor Syndrome, DICER1 syndrome, among many others which also increase the risk of NS neoplasia and are briefly described. Herein, we discuss the NS tumor spectrum seen in the abovementioned cancer predisposition syndromes as with their respective germline genetic abnormalities and recommended surveillance guidelines when applicable. We conclude with a discussion of the importance and rationale for genetic counseling in these patients and their families.

Keywords Brain tumor · Cancer genetics · Cancer predisposition syndromes · Nervous system surveillance of genetic syndromes

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Introduction

Genetic syndromes which develop one or more nervous system (NS) tumors as one of the manifestations can be grouped under the umbrella term of NS tumor predisposition syndromes. Understanding the underlying pathological pathways at the molecular level has led us to many radical discoveries, in understanding the mechanisms of tumorigenesis, tumor progression, interactions with the tumor microenvironment, and development of targeted therapies. Currently, at least 7-10% of all pediatric cancers are now recognized to occur in the setting of genetic predisposition to cancer, or cancer predisposition syndromes [1, 2]. Specifically, the cancer predisposition rate in pediatric patients with NS tumors has been reported to be as high as 15%, though it can approach 50% in certain tumor types (i.e., choroid plexus carcinoma associated with Li Fraumeni Syndrome).

Cancer predisposition syndromes are caused by pathogenic variation in genes that primarily function as tumor suppressors and proto-oncogenes [3, 4]. These variants are found in the germline, or constitutional DNA. Mosaicism is defined as the presence in an individual of at least two cell lines differing in genotype and arising from a single zygote [5]. Mosaicism however, can affect only certain tissues, resulting in varied manifestations [6]. Increased understanding of the genetic underpinnings of cancer predisposition syndromes and the ability of clinical laboratories to offer molecular genetic testing allows for improvement in the identification of these patients. The identification of a cancer predisposition syndrome in a CNS tumor patient allows for changes to medical management to be made, including the initiation of cancer surveillance protocols. Finally, the identification of at-risk biologic relatives becomes feasible through cascade (genetic) testing [7, 8].

These fundamental discoveries have also broadened the horizon of novel therapeutic possibilities and have helped to be better predictors of prognosis and survival. The treatment paradigm of specific NS tumors may also vary based on the patient's cancer predisposition syndrome and may be used to guide therapy (example: immune checkpoint inhibitors in constitutional mismatch repair deficiency [CMMRD] predisposition syndrome and targeted therapies like MEK inhibitors, described later) [9]. Early diagnosis of these cancer predisposition syndromes along with early involvement of genetic counselors is therefore critical, in both unaffected and affected patients [10].

Cancer predisposition syndromes that increase risks for NS tumor development in childhood include classic neurocutaneous disorders like neurofibromatosis type 1 (NF1), schwannomatosis and its subtypes (e.g., NF2, SMARCB1

and LZTR1) and tuberous sclerosis complex (TSC) type 1 and 2 (TSC1, TSC2). Li Fraumeni Syndrome, Constitutional Mismatch Repair Deficiency, Gorlin syndrome (Nevoid Basal Cell Carcinoma), Rhabdoid Tumor predisposition syndrome, Von Hippel-Lindau disease, and Ataxia Telangiectasia will also be discussed given the profound neurological manifestations of these syndromes. In addition, there are other cancer predisposition syndromes like Cowden/PTEN Hamartoma Tumor Syndrome, DICER1 syndrome, among many others which also increase the risk of NS neoplasia and are briefly described.

Herein, we discuss the NS tumor spectrum seen in the abovementioned cancer predisposition syndromes as with their respective germline genetic abnormalities and recommended surveillance guidelines when applicable. We conclude with a discussion of the importance and rationale for genetic counseling in these patients and their families.

Neurofibromatosis Type 1 (NF 1)

NF1 is among the most common genetic disorders. The incidence is about one in every 2500–4000 live births making it one of the most common cancer predisposition syndromes. The incidence does not vary across gender or ethnicity [11, 12]. It is inherited in an autosomal dominant fashion, though 50% of cases are thought to occur *de novo* (are not inherited from an affected parent), with near-complete penetrance, though the clinical manifestations and phenotypes can vary tremendously. Individuals with NF1 are at risk of developing tumors in multiple organ systems, particularly the central and peripheral nervous systems.

Neurofibromin is a negative regulator of the RAS-MAP kinase and mammalian target of rapamycin (mTOR) signaling pathways, and it is a well-known proto-oncogene. The role of the MAPK pathway is well-established in cell growth and proliferation (Fig. 1). The neurofibromin protein is coded by the NF1 gene located on the long arm of chromosome 17. Germline mutations of NF1 result in a multitude of manifestations that can be seen throughout the body [13]. Common nervous system tumors that develop in NF1 include low-grade gliomas (LGGs), cutaneous neurofibromas, plexiform neurofibromas (PNs), malignant peripheral nerve sheath tumors (MPNSTs), and high-grade gliomas (HGGs). Other non-nervous system tumors seen in the NF1 population are juvenile myelomonocytic leukemia (JMML), embryonal rhabdomyosarcoma, gastrointestinal stromal tumors, pheochromocytoma, breast cancers and endocrine tumors among others. The diagnostic criteria for NF1 are well-established, and recommended surveillance guidelines have been published (Table 1).

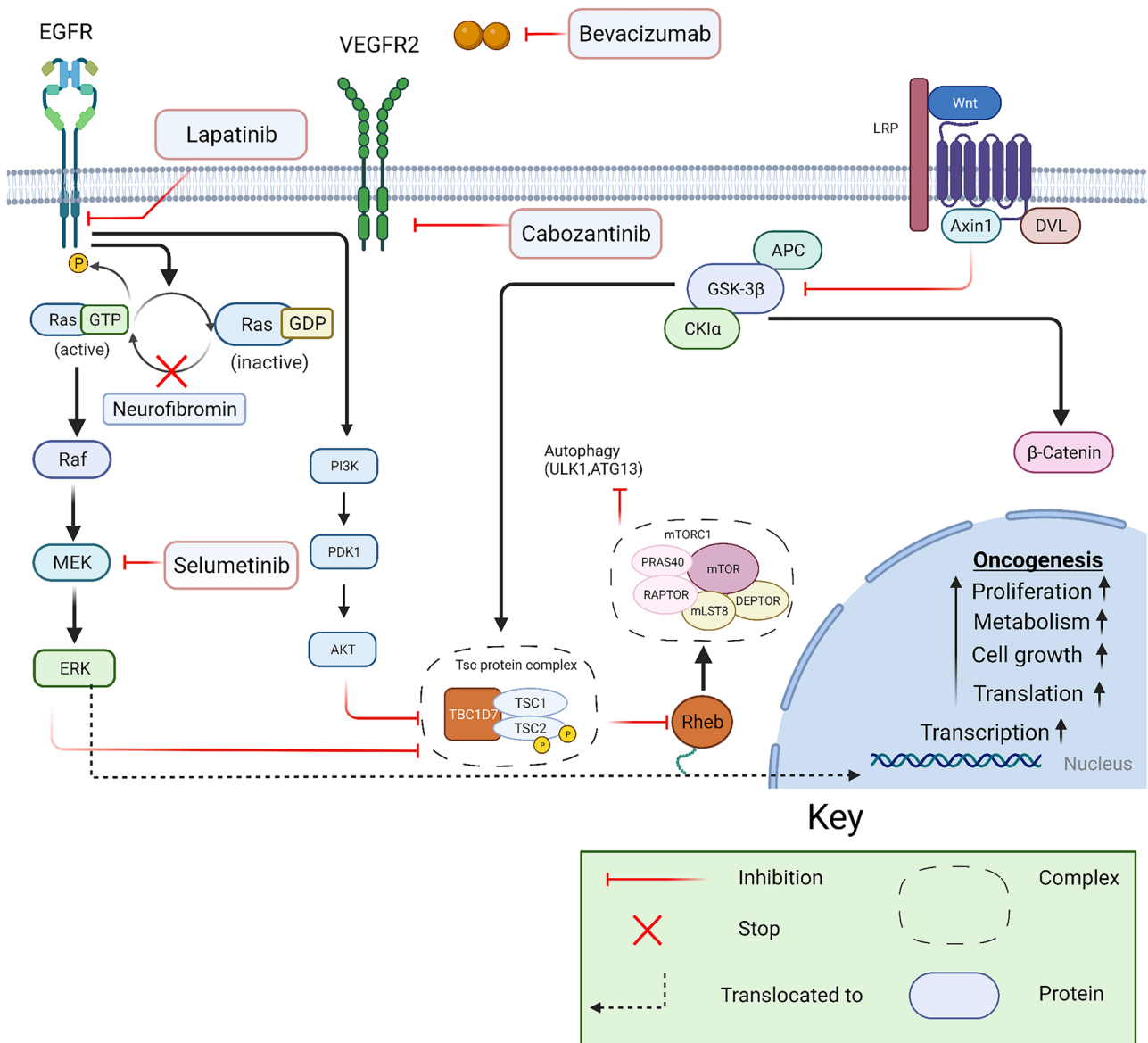


Fig. 1 Molecular pathways associated with pathogenesis of NF and TSC with point of inhibition

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Low-grade gliomas (LGG)

Optic pathway gliomas (OPGs) are the most common central nervous system tumor seen in NF1. OPGs are seen in up to 20% of the patients with NF1 and mainly present within the first 8 years of life, with a mean age of 4.5 years. On rare occasions, OPGs can present at an older age [14]. These lesions can often be indolent, but sometimes, they are progressive and may cause vision impairment. Once diagnosed with NF1, patients should undergo annual age-appropriate vision exams starting at the age of 1 year to 8 years; the recommendations for MRI screening after the age of 8

years are variable and considered on a case by case basis. This recommendation is based upon the relatively lower incidence of OPGs and vision dysfunction in this older age group [14, 15]. Tumors can arise anywhere from the retro-orbital optic nerve to optic radiations. In up to 50% of the patients, there may be no eye or vision symptoms associated with the tumors. Current recommendations are to treat only symptomatic tumors, particularly if there is a worsening in visual acuity. Due to the high degree of variability in symptoms and lack of reliable biomarkers, ophthalmologic exams and quantitative vision testing serve as the primary surveillance tools. Screening baseline MRI evaluations are not indicated for NF1-OPGs, as detecting these tumors rarely changes management in the absence of clinical symptoms

Table 1 Revised diagnostic criteria for neurofibromatosis type 1: an international consensus recommendation (NF1) [19]

Revised diagnostic criteria for neurofibromatosis type 1: an international consensus recommendation

A: The diagnostic criteria for NF1 are met in an individual who does not have a parent diagnosed with NF1 if two or more of the following are present:

- Six or more café-au-lait macules over 5 mm in greatest diameter in prepubertal individuals and over 15 mm in greatest diameter in postpubertal individuals^a
- Freckling in the axillary or inguinal region^a
- Two or more neurofibromas of any type or one plexiform neurofibroma
- Optic pathway glioma
- Two or more iris Lisch nodules identified by slit lamp examination or two or more choroidal abnormalities (CAs)—defined as bright, patchy nodules imaged by optical coherence tomography (OCT)/near-infrared reflectance (NIR) imaging
- A distinctive osseous lesion such as sphenoid dysplasia,^b anterolateral bowing of the tibia, or pseudarthrosis of a long bone
- A heterozygous pathogenic NF1 variant with a variant allele fraction of 50% in apparently normal tissue such as white blood cells

B: A child of a parent who meets the diagnostic criteria specified in A merits a diagnosis of NF1 if one or more of the criteria in A are present

^aIf only café-au-lait macules and freckling are present, the diagnosis is most likely NF1 but exceptionally the person might have another diagnosis such as Legius syndrome. At least one of the two pigmentary findings (café-au-lait macules or freckling) should be bilateral

^bSphenoid wing dysplasia is not a separate criterion in case of an ipsilateral orbital plexiform neurofibroma

or signs [16]. Asymptomatic tumors are hence monitored with annual to biannual ophthalmologic evaluations. Symptomatic tumors are often treated with chemotherapy and monitored with more frequent MRI surveillance and ophthalmologic exams [17, 18]. These practices may vary based on institution-specific protocols.

NF1-associated gliomas can occur in other parts of the central nervous system (CNS), including the brain, spinal cord, and cerebellum. Management options include maximum safe surgical resection symptomatic tumors. If surgery is not a safe option, chemotherapy is often utilized. Typically, it is the same chemotherapy used for LGG not associated with NF1, like the combination of carboplatin and vincristine, vinblastine monotherapy, and in the modern era, MAP kinase-targeted therapies, like Mek inhibitors as discussed in below. NF1-associated LGGs, including OPGs, have a more favorable prognosis when compared to the subgroup without NF1.

Surgery has a limited role in treating OPGs and is typically offered when vision is already significantly compromised and non-visual symptoms need management [16, 20]. Radiation is seldom used as a treatment modality in NF1-associated LGG due to the high risk of radiation-induced secondary high-grade malignancies, moyamoya disease and other radiation toxicities [21]. Chemotherapy is the main therapy option for OPGs and other NF-1-associated LGGs that require treatment. Packer et al. proposed chemotherapy involving vincristine and carboplatin in the early 1990s [22]. This was followed by the vinblastine regimen proposed by Buffet et al. [23] in the 2000s; both the regimens have similar effectiveness in stabilizing tumors and are often offered as the standard of care. Bevacizumab has been reported to improve visual outcomes in select cases; currently, studies

are underway to evaluate an objective response [24]. In many clinical trials, targeted therapies and combinations are being studied with recent advancements in understanding molecular pathways. The MAP kinase pathway can be inhibited at many target points. One such trial studying MEK1/2 inhibitor selumetinib showed 40% sustained partial response and 96% of patients and progression-free survival and was tolerated well with minor side effects; completion of the studies might replace the current first-line chemotherapy protocols with MEK inhibitors especially in OPGs associated with NF1 [25] (Figure 1).

Plexiform Neurofibromas (PNs) and Malignant Peripheral Nerve Sheath Tumors (MPNSTs)

Plexiform neurofibromas and malignant peripheral nerve sheath tumors arise from schwann cells, grow along the nerve, and are seen in up to 50% of patients with NF1. These tumors are rarely encountered outside the diagnosis of NF1. PNs can be present at birth and classically show maximum growth during childhood [26, 27]. While many PNs may be asymptomatic, some can present with significant complications and morbidities such as pain, cosmetic issues, and disfigurement which may compromise motor function. They can also cause pressure symptoms on the surrounding tissue, for example, around the carotid arteries or trachea which may lead to cardiovascular or respiratory compromise [27, 28–32]. Unlike LGGs, PNs are not sensitive to traditional cytotoxic chemotherapy regimens. Surgery is limited to debulking of symptomatic PNs. While it can be curative, complete resection is rarely safely achieved due to a high degree of involvement of the surrounding tissue and the underlying nerves. Various treatment options, including

antihistamines (ketotifen), antifibrotic (pirfenidone), and immunotherapies (thalidomide and interferon), have been studied with limited success [27, 33–38]. However, one recent phase 2 trial using pegylated interferon (PI) showed more than doubling of the time to progression compared with placebo, further studies are needed to understand the role of immunotherapy in treating PNs [39]. Among the targeted therapies, MEK inhibitors have shown the most promising results. Recently completed phase 2 trial, most children with neurofibromatosis type 1 and inoperable PNs had durable tumor shrinkage and meaningful clinical benefit from selumetinib [40, 41]. Selumetinib is currently FDA approved for children with NF1 with inoperable and symptomatic PNs, after the age of 2 years. Another such therapy with limited evidence but early data suggestive of benefits is cabozantinib, a broad receptor tyrosine kinase (RTK) inhibitor. RTK inhibition has been shown to reduce the interaction between schwann cells and the surrounding tumor microenvironment. In a recently published phase 2 trial evaluating the response of PNs to cabozantinib, of the 19 participants assessed for response, 8 (42%) had a partial response (PR), and 11 had stable disease (SD). No patient had disease progression while on the study. PR was defined as $\geq 20\%$ reduction in tumor volume from baseline on MRI [42]. Other trials targeting the mammalian target of rapamycin (mTOR) pathway, which has shown success in treating tumors related to neurofibromatosis 2 (NF2), have failed to show similar benefit in NF1 related PNs [43–45]. Future trials are underway to study the benefit of these medications in combination in addition to evaluating late effects. In approximately 8–13% of cases, PNs can progress to high-grade sarcoma-like tumors known as malignant peripheral nerve sheath tumors (MPNST) which is the leading cause of mortality in NF1 patients [46, 47]. Due to the risk of transformation of PNs to MPNSTs, radiation is not used to treat a classic PN [27, 48, 49] (Fig. 1).

Malignant Peripheral Nerve Sheath Tumors

Sequential inactivation of tumor suppressor genes is proposed as the underlying mechanism of MPNST. Other somatic alterations that contribute to high-grade cancerous behavior of these tumors in addition to biallelic inactivation of the NF1 gene include inactivation of tumor suppressors, namely CDKN2A/B, TP53, PTEN, and polycomb repressor complex 2 (PRC2). Additionally, the involvement of growth-promoting genes, such as PDGFR and EGFR, has also been implicated along with H3K27me3 alteration [27, 50, 51]. Risk factors for PNs transforming to MPNSTs include NF1 gene microdeletion, overall larger PN burden, atypical lesions, nodularity, and prior history of radiation [52]. Clinically, change in consistency or hardening of PNs, development of new neurological symptoms in the area involved,

rapid growth, distinct nodular lesions (DNLs), and refractory pain are considered red flags for transformation and warrant further investigation with MRI sequences including diffusion-weighted imaging. ^{18}F Fluorodeoxyglucose (FDG)-PET can also be considered, and FDG-PET scan can sometimes distinguish between benign growth of PNs versus malignant MPNST by demonstrating increased FDG uptake in the latter. Similarly, a combination of mean apparent diffusion coefficient (ADC) value and absence of split fat is excellent for discriminating malignant from benign PNSTs [12, 52–54]. Though tissue biopsy is still the gold standard for diagnosis, it might not always be safe to perform.

MPNSTs are aggressive and pose significant treatment challenges. Similar to PNs, there is a limited role of cytotoxic chemotherapy and radiation. Surgical options, including those using fluorescence techniques, are considered the first line to reduce tumor burden. Currently, many clinical trials are underway to study the role of targeted therapies alone and in combination. Many immune checkpoint inhibitors and vaccine therapies are also currently being tested, details of which are beyond the scope of the article.

High-Grade Gliomas (HGG)

NF1 patients are also at increased risk of developing HGG and most cases present in adulthood. The somatic molecular landscape of these HGGs is similar to that of MPNSTs, and similar additional secondary mutations are seen mainly in CDKN2A/B, TP53, ATRX, TERT, and PI3 kinase pathways along with the underlying germline NF1 mutation [15, 55]. It should be noted that alterations in IDH1 and histone proteins seen in adult HGGs are not typically seen in NF1-associated HGGs [56]. Overall, the prognosis of NF1-associated HGGs is better than in patients without NF1 [15, 57]. Due to its relative rarity, the treatment paradigm is yet to be universally standardized. Radiation along with temozolomide are often used, similarly to sporadic HGG, as they share similar tumor biology and harbor similar genetic mutations as mentioned above. Other novel therapies are still being tested in this population.

Neurofibromatosis Type 2 (NF 2)

NF2 was previously grouped under the umbrella term of neurofibromatosis and then considered a separate syndrome only after discovering the underlying genetic mutation in the NF2 gene on chromosome 22. NF2 is an autosomal dominant disease with distinct clinical manifestations and therapeutic challenges as compared to NF1.

The constellation of these manifestations along with the family history is used in the Manchester criteria, which remains the standard for diagnosis (Table 2). Due to significant mosaicism, genetic testing might be negative in up to

Table 2 Manchester criteria for diagnosis of NF2, diagnosis of NF2 requires to meet primary and added features as mentioned below

Primary finding	Added features needed for diagnosis
Bilateral Vestibular Schwannoma	None
First-degree relative with NF2	Unilateral Vestibular Schwannoma, or Any two (2) other NF2-Associated lesions: Meningioma, Schwannoma, Ependymoma, or Juvenile Cataracts
Unilateral Vestibular Schwannoma	Any two (2) other NF2-Associated lesions: Meningioma, Schwannoma, Ependymoma, or Juvenile Cataract
Multiple Meningiomas	Unilateral Vestibular Schwannoma, or Any two (2) other NF2-Associated lesions: Schwannoma, Ependymoma, or Juvenile Cataracts

15% of the patients even if they meet the clinical criteria, in such cases testing for NF2 mutation in tumor tissue and germline might have a better yield. Genetic testing is often not required for the diagnosis. NF2 affects 1 in 25000 people worldwide [58, 59]. Common nervous system tumors in NF2 involve the cells covering the cranial nerves (schwannomas), brain (meningiomas), and ventricles and spinal canal (ependymomas). Most of the tumors arising in NF2 are low-grade, and malignant transformation is rare. However, patients may suffer from significant morbidity due to pressure symptoms on surrounding cranial nerves (CNs) and brainstem and spine, creating chronic progressive symptoms which are often challenging to treat and ultimately lead to death [60, 61].

The relationship between the severity of the symptoms and underlying germline NF2 variant is becoming more evident. It is now known that biallelic loss of NF2 gene is necessary for tumor development in schwannomas. Patients with truncated mutations present at an earlier age and have more severe symptoms when compared to missense mutations which have a milder course [62].

Schwannomas

As the name suggests, schwannomas arise from the schwann cells which are responsible for the production of myelin and play a similar role to the oligodendrocyte in the central nervous system [63]. NF2-associated schwannomas most commonly arise within the 8th cranial nerve (CN), and multiple foci along the same nerve can be involved. Symptoms are mainly due to the pressure effects on the surrounding cranial nerves and brainstem. For example, schwannoma of 8th CN often presents clinically with hearing loss due to direct effects upon the cochlear part of the CN 8. This can then progress to vestibular symptoms and facial weakness as the vestibular part of the 8th CN and facial nerve lie in close anatomic proximity. Current data suggests that the size of the tumor does not directly correlate to the severity of hearing loss [64, 65].

Since the schwannoma typically does not lead to rapid mortality, the management is mainly on preserving the function of the nerve and patient functional outcomes such as

hearing whenever possible. This limits the role of surgery and radiation in managing the disease. Currently, bevacizumab a VEGF inhibitor (Figure 1) is considered the first choice in treating symptomatic CN8 schwannomas, as multiple studies have shown benefit in tumor size and hearing [66–68]. Bevacizumab has shown to cause tumor volume reduction in 40% of the tumors and 40% improvement in hearing loss. While bevacizumab may be used to control size for either unilateral (hearing ear) or bilateral tumors, when used for hearing loss alone, it is generally withheld until an only functioning/hearing ear begins to fail. Another phase 2 study using lapatinib, an EGFR and Erb2 inhibitor, has also demonstrated some improvements in both tumor volumes and hearing and may be considered in the management [69].

With recent advancements in the understanding of genetic and molecular underpinnings, several clinical trials are underway investigating different therapeutic options. Some of them include the COX2 inhibitor aspirin (NCT03079999), MEK inhibitor selumetinib (NCT03095248), tyrosine kinase inhibitor brigatinib (NCT04374305), and crizotinib, which inhibits focal adhesion kinase 1 (FAK1) (NCT04283669).

Meningiomas

Meningiomas are the second most common tumor typical of NF2. These can develop anywhere in the craniospinal axis through the length of meninges; however, they typically occur intracranially in the majority of the cases [70]. Upto 80% of patients with NF2 develop meningiomas and multifocal disease is seen in almost all of these patients. The presence of multiple meningiomas or early onset meningiomas should raise suspicion of NF2, as this might be the first presenting sign [71–73]. Treatment options for meningiomas are limited. Surgical resection is typically reserved for symptomatic patients. Radiation therapy is used on a case-by-case basis when complete surgical resection is not possible, radiation is used with caution in the setting of NF2 meningiomas. lapatinib, everolimus, and combination everolimus + octreotide and bevacizumab + everolimus have shown benefit in slowing the tumor growth, none have shown any benefit in tumor shrinkage [74–78].

Meningioangiomas

Meningioangiomas was initially considered a benign hamartomatous lesion with meningeal and vascular components; however, it is increasingly being recognized as a pre-cancerous lesion. It can be unifocal or multifocal in nature and clinicians are advised to follow with serial MRIs [79–82].

Ependymomas

Ependymomas are seen in about 35–50% of the patients with NF2, but only 20% of the patients report symptoms. These most commonly arise in the spinal cord and cervical medullary junction [83, 84]. Ependymomas are more often seen in patients with truncating *NF2* mutations, highlighting that those with truncating mutations have a more aggressive disease course [72]. Surgery is considered the standard of care with a goal of gross total resection when feasible and safe, especially in symptomatic patients. Radiation therapy (both newer proton beam and traditional photon therapy) is well established in the treatment of sporadic ependymomas and is considered with caution considering the underlying NF2 mutation [27, 85, 86].

Schwannomatosis

Schwannomas can also arise in other conditions which have overlapping clinical manifestations with NF2. These conditions are known as schwannomatosis and can be further classified as SMARCB1 (previously INI) and LZTR1-related schwannomatosis based on underlying germline mutation. Schwannomatosis is yet to be well understood. It is an autosomal dominant disorder with incomplete penetrance. About 5% of patients with germline pathogenic SMARCB1 variants might develop meningiomas. Germline pathogenic LZTR1 mutations can present with only unilateral vestibular schwannomas along with other intradermal tumors like NF2 but often lack other ocular manifestations like posterior subcapsular cataracts and retinal findings and skin findings [27, 87, 88]. It is important to make the distinction between these three overlapping conditions as the course of the tumors, management, and surveillance guidelines may change based on the diagnosis. Hence, the revised Manchester criteria differentiate LZTR1 as a separate condition from NF2 [87, 89] (Table 2).

Constitutional Mismatch Repair Deficiency (CMMRD)

CMMRD is one of the most aggressive cancer predisposition syndromes, with typical onset of malignancy in childhood. This rare autosomal recessive syndrome is caused by

bi-allelic pathogenic variants in one of the mismatch repair genes: MLH1, MSH2, MSH6, and PMS2, and rarely EPCAM or MSH3. Single variants in these genes cause autosomal dominant Lynch Syndrome (LS), or Non-polyposis Hereditary Colorectal Cancer Syndrome. LS involves a predisposition to the development of gastrointestinal and genitourinary tract cancers with a mean age of first tumor onset of 45 years old [90–92]. While patients with Lynch syndrome can develop CNS tumors, it is rare. This syndrome is considered an adult-onset cancer predisposition syndrome. Therefore, current LS surveillance guidelines do not recommend screening for CNS tumors or any cancer screening in childhood [93, 94].

The process of DNA replication is a highly conserved process, which relies on DNA polymerases and the mismatch repair (MMR) system. Specifically, the exonuclease domains of DNA polymerases along with the mismatch repair complex are responsible for monitoring and repairing DNA replication errors. As a result, the hallmark of mismatch repair deficiency is an accumulation of point mutations and microsatellite instability [95].

Cancers in constitutional mismatch repair deficiency can arise as young as infancy and throughout childhood. The mean age for the first tumor presentation is reported to be 7.5 years old, though there is a wide range reported (0.4–39) [94]. Brain tumors are the most common tumors reported in patients with CMMRD and present at an average age of 10 years. Although a variety of brain tumors have been reported, including medulloblastomas, high-grade gliomas (HGG) are the most common type seen [93, 94, 96, 97]. Non-CNS malignancies reported in CMMRD include T cell lymphoblastic lymphoma, colorectal carcinoma, and other blood and gastrointestinal malignancies along with some of the neurocutaneous features described below.

Surveillance guidelines for patients with CMMRD have been proposed by both the “Care 4 CMMRD,” an International Biallelic Mismatch Repair Deficiency (BMMRD) Consortium and by the American Association of Cancer Research (AACR) Childhood Cancer Predisposition Workshop. They include brain MRI imaging at diagnosis and every 6 months thereafter. In addition, whole body MRI (WB-MRI) should begin by age 6 y/o (when the need for anesthesia is lessened) and be performed annually. Since body MRIs are not sensitive enough for brain tumors, brain MRIs are still recommended bi-annually [93, 94, 98]. Additional surveillance recommendations are included in Table 3.

Interestingly, there is an overlap between some neurocutaneous findings between CMMRD and NF1; patients with both diseases can develop café au-lait macules (CALM). While 99% of patients with NF1 develop CALMs by the age of 1 year, CALMs are known to develop in 62–71% of the patients with CMMRD [93, 96, 97]. However, many CMMRD patients do not have greater than 5 CALMs

Table 3 Summary of cancer predisposition syndromes and recommended surveillance in childhood

Cancer predisposition syndrome (inheritance)	Genes involved	CNS tumor risk/spectrum	Other neoplasia risk and features	AACR Childhood Cancer Predisposition Workshop: Surveillance Guidelines (for those without active disease)
Neurofibromatosis type 1 (AD)	NF1	Optic pathway gliomas, Gliomas (low and high grade)	Malignant peripheral nerve sheath tumor Leukemia Lymphoma Embryonal rhabdomyosarcoma Breast cancer Pheochromocytoma Gastrointestinal stromal tumor	Annual neurologic exam and physical exam Ophthalmic assessments from birth to 8 years, every 6 months to 12 months MRI surveillance is not currently recommended unless symptomatic or with an already diagnosed tumor Annual Screening for Breast cancer and Hypertension after the age of 30 years Annual history and physical exam (including audiology with measurement of pure-tone thresholds and Word Recognition Scores) Annual (consider twice yearly in first year since diagnosis or signs of rapid growth) brain MRI starting at 10 years of age. If baseline imaging shows no characteristic sites of involvement, reduce frequency of screening to every 2 years Surveillance spinal MRI is recommended at 24– to 36-month intervals beginning at 10 years of age
Li Fraumeni Syndrome (AD)	TP53	High-grade glioma, anaplastic astrocytoma, choroid plexus carcinoma medulloblastoma	Sarcomas Breast Cancer Adrenocortical Carcinoma Leukemia and Lymphoma	Complete physical examination and neurologic exam every 3–4 months US of abdomen and pelvis every 3–4 months Annual brain MRI beginning at diagnosis Annual Whole-Body MRI beginning at diagnosis
Von Hippel Lindau Syndrome (AD)	VHL	CNS Hemangioblastoma	Pheochromocytomas, retinal hemangioblastomas, ELST	Biennial brain and spine MRI beginning at age 8 Annual ophthalmologic examination beginning at birth Annual physical exam Annual plasma free metanephrines beginning at age 2 Biennial audiogram beginning at age 5 Annual abdominal MRI beginning at age 10

Table 3 (continued)

Cancer predisposition syndrome (inheritance)	Genes involved	CNS tumor risk/spectrum	Other neoplasia risk and features	AACR Childhood Cancer Predisposition Workshop: Surveillance Guidelines (for those without active disease)
Constitutional Mismatch Repair Deficiency (CMMRD) (AR)	MLH1, MSH2, MSH6, PMS2, EPCAM	High-grade glioma, Medulloblastoma	Hematological malignancies, Gastrointestinal malignancies	Brain MRI at diagnosis and once every 6 months Whole-body MRI annually beginning at age 6 Abdominal ultrasound q6 months beginning at age 1 CBC q 6 months beginning at age 1 Upper gastrointestinal endoscopy; VCE, ileocolonoscopy annually beginning at age 4–6
Gorlin Syndrome / Nevroid Basal Cell Carcinoma Syndrome (AD)	PTCH1, SUFU	Medulloblastoma, especially desmoplastic and SHH-type	Basal cell carcinoma, cardiac and ovarian fibromas, keratocystic odontogenic tumors	Brain MRI can be considered in those with a SUFU variant every 4 months until age 3 and then every 6 months until age 5 Basic echocardiogram in infancy Dental exams with jaw X-ray every 12 to 18 months beginning at age 8 for PTCH1 mutation carriers Ovarian ultrasound by age 18 Basal cell carcinoma screening annually to begin by age 10
Rhabdoid Tumor Predisposition Syndrome (AD)	SMARCB1, SMARCA4	Atypical teratoid rhabdoid tumors, Schwannomatosis, Meningioma	Rhabdoid tumors, Non-small cell carcinoma of the ovary hypercalcemic type	Brain MRI every 3 months till age 5 Consider whole-body MRI annually till age 5 Abdominal ultrasound every 3 months to age 5 (Recommendations are only made for SMARCB1 loss of function mutations)

AD autosomal dominant, AR autosomal recessive, ELST endolymphatic sac tumor

with the required measurements and age to meet the criteria for NF1 (Table 1). Furthermore, there appears to be a subtle difference between the morphologic appearance of CALMs. Patients with CMMRD develop a jagged “coast of Maine appearance” due to varying degrees of pigmentation and appear different from the more uniformly pigmented and smooth margined CALMs seen in NF 1 patients [93, 99–102]. Axillary freckling, Lisch nodules, cutaneous neurofibromas, and tibial pseudoarthrosis, all of which are features of NF1, are also reported in CMMRD with varying frequency. Hence, if the patient shows phenotypic manifestations of NF1 but tests negative for NF1 and SPRED1 gene alteration (causing legius syndrome), CMMRD should be considered in the differential diagnosis [93, 96, 97, 103–106].

Tuberous Sclerosis Complex (TSC)

TSC is another neurocutaneous syndrome that involves the development of a unique set of symptoms involving multiple organ systems. TSC is the result of suppression of one of the tumor suppressor genes namely TSC1 located on chromosome 9 and TSC 2 on chromosome 16. TSC1 and TSC2 encode for proteins hamartin and tuberlin respectively, both of which are involved in the suppression of the mTOR pathway via RAS homologue enriched in the brain (Rheb). These are inherited in an autosomal dominant fashion with near-complete penetrance, but 2/3 of cases are thought to be from de novo mutations [107–110]. The spectrum of manifestations includes typical skin findings, such as hypomelanotic macules, ungual fibromas, shagreen patch, and angiofibromas as well as angiofibromas lymphangiomyomatosis, and hamartomatous growths in multiple organ systems.

In the brain, hamartomatous growth, focal cortical dysplasia, and other developmental anomalies can present even before birth. Epilepsy develops in 70–80% of patients with TSC and is often refractory to traditional anticonvulsants. Frequently, patients develop epileptic encephalopathy in the form of infantile spasms. Current treatment guidelines recommend using ACTH and vigabatrin as the first-line therapy. Recently, EPISTOP demonstrated that when used prophylactically in patients with TSC (without epilepsy), vigabatrin reduces the risk and severity of epilepsy [111].

Another unique finding in TSC is the clustering of cells lining the ventricles named subependymal nodules (SEN) which occur in 80–90% of the patients. Approximately 15% of subependymal nodules transform into a

more aggressive subependymal giant cell astrocytoma (SEGA) [112–114]. Other features with diagnostic criteria are listed in Table 4 (Washington DC 2012 TSC meeting report). A better understanding of the mTOR pathway and its role in TS led to clinical trials testing mTOR inhibitors, ultimately establishing everolimus as an FDA approved drug in the management of TSC-associated SEGAs as discussed below.

SEGA

Subependymal nodules and low-grade hamartomas are mostly asymptomatic. However, depending on the location within the ventricles, these might cause blockage of CSF flow resulting in obstructive hydrocephalus. While SENs mostly develop in the first 2 decades of life, some congenital cases have been reported. When diagnosed congenitally, they are known to grow at a faster rate when compared to non-congenital SEGAs [115–123]. In one large multicenter study of 2200 patients with TSC, it was found that SEGAs developed in 25% of the patients. There is a higher rate in patients with a TSC2 (33%) compared to patients with a TSC1(13%). SEGAs were symptomatic in 42% of patients, and the most common presenting symptoms included the following: seizures or increased seizure frequency (15%), regression of milestones (10%) or behavioral change (12%) [124].

SEGAs are typically classified as low-grade gliomas and mostly have an indolent course. Currently, they are defined by imaging criteria set in 2013 by an international panel of experts [125]. Once a diagnosis of TSC is made, a baseline MRI is recommended to evaluate for underlying SENs and SEGAs [126]. SEGAs show contrast enhancement, and classically, continued tumor growth on serial surveillance MRIs. These characteristics help differentiate them from SENs. Like many SENs, SEGAs can also be asymptomatic. Current recommendations are to treat symptomatic SEGAs and to monitor asymptotic SEGAs and SNs with serial surveillance MRIs.

Everolimus is FDA approved for the treatment of SEGAs, and in the modern era, surgery is reserved for select patients, typically with acute symptoms such as hydrocephalus [110, 127–130]. Everolimus has shown to be effective in shrinking SEGAs by 30–50% and has also shown benefit in the control of underlying epilepsy which can often be refractory to classic anticonvulsants. Similar effects are also seen on renal angiomyolipomas [129, 131–133]. The role of mTOR in other manifestations of TSC including TSC-associated neuropsychiatric disorder (TAND) is yet to be well understood.

Table 4 Tuberous sclerosis complex clinical features and diagnosis

Diagnostic criteria for tuberous sclerosis complex 2012

Diagnostic criteria for tuberous sclerosis complex 2012			
A. Genetic diagnostic criteria	The identification of either a TSC1 or a TSC2 pathogenic mutation in DNA from normal tissue is sufficient enough to make a definite diagnosis of Tuberous Sclerosis Complex (TSC)		
B. Clinical diagnostic criteria	Major features	<ol style="list-style-type: none"> 1. Hypomelanotic macules (≥ 3, at least 5-mm diameter) 2. Angiofibromas (≥ 3) or fibrous cephalic plaque 3. Ungual fibromas (≥ 2) 4. Shagreen patch 5. Multiple retinal hamartomas 6. Cortical dysplasias^a 7. Subependymal nodules 8. Subependymal giant cell astrocytoma 9. Cardiac rhabdomyoma 10. Lymphangiomyomatosis (LAM)^b 11. Angiomyolipomas (≥ 2)^b 	Definite diagnosis: Two major features or one major feature with ≥ 2 minor features Possible diagnosis: Either one major feature or ≥ 2 minor features
	Minor features	<ol style="list-style-type: none"> 1. “Confetti” skin lesions 2. Dental enamel pits (> 3) 3. Intraoral fibromas (≥ 2) 4. Retinal achromic patch 5. Multiple renal cysts 6. Nonrenal hamartoma 	

^aIncludes tubers and cerebral white matter radial migration lines

^bA combination of the two major clinical features (LAM and angiomyolipomas) without other features does not meet criteria for a definite diagnosis. A combination of these two major clinical features does not meet criteria for a definite diagnosis

Von Hippel–Lindau Syndrome (VHL)

VHL is caused by pathogenic germline variants in the VHL gene, located on chromosome 3. This encodes for the pVHL protein, which regulates the response of a cell to hypoxia via hypoxia-inducible transcription factor (HIF) 1 and 2. The loss of a pVHL allele results in a skewed ratio of HIF 1 to HIF 2 resulting in an imbalance of the VHL/HIF axis and tumorigenesis [134]. VHL is inherited in an autosomal dominant fashion with an incidence of around 1 in 36000 live births, but up to 25% of cases occur due to a de novo mutation, but up to 25% of cases occur due to a de novo mutation [135–142].

VHL is not associated with malignant tumors of nervous system origin; however, hemangioblastomas are seen in 60–80% of patients with VHL and most often develop along the craniospinal axis, often causing neurologic symptoms due to direct pressure effects on adjacent structures or spontaneous hemorrhages [143, 144]. Patients with VHL are also known to develop endolymphatic sac tumors in the temporal bone. About 15% can also present

with hearing loss, vertigo, tinnitus, and headaches [145, 146]. Hemangioblastomas can remain dormant for long periods of time before becoming symptomatic due to sudden growth or hemorrhage [147].

Other manifestations of VHL include neuroendocrine tumors of the pancreases, renal angiomas, renal cell carcinomas, serous cystadenomas, paragangliomas, and pheochromocytoma (PPGLs). VHL can be further classified as type 1 (truncating or missense mutation developing pheochromocytoma) and type 2 (missense mutation usually not developing pheochromocytoma). A better understanding of the VHL axis may lead to unique surveillance guidelines in the future based upon type (1 or 2) [136]. It should be noted that patients with VHL show variable degrees of mosaicism impacting clinical management and counseling [148].

Pheochromocytomas and paragangliomas (PPGLs) arising from adrenal chromaffin cells can also be seen in other mutations along the VHL/HIF axis, including PHD, VHL, HIF-2A (EPAS1), and SDHX (VHL Figure 2). About 40% of PPGLs result from germline mutations [149].

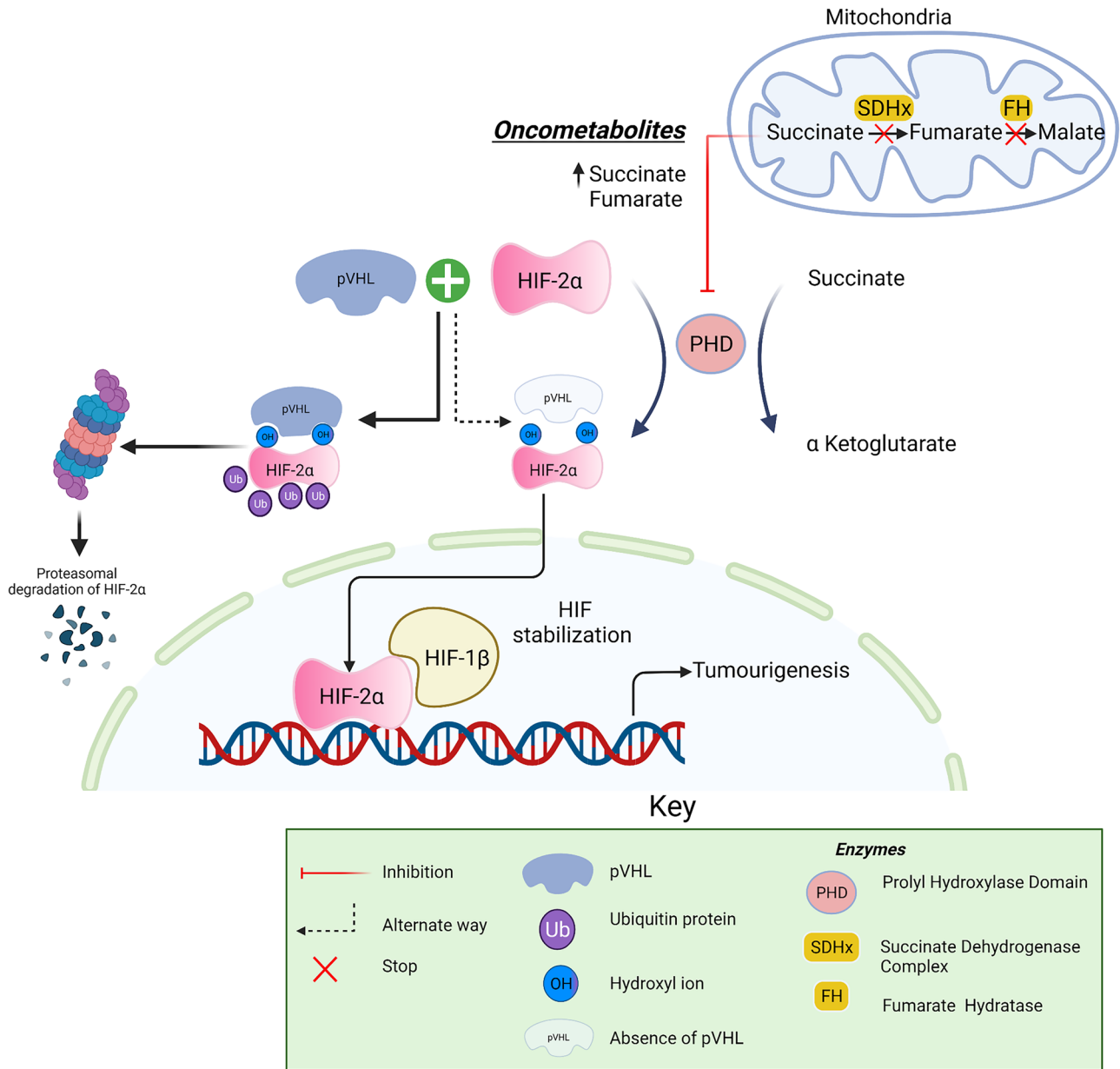


Fig. 2 Pathogenesis of Von Hippel-Lindau syndrome in VHL/HIF axis

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Current NIH and VHL Alliance recommendations for surveillance in individuals with VHL are to obtain MRI imaging of the brain and spinal cord with contrast every two years from the age of 11 years old onward [150]. If neurological symptoms are seen, then imaging should be considered sooner. Additionally, MRI without contrast is recommended around the 4th month of pregnancy to evaluate for the need

for cesarean section in the presence of retinal, brain, or spinal lesions within the developing fetus [136, 140, 147, 151, 152]. This is accompanied by detailed neurological testing, ophthalmologic and hearing tests as mentioned in Table 3.

Recently, Belzutifan a hypoxia-inducible factor inhibitor was approved to treat adults with VHL-associated renal cell carcinoma, central nervous system hemangioblastomas, and pancreatic neuroendocrine tumors that do not require immediate surgery [153].

Genetic counseling should be a part of management and should be offered early. As 80% of the patients have an affected parent and 20% of patients have de novo mutations, all affected individuals have a 50% chance of passing down VHL to their biological children.

Ataxia Telangiectasia (AT)

As the name suggests, this syndrome is primarily characterized by ataxia secondary to progressive cerebellar atrophy and development of telangiectasias. AT is an autosomal recessive disorder with an incidence of 1 in 20,000 to 100,000 people [154]. Mutations in the “ataxia telangiectasia mutated” (ATM) gene located on chromosome 11 are known to cause carcinogenesis and can promote malignant transformation. Gene products of ATM genes are normally involved in repairing double-strand DNA breaks. Although reported, primary CNS tumors are not a cardinal manifestation. Other characteristic cancers of AT are lymphomas, leukemias, breast, and gastric cancers. Patients are particularly radiosensitive and ionizing radiation should be avoided [147, 155, 156].

Surveillance guidelines for individuals with AT have been recommended by the Pediatric Cancer Predisposition Workshop [157]. Routine MRI surveillance in patients with brain and spinal lesions should be considered. If neurological symptoms arise, surgery can be considered with caution. Although no targeted therapies have been successful in treating the underlying disorder or manifestations, beta-methasone has shown some improvement in ataxia and can be tried if ambulation is affected [158]. Patients with AT can have poor ambulation and often develop early wheelchair dependency. Only 50% of the patients will survive into their 20s [159, 160].

Nevoid Basal Cell Carcinoma Syndrome (NBCCS or Gorlin Syndrome)

NBCCS or Gorlin syndrome has an incidence of 1:15,000 to 19,000 births and is caused by an underlying germline mutation in the sonic hedgehog (SHH) pathway, specifically in patched1 (PTCH1) gene and suppressor of fused (SUFU) gene, which are inherited in an autosomal dominant fashion [161–164]. About 5% of patients develop medulloblastomas in the course of the disease, and most of them arise before the age of 3 years old [165–167]. Often medulloblastoma might be the first presenting sign, and therefore, a high level of suspicion is needed. Detailed family history and a detailed skin exam should be performed in those children less than 3 years old presenting with medulloblastoma, particularly if the subgrouping is SHH. Other features that help in the clinical diagnosis include jaw keratocyst, palmar and plantar pits, multiple basal cell carcinomas, and MRI findings of lamellar calcification of falx or clear evidence of calcification in a young child. The Jones criteria are often used to make the diagnosis and is described in Table 5 [168]. Patients who receive radiation for treatment of their medulloblastoma have the risk of developing a large number of basal cell carcinomas within the radiation entrance fields, further stressing the importance of early diagnosis [169, 170].

Rhabdoid Tumor Predisposition Syndrome (RTPS)

RTPS results from germline loss-of-function mutations in the SMARCB1 gene and differs phenotypically from schwannomatosis though both the syndromes

Table 5 Jones criteria for diagnosis of Gorlin syndrome. A diagnosis can be made when 2 major or 1 major and 2 minor criteria are fulfilled

Major criteria	Minor criteria
Lamellar (sheet-like) calcification of the falx or clear evidence of calcification in an individual younger than age of 20 years	Childhood medulloblastoma
Jaw keratocyst	Lympho-mesenteric or pleural cysts
2 or more palmar/plantar pits	Macrocephaly (OFC > 97th centile)
Multiple BCCs (more than five in a lifetime) or a BCC before the age of 30 years	Cleft lip/palate
First degree relative with Gorlin Syndrome	Vertebral/rib anomalies such as bifid/splayed/extra ribs or bifid vertebrae
	Preaxial or postaxial polydactyly
	Ovarian/cardiac fibromas
	Ocular anomalies (cataract, developmental defects, and pigmentary changes of the retinal epithelium)

share common genetic mutation. More recently, germline SMARCA4 mutations have also been associated with RTPS. These are both transmitted in an autosomal dominant fashion. Patients with germline mutations in SMARCB1 are classified as RTPS type 1 and patients with germline mutations in SMARCA4 are categorized as RTPS type 2 [171–174]. Patients with SMARCB1 mutations have shown more aggressive tumor types like atypical teratoid rhabdoid tumors (AT/RT) and intracranial meningiomas, schwannomas, and malignant peripheral nerve sheath tumors in the peripheral nervous system. Whereas, patients with SMARCA4 pathogenic variants harbor more risk of ovarian and thoracic carcinoma, in addition to the rhabdoid tumors. AT/RT can be the first manifestation, and hence, the current recommendation is to test for germline mutations in both the genes in all newly diagnosed cases of AT/RT [175–177]. Underlying germline mutation in SMARCB1 also indicates an overall poor prognosis.

Initial diagnosis of RTPS can be established after genetic confirmation of germline heterozygous mutation in either of the two genes in a patient with rhabdoid tumor anywhere in the body, and/ or a family history of rhabdoid tumors and/ or other typical tumors as described above [178].

Surveillance recommendations have been published by the AACR Pediatric Cancer Predisposition Workshop and suggest MRI brain and ultrasound of the abdomen every three months from the day of diagnosis till the age 5 years and whole-body MRI at the age 5 years [171]. This recommendation was made only for truncating/loss of function mutations in the SMARCB1 gene. No recommendations were made on brain imaging of SMARCA4 genes and germline missense mutations of SMARCB1. The only recommendation for SMARCA4 was abdominal ultrasound every 6 months and consideration of oophorectomy if considered high risk after childhood [171].

Li-Fraumeni Syndrome (LFS)

LFS is an autosomal dominant disorder resulting from germline pathogenic variants in the TP53 gene. The TP53 tumor suppressor gene is located on chromosome 17. In addition to germline variants causing LFS, somatic (tumor) mutations in TP53 are some of the most common genetic alterations in human cancer [179].

Diagnosis of Li-Fraumeni can be made in a patient with a heterozygous germline mutation in TP53 and/or a patient meeting all the 3 diagnostic criteria, including the following: development of sarcoma before 45 years; one first-degree relative with any cancer diagnosed before 45 years; and one first or a second-degree relative with any

cancer diagnosed before age of 45 years *or* a sarcoma at any age [180].

The risk of CNS tumors in LFS is well-established, and numerous brain tumors have been described such as medulloblastomas, astrocytomas, and choroid plexus carcinomas. The incidence of brain tumors in LFS can be estimated to be around 12% and appear to have a bimodal distribution with the first peak occurring around infancy to early childhood and a second, smaller peak at about the 3rd to 4th decade of life [181–183]. Females have a higher risk of tumors including brain tumors when compared to males [184, 185]. The implications of germline TP53 abnormalities on medical decision-making and prognosis remain poorly understood; however, all individuals with LFS are highly susceptible to the DNA-damaging effect of radiation, increasing the risk for secondary malignancies within radiation therapy fields [186].

Currently, the modified Toronto protocol of 2017 recommends annual brain MRIs; the first one should be performed with contrast and the subsequent ones without contrast if no abnormality is observed [187].

Surveillance guidelines for pediatric patients with LFS are available through the AACR Pediatric Cancer Predisposition Workshop and summarized in Table 3 [187].

Other Cancer Predispositions with CNS Tumors

Other rare cancer predisposition syndromes with a risk of developing CNS tumors are described below.

Multiple Endocrine Neoplasia Type 1 (MEN-1)

This autosomal dominant disorder of the tumor suppressor gene MEN1 can present with pituitary adenoma or tumors of the pancreas and parathyroid glands. The pituitary adenomas are likely to be prolactinomas and are more resistant to treatment when compared to the ones seen in the general population [188–190]. Neurologic symptoms mainly arise due to direct local pressure on the pituitary often causing vision loss and disturbances in the hypothalamic-pituitary axis.

Rubinstein Taybi Syndrome (RTS)

RTS is an autosomal dominant disorder linked to CREBBP gene products which plays a regulatory role in SHH and AMP regulated gene expression. An increased risk of medulloblastoma, oligodendroglioma, and meningiomas has been reported. Other morphological features that help raise suspicion of this diagnosis include broad

thumbs, broad great toes, and distinctive facial features [191–194].

Cowden Syndrome (CS)

CS is an autosomal dominant syndrome with an underlying mutation in *PTEN* which has been linked to a higher risk of brain tumors, specifically cerebellar dysplastic gangliocytoma [195–197]. *PTEN* mutations have also been described in various other types of cancers.

Melanoma-Astrocytoma Syndrome

First described by Kaufman et al in 1993, melanoma-astrocytoma syndrome is now known to have mutations in *CDKN2A*. Many families have been described presenting with both melanomas and astrocytomas. Genetic testing should be considered if skin findings are concerning [198–204].

DICER1 Syndrome

DICER1 Syndrome, previously called Pleuropulmonary Blastoma (PPB) Syndrome, is caused by germline pathogenic variants in the *DICER1* gene. This autosomal dominant syndrome increases the risk for a broad spectrum of both benign and malignant neoplasia, including pleuropulmonary blastoma, cystic nephroma, genitourinary embryonal rhabdomyosarcoma, ovarian Sertoli Leydig cell tumors, and thyroid carcinoma. Specifically, CNS manifestations can include metastatic PPB, pituitary blastoma, pineoblastoma, and ciliary body medulloepithelioma [205].

Genetic Counseling

All individuals suspected to have a hereditary cancer predisposition syndrome should be referred to a cancer genetic counselor or a pediatric genetics clinic. Genetic counselors are trained to provide pre-test counseling, obtain informed consent for genetic testing, and select the optimal genetic test for each patient. Once genetic testing results become available, genetic counselors are trained to facilitate a discussion of recommended cancer surveillance for children and their families. Genetic counselors are also trained to provide psychosocial support to patients undergoing an evaluation for a cancer predisposition syndrome, as well as to coping and adapting to living with the information once results become available.

Conclusion

Cancer predisposition syndromes might present to neurologists and oncologists alike in pediatric population or later as adults. Early genetic counseling is imperative and should

be part of routine medical care. A detailed clinical history and examination will help raise suspicion, further confirmed with appropriate genetic testing. Patients and families can thus be appropriately subjected to screening measures, aiding not only in potential early diagnosis and management altering the course of the disease but also in family planning. As we continue to understand the complex interplay of genetic and molecular pathways, more diagnostic and therapeutic possibilities are being uncovered and surveillance guidelines continue to be updated in line with newer research.

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Required Author Form Disclosure form provided by the author are available with the online version of this article.

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