

SHH MEDULLOBLASTOMA GENE PANEL DG 3.2.0 (8 genes)

Releasedate: 16-09-2021

| Gene | Agilent V5 covered >10x | Agilent V5 covered >20x | TWIST covered >10x | TWIST covered >20x | Associated Phenotype Description and OMIM disease ID |
|---------|-------------------------|-------------------------|--------------------|--------------------|--|
| BRCA2 | 99,1 | 98,2 | 100 | 100 | Fanconi anemia, complementation group D1, 605724 Wilms tumor, 194070 |
| ELP1 | 99,8 | 98,9 | 100 | 100 | Dysautonomia, familial, 223900 |
| GPR161 | 100 | 100 | 100 | 100 | No OMIM disease ID |
| PALB2 | 100 | 99,9 | 100 | 100 | Fanconi anemia, complementation group N, 610832 |
| PTCH1 | 99,3 | 96,6 | 100 | 99,9 | Basal cell carcinoma, somatic, 605462 Holoprosencephaly 7, 610828 Basal cell nevus syndrome, 109400 |
| SMARCB1 | 100 | 99,9 | 100 | 100 | Rhabdoid tumors, somatic, 609322 Coffin-Siris syndrome 3, 614608 |
| SUFU | 100 | 100 | 100 | 100 | Joubert syndrome 32, 617757 Medulloblastoma, desmoplastic, 155255 Basal cell nevus syndrome, 109400 |
| TP53 | 99 | 95,2 | 91,7 | 91,7 | Hepatocellular carcinoma, somatic, 114550 Breast cancer, somatic, 114480 Li-Fraumeni syndrome, 151623 Pancreatic cancer, somatic, 260350 Nasopharyngeal carcinoma, somatic, 607107 Bone marrow failure syndrome 5, 618165 |

Gene symbols used follow HGNC guidelines: Gray KA, Yates B, Seal RL, Wright MW, Bruford EA. Nucleic Acids Res. 2015 Jan 43(Database issue):D1079-85.

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Agilent V5 is the default chemistry, and used for all exome analyses apart from the (in-house) TURBO/RAPID WES route.

TWIST is the chemistry used for (in-house) TURBO/RAPID WES analysis.

Covered 10x describes the percentage of a gene's coding sequence that is covered at least 10x.

Covered 20x describes the percentage of a gene's coding sequence that is covered at least 20x.

Genes with coverage denoting NC are non-protein coding genes.

non-protein coding genes are covered, but as coverage statistics are based on protein coding regions, statistics could not be generated.

OMIM release used for OMIM disease identifiers and descriptions : September 16th , 2021.

This list is accurate for panel version DG 3.2.0

Ad 1. "No OMIM Disease ID" signifies a gene without a current OMIM association Ad 2. OMIM phenotype descriptions between {} signify risk factors
