

# Standard-of-Care Tissue Profiling

## Targeted panel for swift results

**Standard40 Tissue NGS (Lung, Colon And Solid Tumors)** is a next-generation sequencing (NGS) test identifying targeted somatic genomic alterations in 40 genes using DNA and RNA isolated from tumor tissue with results in up to 7 working days.

This aids in identification of multiple disease alterations for which FDA/HSA approved and CDL-covered treatments are available, with support for identification of available clinical trials.

The test covers all actionable targets for lung (9 guideline-recommended targets) and colon (*NRAS*, *KRAS*, *BRAF*, *ERBB2*, *NTRK*).

## Targets list

### Genes

|              |                 |              |             |               |               |                 |             |
|--------------|-----------------|--------------|-------------|---------------|---------------|-----------------|-------------|
| <i>AKT1</i>  | <i>CDKN2A</i> # | <i>FGFR1</i> | <i>GNAQ</i> | <i>KIT</i>    | <i>MTOR</i>   | <i>PIK3CA</i> # | <i>SMO</i>  |
| <i>ALK</i> # | <i>CTNNB1</i>   | <i>FGFR2</i> | <i>GNAS</i> | <i>KRAS</i> # | <i>NRAS</i>   | <i>PTEN</i> #   | <i>TP53</i> |
| <i>AR</i> #  | <i>EGFR</i> #   | <i>FGFR3</i> | <i>HRAS</i> | <i>MAP2K1</i> | <i>NTRK1</i>  | <i>RAF1</i>     |             |
| <i>ARAF</i>  | <i>ERBB2</i> #  | <i>FLT3</i>  | <i>IDH1</i> | <i>MAP2K2</i> | <i>NTRK3</i>  | <i>RET</i>      |             |
| <i>BRAF</i>  | <i>ESR1</i>     | <i>GNA11</i> | <i>IDH2</i> | <i>MET</i> #  | <i>PDGFRA</i> | <i>ROS1</i>     |             |

#: Includes detection of gene copy number changes.

### Fusions & Splice variants

|             |             |              |              |              |              |              |             |
|-------------|-------------|--------------|--------------|--------------|--------------|--------------|-------------|
| <i>ALK</i>  | <i>EGFR</i> | <i>FGFR1</i> | <i>FGFR3</i> | <i>NRG1</i>  | <i>NTRK2</i> | <i>NUTM1</i> | <i>ROS1</i> |
| <i>BRAF</i> | <i>ESR1</i> | <i>FGFR2</i> | <i>MET</i>   | <i>NTRK1</i> | <i>NTRK3</i> | <i>RET</i>   |             |

## Test Specifications<sup>1</sup>

### Methodology

Ultra-deep next-generation sequencing

### Biomarkers analyzed

SNVs, Indels, CNVs, Fusions & Splice variants

### Sample type

FFPE tumor tissue

### Turnaround time

Up to 7 working days

### SNVs/Indels

**Sensitivity**

98%

**Specificity**

100%

### Fusions

91.25%

100%

- Results tested at the stated mutant allele frequencies using reference standards, FFPE cell line samples, and FFPE clinical samples.
- Sensitivity and specificity reported for SNVs and Indels are at 5% VAF.

References:

[1] Ng, CC-Y. et al. *Front. Mol. Biosci.* 2022. 9:963243.

This test is facilitated through Parkway Labs | Contact to order: 6933 0801/6248 5807 | [sgapadmin@parkwaypantai.com](mailto:sgapadmin@parkwaypantai.com)

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