

S5 Table. The sequencing information and % VAF enrichment from lung cancer cell lines

No.	Sample	Total reads	Coverage depth	Variant reads	% VAF
1	H1650 exon 19 PCR/-BDA	10,508	10,378	7,232	69.68
2	H1650 exon 19 PCR/+BDA	13,625	13,412	12,524	93.38
3	H1975 exon 20 PCR/-BDA	3,998	3,106	2,448	78.82
4	H1975 exon 20 BDA/+BDA	10,868	9,743	8,973	92.10
5	H1975 exon 21 PCR/-BDA	11,200	11,095	8,278	74.61
6	H1975 exon 21 PCR/+BDA	15,235	15,130	13,312	87.98

BDA, blocker displacement amplification; PCR, polymerase chain reaction; VAF, variant allele

frequency.