



# AVENIO ctDNA Targeted Kit V2

## Tumor profiling: Guideline-driven biomarkers



The **AVENIO ctDNA Targeted Kit V2** is a next-generation sequencing (NGS) liquid biopsy tumor profiling research assay for identifying genomic aberrations in solid tumors. This panel contains **17** genes, including those in the U.S. National Comprehensive Cancer Network (NCCN) Guidelines.<sup>1</sup>

### Benefits

- Report all four mutation classes—SNVs, indels, fusions, and CNVs—from just 10 – 50 ng of cfDNA.
- Delivers exceptional analytical performance supported by integrated digital error suppression (iDES) strategies, combining molecular barcodes with in silico error suppression techniques.<sup>2,3</sup>
- Analyze a variety of solid tumor indications for research using a single optimized DNA workflow for up to 16 samples at a time.
- Reduce operational complexity by obtaining reagents for cfDNA isolation from plasma, library prep, and target enrichment from a single trusted vendor.
- Receive an complete solution with the required reagents, a robust bioinformatics pipeline, and software for analysis and reporting to keep your lab at the forefront of cancer research.<sup>†</sup>

### Research indications

Lung, Colorectal, Breast, Gastric, Glioma, Melanoma, Ovarian, Thyroid, and Pancreatic



### Applications

- Non-invasive tumor profiling
- Non-invasive detection of resistance biomarkers



### Analytical metrics<sup>2</sup>

| Mutation Class                          | SNVs        |      | Indels      |      | Fusions     |      | CNVs** |
|---|-------------|------|-------------|------|-------------|------|--------|
| Mutant Allele Frequency/<br>Copy Number | 0.5%*       |      | 1.0%*       |      | 1.0%        |      |        |
| Sensitivity and PPV                     | Sensitivity | PPV  | Sensitivity | PPV  | Sensitivity | PPV  |        |
|   | >99%        | >99% | >99%        | >99% | >99%        | >99% |        |

\*The reported sensitivity and PPV is for 50 ng DNA input at the stated AF%. The kit has the ability to report variants down to 0.1% AF (with lower sensitivity), and can make libraries with as low as 10 ng.  
 \*\*The kit reports ERBB2, EGFR, and MET CNVs. Performance is dependent on various factors, including sample type, input DNA amount, percent tumor content, and the individual tumor's genomic copy number. Our technical study detected CNVs at dPCR-verified copy numbers, using a commercially available reference sample (50 ng). CNV specificity was high (PPV >99%). ERBB2 detected at 4.5 copies, MET detected at 4.5 copies.  
 †For Research Use Only. Not for use in diagnostic procedures.

Performance samples: Seraseq ctDNA Complete Mutation Mixes (SeraCare), healthy donor cfDNA.  
 Sensitivity and Positive Predictive Value (PPV) metrics based on observed product performance. Sensitivity and PPV performance reported per variant. Sensitivity was determined using commercially available reference samples containing verified mutations at the stated allele frequencies. SNV performance data based on hotspot calls. The AVENIO ctDNA Analysis Kit V2 can achieve >99.99% per base specificity across each of the panels. Stated performance requires at least 40 million reads per sample for all AVENIO ctDNA Panels V2. Sequencing performed on an Illumina NextSeq 500/550/550Dx (Research Use Only mode) instrument.

# Specifications

|             |                |
|-------------|----------------|
| Panel size  | 81 kb          |
| Sample size | 4 mL of plasma |
| cfDNA input | 10 - 50 ng     |

|                         |                                   |
|-------------------------|-----------------------------------|
| Reactions per kit       | 16                                |
| Turn-around time        | 5 days from extraction to results |
| Product/Material Number | 09733736001                       |

## Assay targets

| Gene  | Seq Target         | SNV | Indel* | Fusion** | CNV** |
|-------|--------------------|-----|--------|----------|-------|
| ALK   | Selected Regions   | •   | •      | •        |       |
| APC   | Selected Regions   | •   | •      |          |       |
| BRAF  | Selected Regions   | •   | •      |          |       |
| BRCA1 | All Coding Regions | •   |        |          |       |
| BRCA2 | All Coding Regions | •   |        |          |       |
| DPYD  | Selected Regions   | •   |        |          |       |
| EGFR  | All Coding Regions | •   | •      |          | •     |
| ERBB2 | All Coding Regions | •   | •      |          | •     |
| KIT   | Selected Regions   | •   | •      |          |       |

| Gene      | Seq Target         | SNV | Indel* | Fusion** | CNV** |
|-----------|--------------------|-----|--------|----------|-------|
| KRAS      | All Coding Regions | •   |        |          |       |
| MET       | All Coding Regions | •   | •      |          | •     |
| NRAS      | Selected Regions   | •   |        |          |       |
| PDGFRA    | Selected Regions   | •   |        |          |       |
| RET       | Selected Regions   | •   |        | •        |       |
| ROS1      | Selected Regions   | •   |        | •        |       |
| TP53      | All Coding Regions | •   |        |          |       |
| UGT1A1*** | Selected Regions   | •   |        |          |       |

All coding regions are based on the longest transcript from Ensembl build 82.

\* Indels are limited to variants in a pre-specified list of positions, referred to as "Loci of Interest", except for EGFR exon 19 long deletions, EGFR exon 20 long insertions and MET long insertions, which are not restricted to a pre-defined set of Indels.

\*\* Detection of Fusions and CNVs are limited to variants in a pre-specified list of positions, referred to as "Loci of Interest" in the AVENIO analysis software.

\*\*\* UGT1A1\*28 allele sequenced but not currently called by the AVENIO analysis software.

## AVENIO family of NGS Oncology Assays

AVENIO ctDNA Targeted Kit V2 is a part of the AVENIO family of NGS oncology research assays that include three ctDNA and three corresponding tumor tissue assays. By using the wider family of AVENIO assays, labs can obtain detailed molecular findings across all four mutation classes from plasma or tissue samples.

1. National Comprehensive Cancer Network. <http://www.nccn.org>, October 1, 2025.

2. Data on file with Roche.

3. Newman AM, Lovejoy AF, Klass DM, et al. Integrated digital error suppression for improved detection of circulating tumor DNA. *Nature Biotechnology*. 2016;34(5):547-555. doi:10.1038/nbt.3520.

**For Research Use Only. Not for use in diagnostic procedures.**

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**\*Required hardware:** Illumina® NextSeq™ 500/550/550Dx and Roche Oncology Analysis Server. NextSeq™ 500/550/Dx instruments and associated sequencing reagents are manufactured and sold by Illumina and are not supplied by Roche.

**For more information about the AVENIO ctDNA Analysis Kits, please contact your local Roche representative.**

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