Hemato-oncology

Oncomine Myeloid Assay GX v2 More rapid, automated, and complete myeloid genomic profiling

Myeloid malignancy samples can be challenging to analyze. They are complex, heterogeneous, and have the potential to proliferate rapidly. To get meaningful genetic insights, you need a rapid and streamlined approach for profiling all key mutations.

Traditional single-analyte testing approaches can be laborious and time-consuming—especially as the list of relevant genes continues to grow. With the Ion Torrent[™] Oncomine[™] Myeloid Assay GX v2 on the Ion Torrent[™] Genexus[™] System, you can get a comprehensive profile of myeloid mutations from a single next-generation sequencing (NGS) run and results in as little as one day.

A highly automated workflow lets you go from specimen to report with only 20 minutes of hands-on time. Integrated bioinformatic tools allow you to seamlessly generate a variant report with annotations based on the latest evidence.

Target important biomarkers associated with major myeloid disorders

With the ability to profile both DNA and RNA targets in a single test, the Oncomine Myeloid Assay GX v2 allows you to dramatically consolidate the number of individual tests typically required for profiling a broad spectrum of mutations, including both DNA mutations and translocations detected from RNA targets.

Simultaneously interrogate 45 DNA target genes and 35 RNA fusion driver genes. This broad fusion panel allows you to sequence over 700 unique fusion transcripts. Gene content on the panel has been curated to cover relevant targets for all the major myeloid disorders—AML, MDS, MPN, CML, CMML, and JMML.

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Highlights



Rapid turnaround time—from specimen to report in as little as one day



Highly automated workflow—process samples with just 20 minutes of hands-on time required



Comprehensive target coverage—simultaneously profile 45 key DNA genes and 35 fusion drivers (>700 unique fusions) relevant for the spectrum of major myeloid disorders



Trusted performance—reliably detect a range of variants, including challenging single nucleotide variants (SNVs) in homopolymer regions and important internal tandem duplications (ITDs) and partial tandem duplications (PTDs)



Integrated reporting—get annotated variants, including links to the latest relevant evidence, to help inform biological significance

Trusted performance for detecting key mutations

Myeloid samples frequently harbor mutations in difficult-to-sequence regions of the genome. Our assay primer design and chemistry has been carefully engineered to:

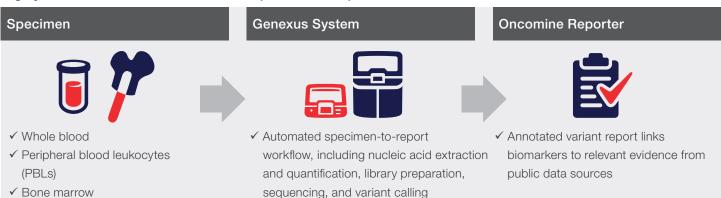
- Reliably detect all important mutations associated with myeloid malignancies
- Accurately detect SNVs in genes with long homopolymer regions like *CEBPA* and *ASXL1*

Table 1. Oncomine Myeloid Assay GX v2 gene targets.

- Detect a range of insertions and deletions, even in challenging genes like *CALR*
- Identify ITDs in *FLT3* and PTDs in *KMT2A*
- Get optimal performance for challenging insertions with integrated FTL3-ITD detection software

DNA panel: hotspot genes (28)		DNA panel: full genes (17)		RNA panel: fusion driver genes (35)			RNA panel: expression genes (5)	RNA panel: expression control genes (5)
ANKRD26	KRAS	ASXL1	PRPF8	ABL1	HMGA2	NUP98	BAALC	EIF2B1
ABL1	MPL	BCOR	RB1	ABL2	JAK2	NUP214	MECOM	FBXW2
BRAF	MYD88	CALR	RUNX1	BCL2	KAT6A (MOZ)	PAX5	MYC	PSMB2
CBL	NPM1	CEBPA	SH2B3	BRAF	KAT6B	PDGFRA	SMC1A	PUM1
CSF3R	NRAS	ETV6	STAG2	CCND1	KMT2A	PDGFRB	WT1	TRIM27
DDX41	PPM1D	EZH2	TET2	CREBBP	KMT2A PTDs	RARA		
DNMT3A	PTPN11	IKZF1	TP53	EGFR	MECOM	RUNX1		
FLT3 (ITD,	SMC1A	NF1	ZRSR2	ETV6	MET	TCF3		
TKD)	SMC3	PHF6		FGFR1	MLLT10	TFE3		
GATA2	SETBP1			FGFR2	MRTFA (MKL1)	ZNF384		
HRAS	SF3B1			FUS	MYBL1			
IDH1	SRSF2				MYH11			
IDH2	U2AF1				NTRK2			
JAK2	WT1				NTRK3			
KIT								

Highly automated NGS workflow-from specimen to report



Start with any common myeloid specimen type. From there, the Genexus System integrates and automates the entire workflow. With only 20 minutes of hands-on time required, your lab can be more efficient and free up staff's time to focus on other applications.

Each lane on the lon Torrent[™] GX5[™] Chip can accommodate up to eight samples in a sequencing run. The four-lane chip and reagents are stable on the instrument for two weeks, giving you the flexibility to batch samples across multiple runs without having to waste unused sequencing capacity.

Onboard, integrated analysis provides robust variant calling and reporting without the need for an external server. The simplified user experience helps minimize the learning curve and avoid human error. The Ion Torrent[™] Oncomine[™] Reporter is a curated knowledge-base and reporting software that links variants to relevant evidence and enables custom reporting. These tools help simplify the bioinformatics workflow and enable you to focus on finding the biological meaning of your data.

Specifications

Gene targets	28 hotspot genes (DNA), 17 full genes (DNA), 35 fusions drivers (RNA), >700 unique fusions, 5 expression genes, 5 expression controls				
Variants	SNVs, insertions, deletions, ITDs, PTDs, gene fusions				
Input	20 ng DNA, 10 ng RNA				
Turnaround time	22 hours for 8 samples (automated library prep, sequencing, reporting)				
Turnaround time	27 hours for 8 samples (automated extraction, purification, library prep, sequencing, reporting)				
Throughput	8 DNA and 8 RNA samples per lane, per run				
Throughput	32 DNA and RNA samples per chip				
Coverage	≥99%				
Sensitivity	≥95%				

We offer different assay configurations to meet your lab's unique needs. Profile DNA and RNA targets simultaneously or have the option to profile each independently.

Ordering information

Product	Cat. No.
Oncomine Myeloid Assay GX v2	A50694
Oncomine Myeloid DNA Assay GX v2	A50753
Oncomine Myeloid RNA Assay GX v2	A50754

Learn more about the Oncomine Myeloid Assay GX v2 at oncomine.com/myleoid