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Oncomine Comprehensive Assay v3

Empower your oncology research with proven Ion Torrent technology



The Oncomine Comprehensive Assay v3 is available for use on the Ion GeneStudio S5 Systems with Ion Chef Instrument or for use on the Genexus System

The Ion Torrent™ Oncomine™ Comprehensive Assay v3 is a member of the family of Ion Torrent™ Oncomine™ assays for clinical cancer research. Oncomine assays are multiple-biomarker assays based on next-generation sequencing (NGS). They have been adopted by leading cancer institutions around the world, have been used to profile thousands of samples in different translational and clinical research projects, and have consistently delivered reliable results.

Oncomine Comprehensive Assay v3

- Content is based on the latest advances in clinical oncology research and also enriched for targets known to be associated with (or drive) childhood cancers
- Based on robust Ion AmpliSeq[™] technology, this assay only requires 10 ng of DNA or RNA per pool, enabling analysis of even small and challenging formalin-fixed, paraffin-embedded (FFPE) samples

- Detects relevant SNVs, CNVs, gene fusions, and indels from 161 unique cancer driver genes in one streamlined workflow
- Optimized and verified for the lon Chef[™] Instrument and lon GeneStudio[™] S5 Systems with the lon 540[™] Chip, or the new lon Torrent[™] Genexus[™] System with the lon Torrent[™] GX5[™] Chip, both enabling full automation including automated library prep

"The requirement of a lower DNA input for the Oncomine assay is a significant advantage when primary samples are becoming increasingly limited."

John Bartlett, PhD

Director of Transformative Pathology Platform
Ontario Institute for Cancer Research



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Hotspot genes				Full-length genes			Copy number genes		Gene fusions (inter- and intragenic)		
AKT1 AKT2 AKT3 ALK AR ARAF AXL BRAF BTK CBL CCND1 CDK4 CDK6 CHEK2 CSF1R CTINNB1 DDR2 EGFR ERBB2 ERBB3 ERBB4 ERCC2	ESR1 EZH2 FGFR1 FGFR2 FGFR3 FGFR4 FLT3 FOXL2 GATA2 GNA11 GNAQ GNAS H3F3A HIST1H3B HNF1A HRAS IDH1 IDH2 JAK1 JAK2 JAK3 KDR	KIT KNSTRN KRAS MAGOH MAP2K1 MAP2K2 MAP2K4 MAPK1 MAX MDM4 MED12 MET MTOR MYC MYCN MYD88 NFE2L2 NRAS NTRK1 NTRK2 NTRK3 PDGFRA	PDGFRB PIK3CA PIK3CA PPP2R1A PTPN11 RAC1 RAF1 RET RHEB RHOA ROS1 SF3B1 SMAD4 SMO SPOP SRC STAT3 TERT TOP1 U2AF1 XPO1	ARID1A ATM ATR ATRX BAP1 BRCA1 BRCA2 CDK12 CDKN1B CDKN2A CDKN2B CHEK1 CREBBP FANCA FANCD2 FANCI	FBXW7 MLH1 MRE11 MSH6 MSH2 NBN NF1 NF2 NOTCH1 NOTCH2 NOTCH3 PALB2 PIK3R1 PMS2 POLE PTCH1	PTEN RAD50 RAD51 RAD51B RAD51C RAD51D RNF43 RB1 SETD2 SLX4 SMARCA4 SMARCB1 STK11 TP53 TSC1 TSC2	AKT1 AKT2 AKT3 ALK AXL AR BRAF CCND1 CCND2 CCND3 CCNE1 CDK2 CDK4 CDK6 EGFR ERBB2 ESR1 FGF19 FGF3 FGFR1 FGFR2 FGFR3	FGFR4 FLT3 IGF1R KIT KRAS MDM2 MDM4 MET MYC MYCL MYCN NTRK1 NTRK2 NTRK3 PDGFRA PDGFRB PIK3CB PIK3CB PIK3CA PPARG RICTOR TERT	AKT2 ALK AR AXL BRCA1 BRCA2 BRAF CDKN2A EGFR ERBB2 ERBB4 ERG ESR1 ETV1 ETV4 ETV5 FGFR1	FGFR2 FGR7 FGR FLT3 JAK2 KRAS MDM4 MET MYB MYBL1 NF1 NOTCH1 NOTCH4 NRG1 NTRK1 NTRK2 NTRK3	NUTM1 PDGFRA PDGFRB PIK3CA PRKACA PRKACB PTEN PPARG RAD51B RAF1 RB1 RELA RET ROS1 RSP02 RSP03 TERT

List of gene targets in the Oncomine Comprehensive Assay v3.

Intelligent NGS assay design

The content covers 161 of the most relevant cancer driver genes, including increased kinase domain coverage and representation of genes involved in DNA repair.

Optimized for challenging FFPE samples

Based on Ion AmpliSeq technology, the Oncomine Comprehensive Assay v3 typically requires three or fewer FFPE slides or as little as 10 ng of input DNA or RNA per reaction. This enables analysis of small and inferior-quality samples, such as fine-needle aspirates. Alternative methods require numerous FFPE slides and hundreds of nanograms of DNA or RNA, making them less practical for routine analysis of FFPE tumor samples.

Oncomine informatics

The Oncomine NGS oncology assay workflow provides a complete solution for NGS data analysis to go from hundreds of variants to a few relevant cancer drivers.

Ordering information

Product	No. of react	ions Cat. No.
For Ion GeneStudio S5 Systems		
Oncomine Comprehensive Assay v3C (automated library preparation with the lon Chef Instrument)	32	A35806
Oncomine Comprehensive Assay v3M	24	A35805
(manual library preparation)	96	A36111

The Oncomine informatics workflow enables you to get high-quality data, prioritizes and annotates driver variants, and comes complete with software that links relevant biomarkers to information available from labels, guidelines, and clinical studies for clinical/translational research support.

Different configurations to meet the needs of any laboratory

The Oncomine Comprehensive Assay v3 has been optimized for throughput of eight samples on the Ion GeneStudio S5 Systems or six samples on the Genexus System. The assay is available in two configurations for use on the Ion GeneStudio S5 Systems: manual and automated library preparation (on the Ion Chef Instrument). The Genexus System combines prep, templating, and sequencing all in one system, with two user touchpoints and a single-day turnaround time.*

Product	No. of reactions	Cat. No.
For Ion Torrent Genexus System		
	16 DNA/RNA	
Oncomine Comprehensive Assay v3 GX	32 DNA	A46296
	32 RNA	_
Oncomine Comprehensive Assay v3 DNA GX	16	A46294
Oncomine Comprehensive Assay v3 RNA GX	16	A46295

^{*} Specimen-to-report workflow will be available after the Ion Torrent™ Genexus™ Purification System and integrated reporting capabilities are added in 2020.

