

Table S3. Control experiment

	Average input DNA (ng)	Somatic mutations	False Discovery Rate (%)	LOH of heterozygous SNPs (%)	False LOH (%)
Germline	34.1 (10-103)	-	-	-	-
Primary Tumor	22.4 (6.5-88.6)	222	0	5.2 (0.1-18)	1.6 (0.1-8.3)
Metastasis	24.9 (1.3-89.4)	706	2.2 (0-7)	18.9 (0-48) *	5.7 (0-16.2) *
Control 6ng	6.0	0	100	1	100
Control 0.6ng	0.6	27	100	0.6	100