



The majority of pediatric cancers are sporadic; however, about 5% are thought to be due to a hereditary predisposition.<sup>1</sup> The disorders represented on this panel are typically *de novo* or inherited in an autosomal dominant fashion, with the exception of mosaic variegated aneuploidy syndrome (autosomal recessive inheritance).<sup>2</sup>

Individuals who carry a pathogenic variant in a hereditary cancer gene have an increased risk of certain cancers compared to the general population. Cancer risks depend on the gene(s) in which the variant(s) is identified. These individuals are eligible for increased cancer screening and/or risk reducing surgeries and therapeutic interventions. In addition, results may influence treatment plans for individuals with cancer. If a pathogenic variant is identified in one of these genes, the patient and/or their family members may be at increased risk for specific cancers and / or genetic conditions. Genetic counselling is recommended for these families.

**Indications for Testing**

Patients presenting with features suggestive of one of the below conditions are eligible for testing.

**Ordering privileges**

This panel may be ordered by Clinical Geneticists.

**Pediatric cancer NGS panel**

<b>Gene(s)</b>	<b>Associated cancers and/or clinical features<sup>2,3</sup></b>	<b>Associated Hereditary Syndrome(s)<sup>2,3</sup></b>
<i>ALK, PHOX2B</i>	Susceptibility to neuroblastoma, increased risk for other cancers	Hereditary neuroblastoma
<i>APC</i>	Colonic polyps, colon cancer, hepatoblastoma	APC-associated polyposis conditions
<i>BUB1B, CEP57</i>	Intrauterine growth restriction, intellectual disabilities, CNS anomalies and cancer predisposition such as Wilms tumor, rhabdomyosarcoma and leukemia	Mosaic variegated aneuploidy syndrome (types 1 & 2)
<i>CDKN1C</i>	Neonatal hypoglycemia, macrosomia, macroglossia, hemihyperplasia, omphalocele, embryonal tumors, visceromegaly, renal anomalies and ear creases / pits	Beckwith-Wiedemann syndrome
	<i>Of note: If a patient is suspected of having BWS, the first line test should be assessment for uniparental disomy of 11p15 or abnormal methylation or deletions of this region, with reflex to CDKN1C sanger sequencing. This testing is available through the Calgary Molecular Genetics Lab.</i>	
<i>DICER1</i>	A tumor susceptibility syndrome that confers an increased risk for pleuropulmonary blastoma (most commonly), ovarian sex cord-stromal tumors, juvenile granulosa cell tumor and thyroid gland neoplasia	DICER1-related disorders
<i>PTCH1, SUFU</i>	Multiple jaw keratocysts and/or basal cell carcinoma, characteristic facial appearance (~60%), skeletal anomalies, medulloblastoma	Nevoid Basal Cell Carcinoma syndrome
<i>SMARCB1</i>	Susceptibility to rhabdoid tumors, onset is typically around 6 months of age	Rhabdoid tumor predisposition syndrome
<i>TP53</i>	Lifetime risk of 68%-93% to develop cancer. The most common tumor types include soft tissue, osteosarcomas, breast cancer and brain cancer	Li-Fraumeni syndrome



WT1	Genital anomalies, cardiac malformations, Wilms tumor and renal disease.	Denys-Drash syndrome Frasier syndrome Meacham syndrome
	Wilms tumor, anirida, genitourinary anomalies, intellectual disability	WAGR
	Susceptibility to Wilms tumor	

**Associated Disorders<sup>2,3</sup>**

Some of the genes on this panel are associated with other rare disorders including:

**Congenital central hypoventilation syndrome (CCHS)** is an autosomal recessive disorder caused by pathogenic variants in *PHOX2B*. It is a disorder of respiratory and autonomic regulation and most commonly presents in the newborn period with a classic presentation, however, it can present more mildly at later ages.

**Schwannomatosis** is an autosomal dominant disorder characterized by multiple schwannomas without the vestibular schwannomas diagnostic of neurofibromatosis type 2. Pathogenic variants in *SMARCB1* are responsible for 30-60% of familial schwannomatosis but only a small percentage of isolated cases.

Pathogenic variants in *PTCH1* have been described in some individuals with **holoprosencephaly** and **microcephaly**.

**When can I expect results?**

Results may take up to 4 months.

**How are results reported?**

Results are sent to the ordering provider and available in Netcare and Connect Care.

**Contact Information**

Genetic Counsellors, Genetics & Genomics  
Calgary: 403-955-3097

**Requisition forms, contact information and other resources can be found at:**

<http://ahsweb.ca/lab/if-lab-genetics-and-genomics>

**References**

1. National Cancer Institute [Internet]: National Institutes of Health (US). Childhood Cancers. [cited 2022 Sept 1] Available from: <https://www.cancer.gov/types/childhood-cancers#:~:text=About%205%20percent%20of%20all,cell%20growth%20and%20eventually%20cancer>.
2. MedlinePlus [Internet]. Bethesda (MD): National Library of Medicine (US); [cited 2022 Sept 1]. Available from: <https://medlineplus.gov/>;
3. Adam MP, Ardinger HH, Pagon RA, et al., editors. GeneReviews® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2018 [cited 2017 Dec]. Available from: <https://www.ncbi.nlm.nih.gov/books/NBK1116/>