

**Table S3 Description of primers used for pyro-sequencing-based allelic imbalance analyses.**

<b>Chromosome</b>	<b>SNP position</b>	<b>Forward primer</b>	<b>Reverse primer</b>	<b>Sequencing primer</b>
<b>3</b>	46581638	[BTN]--GTGCAGCTAGTTACAACAG	CCTTGCCAAAGTGAGTGTT	TTACATTCATTGCATCG
<b>3</b>	46581695	[BTN]--GTGCAGCTAGTTACAACAG	CCTTGCCAAAGTGAGTGTT	ATTCTTGCAAAAAACATAAA

[BTN]-- Biotinylated on 5' end.