



AVENIO Tumor Tissue Targeted Kit V2

Tumor Profiling: Guideline-driven biomarkers



The **AVENIO Tumor Tissue Targeted Kit V2** is a next-generation sequencing (NGS) 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.¹

Benefits

- Report all four mutation classes—SNVs, indels, fusions, and CNVs—from just 20 ng of amplifiable DNA.²
- Delivers exceptional analytical performance supported by enzymatic error suppression and molecular barcodes.²
- Analyze a variety of solid tumor indications for research using a single optimized DNA workflow for up to 24 samples at a time.
- Reduce operational complexity by obtaining reagents for DNA isolation from tissue, library prep, and target enrichment from a single trusted vendor.
- Receive a 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.[†]

Research indications

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



Applications

- Comprehensive genomic profiling of tumor tissue
- Detection of resistance biomarkers



Analytical metrics²

Mutation Class	SNVs		Indels		Fusions		CNVs*
Mutant Allele Frequency/ Copy Number	5%		5%		5%		Sample Dependent*
Sensitivity and PPV	Sensitivity	PPV	Sensitivity	PPV	Sensitivity	PPV	
	>99%	>99%	>99%	>99%	>99%	>99%	

Samples: FFPE tissue curls/sections, DNA input: ≥20 ng of FFPE DNA, total DNA amount for each sample determined by input QC.
 *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.

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 is panel wide. Indel, Fusion based on whitelist variants. AVENIO Tumor Tissue Analysis Kits V2 achieve >99.99% per base specificity across each of the panels. Stated performance requires at least 20 million reads per sample for Targeted, Expanded, and Surveillance Kits V2. Sequencing performed on an Illumina NextSeq 500/550/550Dx (Research Use Only mode) instrument.

Specifications

Panel size	81 kb
Sample size	2 x 10 µm FFPE curls/sections
DNA input*	20 ng of amplifiable DNA

* Total DNA amount for each sample was determined by input QC.

Reactions per kit	24
Turn-around time	5 days from extraction to results
Product/Material Number	Plate A version - 09733710001 Plate B version - 09733728001

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

1. National Comprehensive Cancer Network. <http://www.nccn.org>. October 1, 2025.
2. Data on file with Roche.

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™ 500/550/550Dx instruments and associated sequencing reagents are manufactured and sold by Illumina and are not supplied by Roche.

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

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