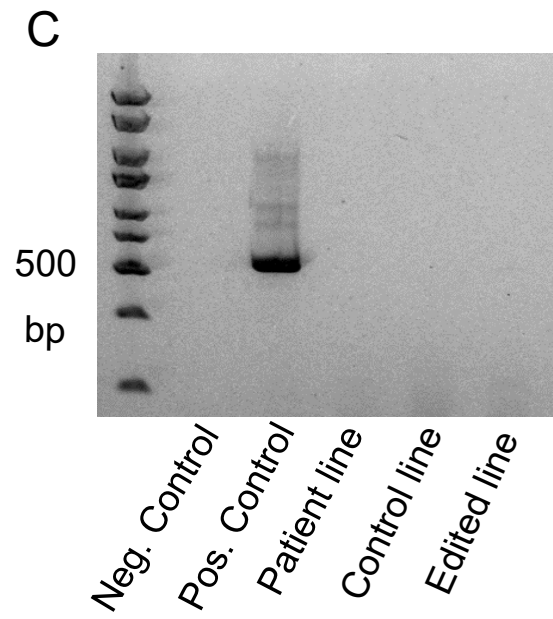
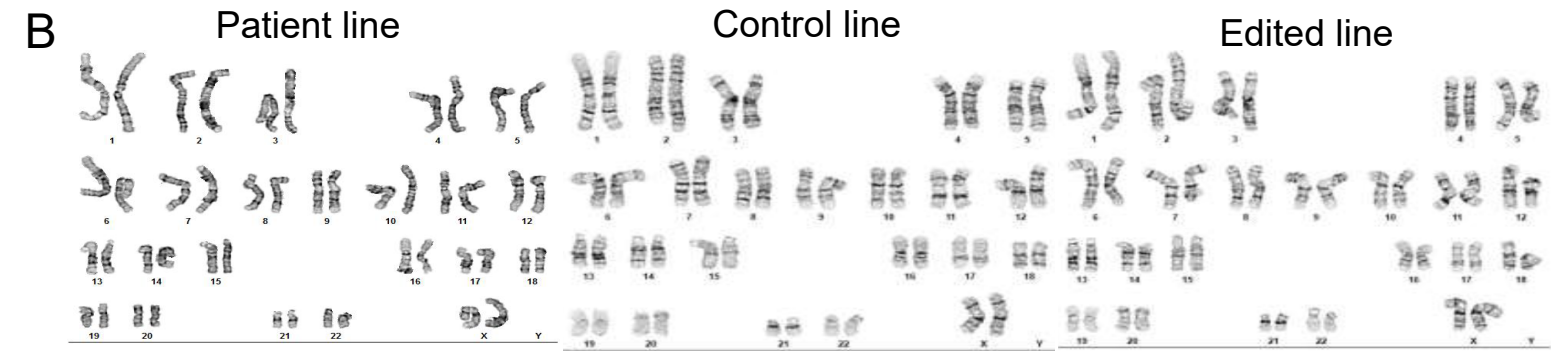
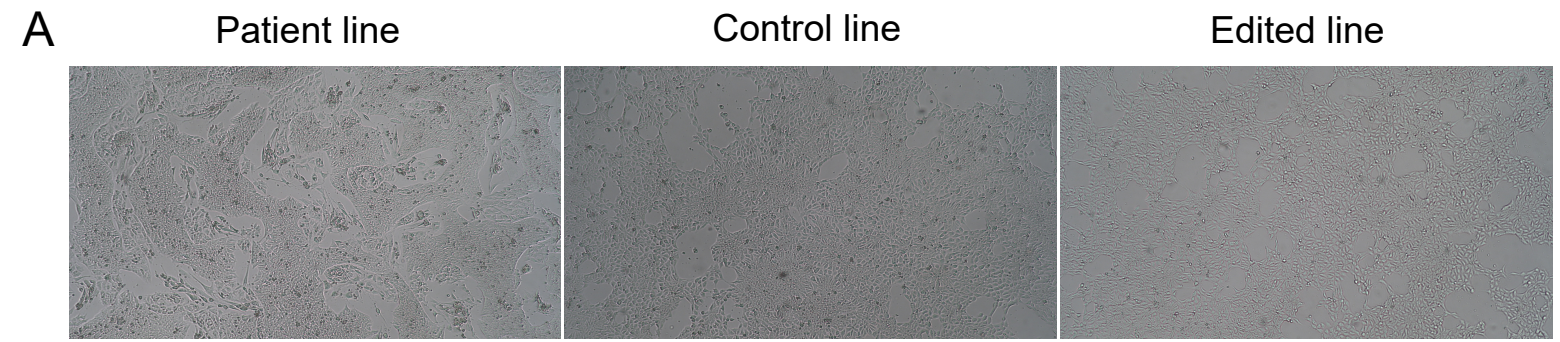


Supplemental Figure 1



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Chromosome	Position	Sequence	Strand	Mismatches
chr1	35931977	CTTAGGcCaTACTGCaCTGT(GGG)	+	3
chr1	244191886	CTTAGtTCCTACTGCTtTGc(TGG)	-	3
chr2	162266236	CTTtGccCCTACTGCTCTGT(TGG)	-	3
chr8	104220814	CTTAGGaCCTAaTGCTCTGc(AGG)	+	3
chr15	65034637	CTTtGGggCTACTGCTCTGT(GGG)	-	3
chr19	28389307	CTtGGTCCTgCTGCTCTGT(AGG)	+	3
chrY	8798586	CTcAGGTCTCaGCTCTt(GGG)	+	3