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# **Data Sheet**

Product Name: CancerSeq AMS Paraffin Tissue Curl

Catalog No.: T2235152-AC

Lot No.: C202119

Species: ■Human □ Mouse □ Rat □ Monkey (Rh) □ Guinea Pig □ Porcine □ Bovine □ Hamster □ Dog □ Monkey (Cy) □ Rabbit Plant

Tissue Type: □ Normal ■ Adult □ Fetal ■ Tumor □Disease □ Cell line

Tissue Name: Lung

**Donor Information:** 

Male: \_\_\_\_\_ year(s) old Female: \_\_\_\_\_ year(s) old

Pathological Diagnosis: Adenocarcinoma

Tumor Size: diameter 5 cm

Location: right upper lobe

**Components:** 

- 1. 5 curls per package
- 2. Certificate of Analysis

APPROVED BY:

FOR IN VITRO RESEARCH USE ONLY



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## **35 Genes Targeted**

AKT1	FGFR2	MAP2K1		
ALK	FGFR3	MAP2K2		
AR	GNA11	MET		
BRAF	GNAQ	MTOR		
CDK4	HRAS	NRAS		
CTNNB1	IDH1	PDGFRA		
DDR2	IDH2	PIK3CA		
EGFR	JAK1	RAF1		
ERBB2	JAK2	RET		
ERBB3	JAK3	ROS1		
ERBB4	КІТ	SMO		
ESR1	KRAS			

### **Details of Variants**

Column Header	Definition						
Gene ID	The Gene symbol for the gene located at this position						
Chrom	The chromosome where the target region is located						
Position	The genomic position of the variant in the build of the genome database						
Ref	The reference allele of the variation						
Variant	The alternate allele of the variation						
Allele Call	The type of variation, either heterozygous or homozygous						
Frequency	The percentage of reads for the sample that includes the variant						
Quality	The quality score of the variant						
Туре	The variant type, which can be SNP, MNP, Ins, Del, and Complex						
Allele Source	Listed as Hotspot for alleles found within the hotspots sequencing file and Novel for all other alleles						
Coverage	The number of reads that cover the region						
Allele Name	The allele name that is defined within the hotspots sequencing file (if Novel allele, then there is no name)						

#### C202119

Gene ID	Chrom	Position	Ref	Variant	Allele Call	Frequency	Quality	Туре	Allele Source	Coverage	Allele Name
ESR1	chr6	152420011	С	Т	Homozygous	100	262	SNP	Novel	20	
RET	chr10	43613843	G	Т	Homozygous	100	262	SNP	Novel	20	
KRAS	chr12	25380290	G	А	Homozygous	100	297	SNP	Novel	22	