Table 2. Mutational Changes for p16 identified by Direct Sequencing

Cell line	Base	Codon	Sequence	Base change	Amino acid change
UACC-1097	247	83	gtgCacg	C→T	His→Tyr
UACC-1529	301	101	gccGggg	G→T	Gly→Trp
UACC-827	379	127	gtcGCACggt	GCAC→TAA	Frameshift (stop at codon 145)
M93-047	149	50	tccAggt	A→C	Gln→Pro
UACC-1273	Int1 –1/ 151	exon 2 splice	gcaGGtca	GG→AA	splicing defect
A-375	181	61	gcgGagc	G→T	Glu → stop
	205	69	gcgGagc	G→T	Glu → stop