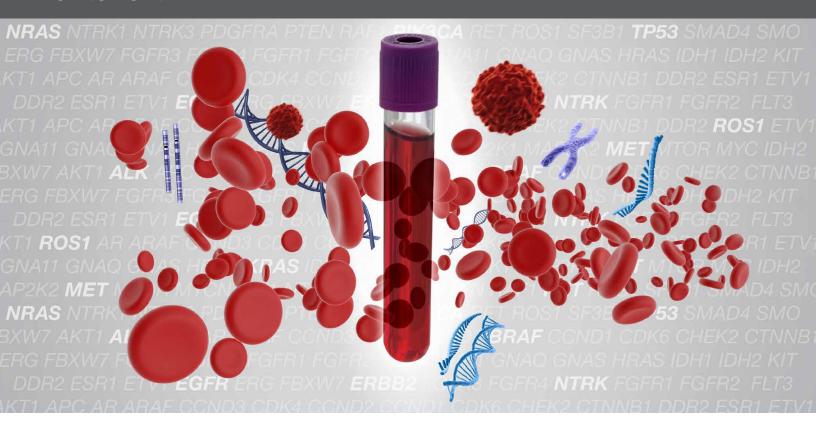
ion torrent



Targeted sequencing for liquid biopsy cancer research

Growing menu of assays provides new content for more relevant insights

Ion Torrent™ Oncomine™ liquid biopsy assays are highly sensitive, multibiomarker next-generation sequencing (NGS) assays that enable molecular insights into tumor evolution and detection of primary driver and resistance mutations for clinical research. The high-value gene content includes key mutations identified by the OncoNetwork consortium and other clinical researchers around the world.

Based on multiplex PCR, our proprietary technology together with Ion Torrent™ sequencing enables researchers to study biomarkers that may impact treatment selection, treatment monitoring, and recurrence monitoring in the future.

Key benefits

- Streamlined workflow—complete NGS research workflow, from blood sample to variant data in 1 to 3 days
- Optimized content—SNVs, indels, CNVs, and fusions for multibiomarker analysis
- Flexible input amounts—results enabled from one tube of blood
- Low limit of detection—variant detection down to 0.1%



The family of Oncomine liquid biopsy assays

Oncomine liquid biopsy assays enable research studies on tumor heterogeneity and recurrence, from minimal sample input (Figure 1). The assays achieve high correlation between variants called in formalin-fixed, paraffin-embedded (FFPE) samples and in cell-free DNA (cfDNA) from plasma (Table 1).

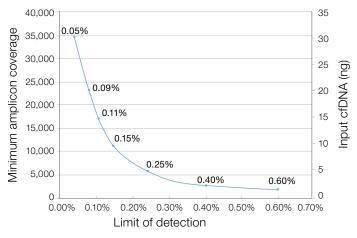


Figure 1. Amplicon coverage and input material determine limit of detection using Oncomine liquid biopsy assays.

Built on proprietary, amplification-based technology, Oncomine liquid biopsy assays enable detection of driver and resistance mutations from cell-free nucleic acids, down to 0.1%. Select from five focused, tumor type—specific assays predesigned with key gene content, or two broad pan-cancer assays that cover all classes of mutations across multiple cancer types (Figure 2).

Table 1. Correlation between results from FFPE and matched plasma samples (late-stage lung cancer samples).

Sample	Variant	FFPE	Plasma
1	EGFR-L858R	71.42%	2.62%
2	TP53-R158L	51.89%	4.32%
3	MET-T1010I	43.87%	51.75%
	KRAS-G12C	34.62%	0.28%
4	NA	No detection	No detection
	EGFR-L858R	58.44%	7.28%
5	MET-T1010I	41.93%	48.72%
	TP53-Y220C	35.54%	1.93%
6	TP53-R158L	10.19%	1.26%

Values in boldface indicate somatic mutations; values not in boldface indicate germline mutations. As expected, there is a higher fraction of somatic mutations in FFPE samples than in plasma samples. Germline variants are seen at the expected levels of ~50% in both sample types. Data were obtained using the Oncomine Lung cfDNA Assay.

4				В	[S	Colon				
Oncomine Lung cfDNA Assay		Oncomine Lung Cell-Free Total Nucleic Acid Assay		Oncomine Breast cfDNA Assay		Oncomine Breast cfDNA Assay v2		Oncomine Colon cfDNA Assay		
ALK	MET	ALK	MET	AKT1	FBXW7	AKT1	FBXW7	AKT1	ERBB2	NRAS
BRAF	NRAS	BRAF	NRAS	EGFR	KRAS	CCND1	FGFR1	APC	FBXW7	PIK3CA
<i>EGFR</i>	PIK3CA	EGFR	PIK3CA	ERBB2	PIK3CA	EGFR	KRAS	BRAF	GNAS	SMAD4
ERBB2	ROS1	ERBB2	RET	ERBB3	SF3B1	ERBB2	PIK3CA	CTNNB1	KRAS	TP53
KRAS	TP53	KRAS	ROS1	ESR1	TP53	ERBB3	SF3B1	EGFR	MAP2K1	
MAP2K1		MAP2K1	TP53			ESR1	TP53			

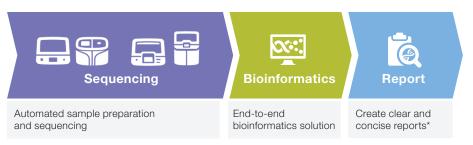
Pan-cancer

						, I dil Gdil	001				
				0	ncomine l	Pan-Cancer	Cell-Free Ass	say			
DNA hots	spots					Tumor sup	opressors	CNVs		Fusions	
AKT1	EGFR	FLT3	KRAS	PDGFR/	4	APC		CCND1	ERBB2	ALK	FGFR3
ALK	ERBB2	GNA11	MAP2K1	PIK3CA		FBXW7		CCND2	FGFR1	BRAF	MET
AR	ERBB3	GNAQ	MAP2K2	RAF1		PTEN		CCND3	FGFR2	ERG	NTRK1
ARAF	ESR1	GNAS	MET	RET		TP53		CDK4	FGFR3	ETV1	NTRK3
BRAF	FGFR1	HRAS	MTOR	ROS1				CDK6	MET	FGFR1	RET
CHEK2	FGFR2	IDH1	NRAS	SF3B1				EGFR	MYC	FGFR2	ROS1
CTNNB1	FGFR3	IDH2	NTRK1	SMAD4							
DDR2	FGFR4	KIT	NTRK3	SMO							
					Oncor	nine Precisi	on Assay GX				
DNA hots	spots						CNVs		Fusions		
AKT1	CDKN2A	FGFR1	HR/	IS	MTOR	RAF1	ALK	FGFR1	ALK	FGFR3	RET
AKT2	CHEK2	FGFR2	? IDH	1	NRAS	RET	AR	FGFR2	AR	MET	ROS1
AKT3	CTNNB1	FGFR3	B IDH2	2	NTRK1	ROS1	CD274	FGFR3	BRAF	NRG1	RSPO2
ALK	EGFR	FGFR4	t KIT		NTRK2	SMO	CDKN2A	KRAS	EGFR	NTRK1	RSPO3
AR	ERBB2	FLT3	KRA	S	NTRK3	TP53	EGFR	MET	ESR1	NTRK2	
ARAF	ERBB3	GNA11	MAF	P2K1	PDGFRA		ERBB2	PIK3CA	FGFR1	NTRK3	
BRAF	ERBB4	GNAQ	MAF	P2K2	PIK3CA		ERBB3	PTEN	FGFR2	NUTM1	
CDK4	ESR1	GNAS	ME7	-	PTEN						

Figure 2. Gene content of Oncomine liquid biopsy assays. (A) Gene lists for five assays that target breast, colon, and lung cancer. (B) Gene lists for two pan-cancer assays, Oncomine Pan-Cancer Cell-Free Assay and Oncomine Precision Assay GX.

Streamline your targeted sequencing workflow

The Oncomine liquid biopsy NGS workflow consists of three key steps (Figure 3). First, cell-free nucleic acids are extracted, enriched, and amplified from one tube of blood. These amplicon-based libraries are then assembled and prepped for targeted resequencing. Our integrated informatics solution then takes you from variant caller to a clear and concise report that links biomarkers to relevant evidence (Figure 4). This process transforms data into knowledge, helping you gain efficiency for cancer research and future drug development.



^{*} Specimen-to-report workflow will be available after the lon Torrent™ Genexus™ Purification System and integrated reporting capabilities are added in 2021. The content herein may relate to products or workflows that have not been officially released or fully validated and is subject to change without notice.

Figure 3. Our 1- to 3-day comprehensive NGS workflow for liquid biopsy provides streamlined detection and analysis of genes and key mutations.

Example Labs 123 Street City, ST 12345 USA email@examplelabs.com www.exampleiabs.com Sample ID: 00-123456789 Date: 29 Sep 2020 Sample Type: Primary Tumor Site: Lung 010101 Sample Collected: Sample Cancer Type: Non-Small Cell Lung Cancer Relevant Non-Small Cell Lung Cancer Findings Finding Not detected NTRK1 Not detected ALK BRAF Not detected NTRK2 Not detected EGFR. EGFR E746 A750del (EGFR expn 19 NTRK3 Not detected deletion), EGFR T790M ERBB2 Not detected RET Not detected KRAS Not detected ROS1 Not detected MET Not detected Relevant Biomarkers Genomic Alteration EGFR E746_A750del EDDIT even 18 deletions Long dy 7:55242464 Transcript: NVL,005228.4 EGFR T790M Transcript: NW, 005229 f. IIC 7P53 P60fs tumor protein p53 Lacux 89/17/7979508 Transcript: NVL,000545-5

"What's in the blood is actually what's relevant. One could argue it's not the overall tumor composition that we really want. We want the biologically relevant population. How do we define that? If that's in the blood, that's what we should be sampling—it's that global representation of tumor biology that we need."

—Minetta Liu, MDAssociate Professor of Oncology,Mayo Clinic

"Genomic heterogeneity after anti-EGFR therapy was successfully detected in 94% of the metastatic colorectal cancer research samples using a next-generation sequencing multibiomarker approach. The Oncomine Colon cfDNA Assay enabled detection of somatic mutations with MAF >0.1%."

-Beatriz Bellosillo, PhD Hospital del Mar, Spain

Figure 4. Sample report from Ion Torrent™ Oncomine™ Reporter. The final report will provide relevant and detailed biomarker insights for your sample(s).

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Ordering information

Product	Cat. No.
Liquid biopsy solutions on Ion GeneStudio S5 Systems	
Oncomine Pan-Cancer Cell-Free Assay	A37664
Oncomine Lung Cell-Free Total Nucleic Acid Assay	A35864
Oncomine Lung cfDNA Assay	A31149
Oncomine Breast cfDNA Assay v2	A35865
Oncomine Breast cfDNA Assay	A31183
Oncomine Colon cfDNA Assay	A31182
Tag Sequencing Barcode Set 1-24	A31830
Tag Sequencing Barcode Set 25-48	A31847
Ion GeneStudio S5 Prime System	A38196
Ion GeneStudio S5 Plus System	A38195
Ion GeneStudio S5 System	A38194
Ion 550 Chip Kit	A34538
Ion 540 Chip Kit	A27766
Ion 530 Chip Kit	A27764
Ion Chef System	4484177
Ion 550 Kit-Chef	A34541
Ion 540 Kit-Chef	A30011
lon 510 & lon 520 & lon 530 Kit-Chef	A34461
Liquid biopsy solutions on Ion Torrent Genexus System	
Oncomine Precision Assay GX	A46291
Ion Torrent Genexus System	A45727
Ion Torrent Genexus GX5 and Coupler	A40269
Ion Torrent Genexus GX5 Starter Pack-AS	A40279
Oncomine informatics	
Ion Reporter Server System	4487118
Oncomine Reporter	A33109

