

Additional file 3: Table S3. The internal consistency reliability on 14 samples with known genotypes

Gene	Variant	Zygosity	Reads ratio		
			time 1	time 2	time 3
<i>SLC26A4</i>	c.919-2A>G	heterozygous	0.54	0.55	0.51
<i>GJB2</i>	c.235delC	heterozygous	0.54	0.51	0.51
<i>GJB2</i>	c.109G>A	heterozygous	0.46	0.46	0.46
<i>GJB2</i>	c.109G>A	heterozygous	0.49	0.44	0.44
<i>GJB2</i>	c.109G>A	heterozygous	0.47	0.45	0.46
<i>GJB2</i>	c.176_191del	heterozygous	0.54	0.56	0.65
<i>GJB2</i>	c.109G>A	heterozygous	0.46	0.48	0.48
<i>GJB2</i>	c.299_300delAT	heterozygous	0.56	0.48	0.46
<i>MT-RNR1</i>	m.1095T>C	homoplasmy	1	0.99	1
<i>GJB2</i>	c.299_300delAT	heterozygous	0.49	0.51	0.51
<i>GJB2</i>	c.235delC	heterozygous	0.49	0.57	0.66
<i>GJB2</i>	c.109G>A	heterozygous	0.46	0.44	0.44
<i>SLC26A4</i>	c.919-2A>G	heterozygous	0.6	0.52	0.54
<i>GJB2</i>	c.109G>A	heterozygous	0.48	0.48	0.43