

Solid Tumor v2 Panel

The **oncoReveal™ Solid Tumor v2 Panel** is a robust 48-gene assay that simultaneously tests for key mutations present in solid tumors, including NSCLC, colorectal, melanoma, endometrial, pancreatic, GIST, bladder, thyroid, and gliomas. Additionally, genes with potential importance in immuno-oncology such as POLD1 and POLE are analyzed. The panel uses proprietary Stem-Loop Inhibition-Mediated amplification (SLIMamp®) technology, a tiled amplicon-based library prep chemistry for efficient single-tube target enrichment.

oncoReveal[™] Solid Tumor v2 Panel (48 genes)

| AKT1 | CYSLTR2 | FBXW7 | GNAS | KEAP1 | NTRK1 | PTEN* | SMAD4 |
|--------|---------|-------|-----------------|--------|--------|--------|-------|
| ALK | DDR2 | FGFR1 | H3F3A (H3-3A) | KIT | PDGFRA | PTPN11 | SRSF2 |
| ARAF | EGFR | FGFR2 | HIST1H3B (H3C2) | KRAS | PIK3CA | RAC1 | STK11 |
| BRAF | EIF1AX | FGFR3 | HRAS | MAP2K1 | PLCB4 | RAF1 | TERT |
| CDKN2A | ERBB2 | GNA11 | IDH1 | MET | POLD1 | RET | TP53 |
| CTNNB1 | ERBB4 | GNAQ | IDH2 | NRAS | POLE | SF3B1 | TSHR |

Genes marked in green indicate full CDS coverage

Simple NGS library prep workflow

Maintain control of samples and results with single-tube, tiled amplification that can be performed in-house by any NGS lab

Sensitive and robust chemistry

Achieve variant detection as low as 1% VAF[†] without UIDs[‡] even with limited DNA input or poor sample quality

Reduced fullyloaded lab costs

Improve lab efficiency and reduce "no calls", repeat testing, and difficult interpretation decisions

Simple, one-day workflow



Gene-specific PCR & cleanup

Indexing PCR & cleanup

Quantitation & normalization

Load sequencer

^{*} indicates full CDS with exception of exon 9, chr10; 89725157-89725229

Panel specifications*

| Enrichment chemistry | Multiplex PCR using tiled amplicons | | |
|--|--|--|--|
| Number of pools | 1 pool | | |
| Number of genes/amplicons | 48/246 | | |
| Total panel size | Full CDS coverage of 3 genes; hotspots in 45 additional genes; 25.2kb total size | | |
| Variant types | SNVs, small and medium indels | | |
| Average amplicon size | 154bp | | |
| Recommended DNA input range | 20ng to 80ng | | |
| Sample types | DNA from tissue, blood, or FFPE | | |
| Mapping rate | 98.0% | | |
| % on-target aligned reads | 98.0% | | |
| Coverage uniformity (% targets with >0.2X mean coverage) | 98.0% | | |
| Recommended Reads Per Sample | ~2 million paired-end reads | | |
| Total assay time (from DNA to sequencer) | <8 hours | | |
| | | | |

^{*} Mapping rate, percentage of on-target aligned reads, and coverage uniformity metrics are based on internal testing performed using reference standard materials

Ordering information

Select the panel AND one of the index kit options listed below.

| Panel | Part number |
|---|----------------|
| oncoReveal™ Solid Tumor v2 Panel (24 reactions) | HDA-CH-3003-24 |

| Pillar Index Kit Options | Reactions | Part number |
|-----------------------------------|--------------------------------|-----------------|
| Pillar Custom Index Primers Kit A | 32 Combinations, 96 reactions | IDX-PI-1001-96 |
| Pillar Custom Index Primers Kit D | 96 Combinations, 192 reactions | IDX-PI-1004-192 |

TO ORDER OR LEARN MORE: pillarbiosci.com

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