

This table includes regions of low-quality coverage for the Hearing Loss panel and therefore variant calling may be limited.

Gene	Genomic Region (Hg19)	Gene Region	Canonical Transcript
<i>ADCY1</i>	chr7:45614143-45614412	Exon 1	NM_021116.4
<i>CDKN1C</i>	chr11:2906063-2906229	Exon 2	NM_001122631.2
<i>DNMT1</i>	chr19:10290863-10290910	Exon 5	NM_001318731.2
<i>ESPN</i>	chr1:6485016-6485114	Exon 1	NM_031475.3
<i>ESPN</i>	chr1:6508865-6509069	Exon 8	NM_031475.3
<i>FOXC1</i>	chr6:1611259-1611904	Exon 1	NM_001453.3
<i>GIPC3</i>	chr19:3585596-3585679	Exon 1	NM_133261.3
<i>PEX6</i>	chr6:42946642-42946677	Exon 1	NM_001316313.2
<i>ROR1</i>	chr1:64240089-64240179	Exon 1	NM_005012.4
<i>SIX5</i>	chr19:46271300-46272102	Exon 1	NM_175875.5
<i>SPNS2</i>	chr17:4402343-4402481	Exon 1	NM_001124758.3
<i>TBX1</i>	chr22:19754013-19754390	Exon 7	NM_001379200.1
<i>TPRN</i>	chr9:140094597-140095162	Exon 1	NM_001128228.3

If you are a clinician and have a strong clinical suspicion of a disorder associated with a condition in a gene listed in this table, please contact the laboratory and additional orthogonal sequencing may be considered.