

OVERVIEW

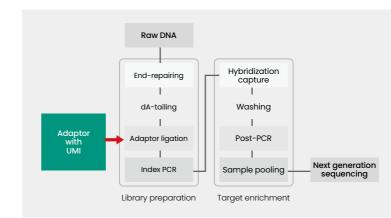
The detection sensitivity for low-frequency variants from a limited amount of sample is of great importance to ctDNA analysis kits. Celemics has developed ctDNA kits for colon, breast, and lung cancer assay through collaborative research with Seoul National University Hospital (SNUH) since 2017. We have integrated our market-leading proprietary technologies including probe design algorithms, noise removal techniques, and reagents optimization. The panels are thoroughly validated and ready to use for clinical diagnosis.

KEY FEATURES

1. Detects ctDNA for colorectal cancer, breast Assess 15 key genes for colorectal cancer, 27 for breast cancer, cancer, and lung cancer 28 for lung cancer Complete validated panel performance conducted with patient 2. Highly optimized panel for clinical testing with samples through collaborative research with Seoul National exceptional accuracy University Hospital Receive high-quality data supported by Celemics proprietary UMI 3. Provides Unique Molecular Identifiers (UMI) and algorithms and analysis software, enabling efficient duplication **Bioinformatics Software**

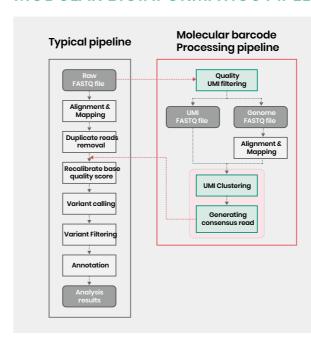
removal and minimizing sequencing noise

MODULAR UNIQUE MOLECULAR IDENTIFIER



- 1. Able to assess ctDNA with ultra-low variant allele frequency (VAF)
- 2. Retrieves more unique reads than that from conventional duplication removal algorithm, reducing sequencing costs
- 3. Noise removal and accurate calls possible due to proprietary consensus sequence generation algorithm
- 4. Modular algorithm to be applied to the existing pipeline.

MODULAR BIOINFORMATICS PIPELINE



- 1. Molecular Barcode Analysis Program provided to the customers using Celemics ctDNA Panels. (Linux, CLI program)
- 2. Streamlined application of the Molecular Barcode Analysis Program to the standard

PACKAGE COMPOSITION

Package name	Co	mposition	ns
Target Enrichment	Target capture Probe		-
Standard	Target Enrichment	Library	-
All-In-One	reagents	prep Kit	Beads / Polymerase

Package option	Options						
Pooling method	Single Reaction	Pre-capture Pooling					
Library Preparation kits	Standard Kit	EP-kit					
Hybridization Enhancer	Included	Not included					



Circulating-tumor DNA Colorectal Cancer Panel

SPECIFICATION

Gene count*	15 genes							
Covered region	Whole CDS							
Target size	49 kb							
Mutation type	SNV, Indel							
Sample type (amount)	Plasma (> 20 ng of cfDNA)							
Platform	All sequencers from Illumina and MGI							
Bioinformatics pipeline	1. Primary and Secondary analysis result (FASTQ to VCF) 2. Tertiary analysis result (VCF to Clinical report) 3. Linux-based consensus read generation software provided							

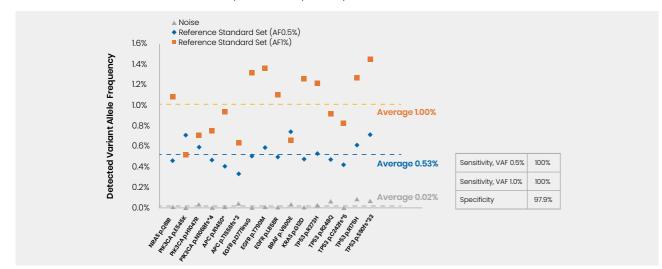
^{*} Gene Add-On Service: Genes can be added by customer's request

GENE LIST

ctDNA Panel	APC	ATM	BRAF	EGFR	ERBB2	FBXW7	KRAS	MET	NRAS	PDGFRA	PIK3CA	PTEN	SMAD4
Colorectal Cancer	TCF7L2	TP53											

PANEL PERFORMANCE

Detection of 16 variants with 100% sensitivity and 97.9% specificity at 0.5% VAF and 1% VAF



Circulating-tumor DNA Breast Cancer Panel

SPECIFICATION

Gene count*	27 genes								
Covered region	Whole CDS								
Target size	99 kb								
Mutation type	SNV, Indel								
Sample type (amount)	Plasma (> 20 ng of cfDNA)								
Platform	All sequencers from Illumina and MGI								
Bioinformatics pipeline	 Primary and Secondary analysis result (FASTQ to VCF) Tertiary analysis result (VCF to Clinical report) Linux-based consensus read generation software provided 								

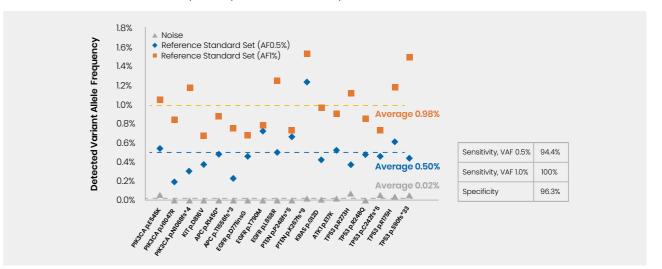
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GENE LIST

	AKT1	APC	AR	BRCAI	BRCA2	CCNDI	CDHI	EGFR	ERBB2	ESRI	FGFR1	FGFR2	GATA3
ctDNA Panel Breast Cancer	IGFIR	KIT	KRAS	MAP2K4	MAP3K1	MDM2	MYC	NFI	PIK3CA	PIK3R1	PTEN	RB1	TOP2A
	TP53												

PANEL PERFORMANCE

Detection of 27 variants with 96.3% specificity and 94.4% sensitivity at 0.5% VAF and 100% at 1% VAF



Circulating-tumor DNA Lung Cancer Panel

SPECIFICATION

Gene count*	28 genes						
Covered region	Whole CDS for 8 genes and Hotspot exonic region for 20 genes						
Target size	47 kb						
Mutation type	SNV, Indel						
Sample type (amount)	Plasma (> 20 ng of cfDNA)						
Platform	All sequencers from Illumina and MGI						
Bioinformatics pipeline	 Primary and Secondary analysis result (FASTQ to VCF) Tertiary analysis result (VCF to Clinical report) Linux-based consensus read generation software provided 						

^{*} Gene Add-On Service: Genes can be added by customer's request

GENE LIST

	AKT1	ALK	ARAF	ARID1A	BRAF	CBL	CDKN2A	EGFR	ERBB2	HRAS	KEAPI	KRAS	MAP2K1
ctDNA Panel Lung Cancer	MET	MTOR	NFI	NRAS	NTRK1	NTRK2	PIK3CA	PTEN	RBI	RIT1	ROSI	SETD2	STKII
	TP53	U2AF1											

PANEL PERFORMANCE

Detection of 28 variants with 100% sensitivity and 100% specificity at 0.5% VAF and 1% VAF detection

