

Oncomine Solutions for precision oncology research

True end-to-end workflows for your laboratory

Explore our educational website

Learn more about precision oncology research, next-generation sequencing, and Oncomine[™] Solutions at **oncomine.com**

- Learn about our comprehensive portfolio of products
- Hear from our customers
- Access on-demand webinars, infographics, and articles

Dr. Philip Jermann, PhD

Head of the Molecular Assay Development Unit of the Institute of Medical Genetics and Pathology, University Hospital Basel

"The Ion Torrent Genexus System is based on technology that has evolved over the last 10 years. I was an early adopter, so I have followed its entire development. The method, to the best of my knowledge, has been cited in more than 6,000 scientific papers. Recently, a seminal retrospective study from the University of Heidelberg in Germany that used the Ion Torrent technology demonstrated a nearly 97% sequencing success rate across more than 3,000 non-small cell lung cancer samples, and a rejection rate, mostly due to quantity-not-sufficient errors, of only around 3%."

Retrieved from "What is the real impact?" in *The Pathologist*, December 2019

- Subscribe to get the news first
- Follow our blog to read what we and our customers think about hot topics

Dr. Davide Soldini, MD, PhD

Director of the Institute of Molecular Pathology MEDICA, Zurich

"Oncomine Solutions means to me that I can perform the analysis, basically for all the samples that we get, even with a small amount of material".

Retrieved from "What Oncomine Solutions mean to you" video

Catch up on the Oncomine Clinical Research Grant Program, new calls and awardees

The Program supports investigator-initiated studies (IIS) and education projects, on molecular testing in oncology and reproductive health, with the goal to increase high quality molecular profiling for patients and to improve clinical outcomes in the future.

Follow where you can meet us and each other

We host and participate at numerous events around the globe, and recently virtually.

See where you can meet us to see our latest solutions, discuss with our product experts, and hear from your peers about their experiences at oncomine.com/events.



Watch videos about Oncomine Solutions





Download key biomarker testing guides



NSCLC testing guide



PIK3CA testing guide



testing guide



Oncomine Solutions

Oncomine Solutions offer end-to-end workflows that address specific challenges when implementing next-generation sequencing (NGS) for clinical oncology research. We understand that NGS applications and content need to be tailored for your lab's needs. That's why we offer two types of workflows, a broad assay portfolio, complete bioinformatics, and technical support, to suit the way you work.

conestudio S5 System 40

Specimen to repo

Nucleic acid to repo

Key genes for one-day genomic profiling

Minimum sample input requirement

2

Ion Torrent[™] Oncomine[™] Reporter software

22

Specialized implementation support

End-to-end protocols tested on clinical research samples

Broad assay menu across key applications

One size does not fit all—Overview of all Oncomine assays across key applications

Oncomine Solutions give you options because each individual, sample, and lab is unique

Hemato-oncology research solutions

- Ion Torrent[™] Oncomine[™] Myeloid Assay GX
- Ion Torrent[™] Oncomine[™] Myeloid Research Assay
- Ion Torrent[™] Oncomine[™] BCR Pan-Clonality Assay
- Ion Torrent[™] Oncomine[™] TCR Pan-Clonality Assay
- Ion Torrent[™] Oncomine[™] IGHV Leader-J Assay
- Ion Torrent[™] Oncomine[™] Lymphoma Assay
- Ion Torrent[™] Oncomine[™] Childhood Cancer Research Assay

Immuno-oncology research solutions



- Ion Torrent[™] Oncomine[™] Comprehensive Assay Plus
- Ion Torrent[™] Oncomine[™] Tumor Mutation Load Assay
- Ion Torrent[™] Oncomine[™] TCR Beta-LR Assay (GX)
- Ion Torrent[™] Oncomine[™] TCR Beta-SR Assay
- Ion Torrent[™] Oncomine[™] Immune Response Research Assay
- Ion Torrent[™] Oncomine[™] TCR Pan-Clonality Assay

*The content here as key to notes above for the additions to the FFPE section provided herein may relate to products that have not been officially released and is subject to change without notice. For Research Use Only. Not for use in diagnostic procedures.

Liquid biopsy research solutions

- Ion Torrent[™] Oncomine[™] cfDNA Assays for Lung, Breast, Colon
- Ion Torrent[™] Oncomine[™] Breast cfDNA Assay v2
- Ion Torrent[™] Oncomine[™] Lung Cell-Free Total Nucleic Acid Research Assay
- Ion Torrent[™] Oncomine[™] Pan-Cancer Cell-Free Assay
- Oncomine[™] Precision Assay

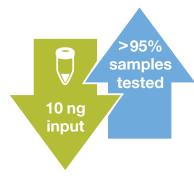
FFPE tissue testing solutions



- Oncomine[™] Precision Assay
- Ion Torrent[™] Oncomine[™] Comprehensive Assay Plus
- Ion Torrent[™] Oncomine[™] tumor-specific panels
- Ion Torrent[™] Oncomine[™] Comprehensive Assay v3
- Ion Torrent[™] Oncomine[™] Focus Assay
- Ion Torrent[™] Oncomine[™] BRCA Research Assay
- Ion Torrent[™] Oncomine[™] Comprehensive Assay Plus GX*
- Ion Torrent[™] Oncomine[™] BRCA Research Assay GX*

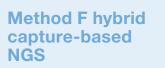
The importance of choosing the best technology for the purpose

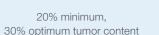
Low FFPE tissue sample input requirement is crucial



Formalin-fixed, paraffin-embedded (FFPE) tissue samples are often small, and the tumor content is limited in guantity or guality*. Oncomine assays require the least input of any commercially available assay and they have a proven record of high sequencing success**. This enables labs to get results for maximum possible number of samples and save tissue sample for possible future further analysis.

Not all NGS is the same-sample requirements can differ greatly from one test to the next NGS-based testing input requirements are typically expressed in nanograms of nucleic acid, and can differ significantly between different NGS-based tests. The figures below explain the practical implications of these different requirements in terms of tissue, tumor area, and content. Even if similar numbers of slides are required for both tests, the tumor area and percentage of tumor content required are significantly higher for method F, in order for testing to be successful.





10% minimum tumor content



25 mm² minimum sample surface area, entire block or 10 slides required



50–1,000 ng minimum input

No minimum surface area requirement; 2 slides for resection, 9 for CNB required



42.7%

Method O

NGS

amplicon-based

Samples eligible for hybrid capture-based NGS



eligible for amplicon-based NGS

According to data from a multicentric study on more than 20,000 samples, more than half of all samples would not be suitable for hybrid capture-based NGS, while 93.8% of samples including those with less than 25 mm2 surface area were successfully tested by amplicon-based NGS methods***.

Samples

* Fernandes, et al. (2021) Clinical application of next-generation sequencing of plasma cell-free DNA for genotyping untreated advanced non-small cell lung cancer Cancers 13(11) 2707. https://doi.org/10.3390/cancers13112707

** Volckmar, A.-L. et al. (2019) Combined targeted DNA and RNA sequencing of advanced NSCLC in routine molecular diagnostics: Analysis of the first 3,000 Heidelberg cases. Int. J. Cancer, 145: 649-661. https://doi.org/10.1002/ijc.32133

*** Scott, A et al. (2020) Actionable CR-based comprehensive genomic profiling (PCR-CG P): Feasibility from >20,000 tissue specimens and predicted impact on actionable biomarker identification vs. hybrid capture (H)-CG P and plasma (P)-CGP. Presented at ASCO 2020.

Translocation and fusion detection: two sides of the story

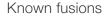
The Ion Torrent[™] Oncomine[™] assays include a multifunctional approach to detect and identify gene fusions in order to achieve a proper balance of strong performance on FFPE tissue with limited RNA input and comprehensive fusion detection.

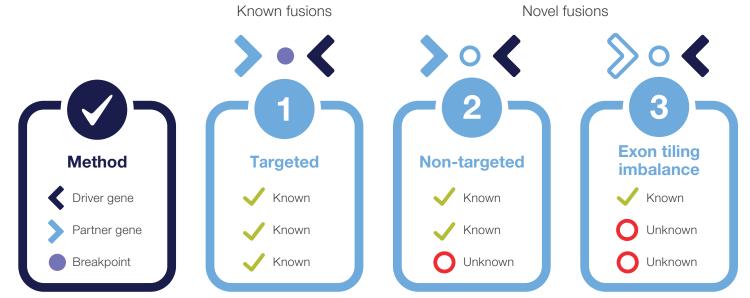


- 1. Detection of fusion in low sample inputs
- 2. Detection of fusion transcripts at low levels
- 3. Detection of novel fusions for driver genes
- 4. Many technologies emphasize No. 3 above but ignore Nos. 1 and 2. With FusionSync[™] detection, we address Nos. 1, 2, and 3.

FusionSync detection technology

This new technology for routine fusion detection in clinical research samples is designed to improve novel fusion detection and is available in Oncomine Precision Assay and Oncomine Comprehensive Assay Plus. The targeted approach employs hundreds of fusion isoforms across varying partners and breakpoints to help ensure sensitive and specific detection of known fusions. In addition, exon tiling imbalance is available for ALK, FGFR1, FGFR2, FGFR3, NTRK1, NTRK2, NTRK3, and RET fusion drivers, which indicates not only the presence of an unknown fusion but also whether it will be a functional or nonfunctional fusion based on its impact on kinase domains.



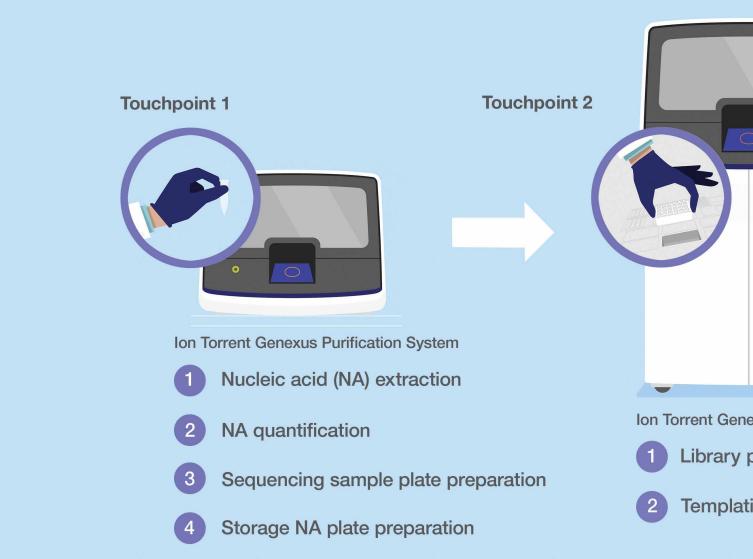


Key considerations for optimal fusion detection:

The one and only end-to-end NGS solution for any laboratory—The Genexus System



All operated by one Ion To



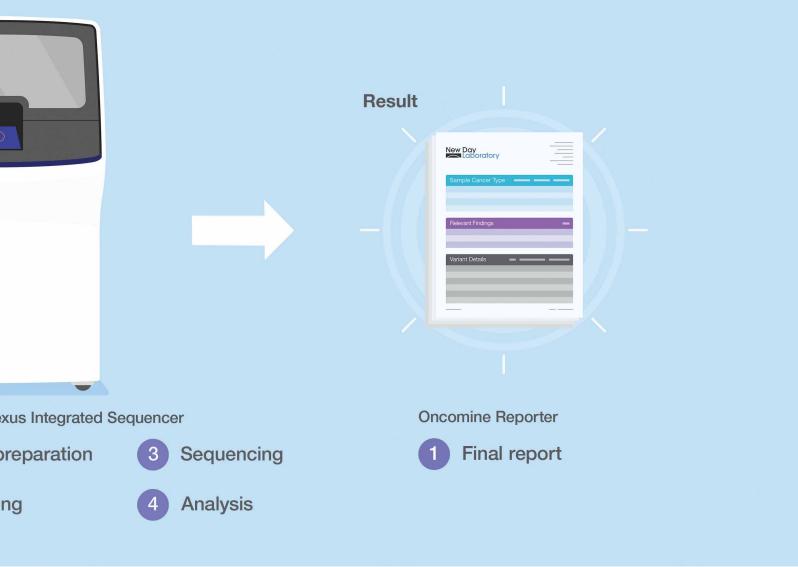
Currently including

- Oncomine Precision Assay
- Oncomine Comprehensive Assay v3
- Oncomine Myeloid Assay

Oncomine TCR Beta-LR Assay Coming Soon

- Oncomine BRCA Research Assay
- Oncomine Comprehensive Assay Plus

rrent[™] Genexus[™] Software



Two Oncomine flagship assays on the Genexus System– Key biomarkers in less than 1 day, with 10 min hands-on time

Oncomine Precision Assay

The Oncomine Precision Assay on the new Ion Torrent[™] Genexus[™] System is our next-generation solution for genomic profiling that empowers labs with an automated, hands-off workflow that takes you from FFPE tissue or plasma to report in one day.* It can be easily implemented in every lab, even without previous NGS experience, and will enable you to combine your lab's immunohistochemistry (IHC) results with an NGS profile of 50 key genes for a fast, comprehensive report.

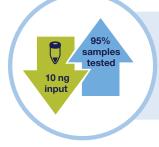


Key relevant biomarkers across 50 key genes enable you to generate a complete relevant profile

Tissue and plasma

Enabling you to generate a profile even with a lack of tissue sample



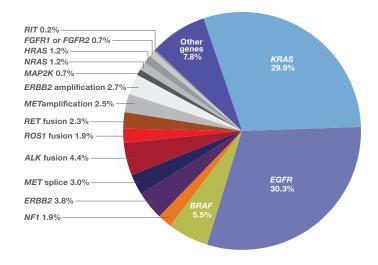


Low sample requirement

of 10 ng means many more samples successfully tested

With the Oncomine Precision Assay

you can test all key relevant biomarkers for NSCLC clinical research either from FFPE tissue or liquid biopsy sample



* Specimen-to-report workflow will be available after the Ion Torrent[™] Genexus[™] Purification System and integrated reporting capabilities are added in H2 2021.

Oncomine Myeloid Assay GX

With the Oncomine Myeloid Assay GX on the Genexus System, you can get a comprehensive myeloid mutational profile and results from a single NGS run in just one day. The highly integrated workflow lets you go from specimen to report with only 10 minutes of hands-on time and two user touchpoints. Simultaneously profile 40 DNA target genes and 29 RNA fusion driver genes with a single assay. The broad fusion panel allows you to detect over 600 unique fusion isotypes.



Rapid turnaround results in 1 day to help provide critical answers fast

Comprehensive panel of biomarkers DNA mutations and RNA fusion transcripts with one assay





Study all major myeloid disorders AML, MDS, MPN, CMML, and JMML



40 total genes 23 hotspot genes 17 full genes

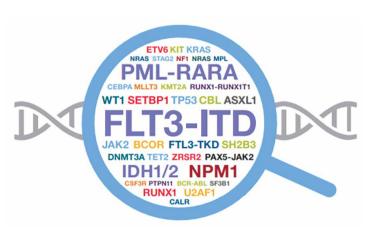


29 RNA fusion drivers >600 fusion isotypes



5 gene expression genes

5 gene expression control genes



Comprehensive genomic profiling without compromises **Oncomine Comprehensive Assay Plus**

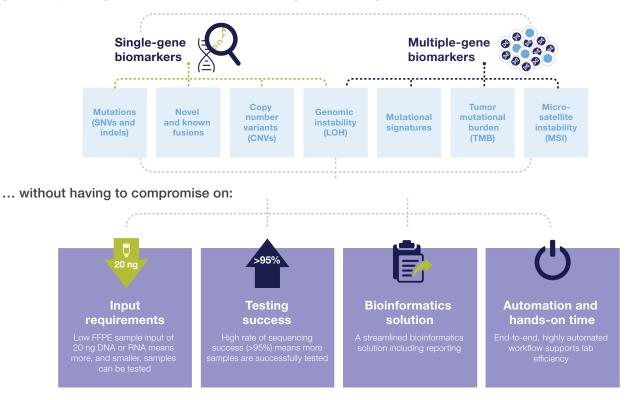
Comprehensive genomic profiling (CGP) is advancing precision oncology research through the analysis of multiple relevant biomarkers in a single NGS test*. However, the high sample input volume requirements and other technical limitations of some hybrid capture-based NGS assays, mean that >57% of clinical research samples are not even eligible for testing**. The Oncomine Comprehensive Assay Plus provides CGP without these limitations.

Existing tools are limited



Oncomine Comprehensive Assay Plus

From one sample, in one assay run, you can deliver truly comprehensive genomic profiling based on DNA and RNA analysis of >500 genes ...



* Bubendorf, L et al. (2017) Non-small cell lung carcinoma: diagnostic difficulties in small biopsies and cytological specimens. European Respiratory Review. 26(144)17007 DOI: 10.1183/16000617.0007-2017 ** Rhodes, D (2020) PCR-based comprehensive genomic profiling (PCR-CGP): feasibility from >20,000 tumor tissue specimens (TTS) and predicted impact on actionable biomarker identification versus hybrid capture (H)-CGP and plasma (P)-CGP. Journal of Clinical Oncology. 38(15) 3574.

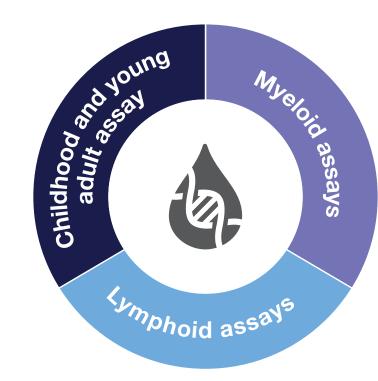
Simplified journey to answers in hemato-oncology research The tools you need to expedite your path to hemato-oncology research insights in one place

Hemato-oncological disorders are complex, heterogenous diseases with many subtypes and oncogenic drivers.

Traditionally, the molecular research of myeloid and lymphoid malignancies has involved numerous single-gene tests, multiple technologies, and laborious workflows.

In-house NGS testing enables you to consolidate and streamline your workflows, providing faster turnaroundtime (TAT), higher sensitivity, and potentially higher detection yield. Simplify and accelerate your research with our growing portfolio of hemato-oncology research assays. We make it easy for you to get everything you need in one place.

- 1 dedicated NGS partner
- 1 familiar workflow
- 1 comprehensive assay menu for hemato-oncology research





Oncomine Childhood Cancer Research Assay



Oncomine Myeloid Research Assay Oncomine Myeloid Assay GX



Oncomine IGHV Leader-J Assay Oncomine Lymphoma Assay* Oncomine TCR Beta-SR Assay

End-to-end bioinformatics solution

The Oncomine informatics workflow presents a sample-to-report solution for data analysis, from initial sequence analysis of many variants to annotation of relevant cancer drivers, and a final report by Oncomine Reporter*. This creates a simple, streamlined solution that doesn't require any specialized bioinformatics expertise.

Oncomine informatics workflow



Analysis of the sequence: Torrent Suite[™] Software streamlines sequencing runs and creates BAM files



Annotation and filter: Ion Reporter[™] Software calls variants; identifies and annotates cancer drivers



Oncomine Reporter links variants to relevant evidence in a curated knowledgebase



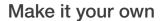
123 Street City, State USA 000000 Tel +1 000-000-0000 email@examplehealth.com www.examplehealth.com

 Case Number:
 00-123456789
 Sample Collected:

 Sample Cancer Type: Non-Small Cell Lung Cancer
 Sample Collected:
 Sample Collected:

| Gene | Finding | Gene | Finding |
|-------|-----------------------|-------|--------------|
| ALK | Not detected | NRAS | Not detected |
| BRAF | Not detected | NTRK1 | Not detected |
| EGFR | EGFR exon 19 deletion | NTRK2 | Not detected |
| ERBB2 | Not detected | NTRK3 | Not detected |
| KRAS | Not detected | RET | Not detected |
| MET | Not detected | ROS1 | Not detected |

| Relevant Biomarkers | | | | | | | |
|---------------------|-----------------------|-------------|--|--|--|--|--|
| Tier | Genomic Alteration | Annotations | | | | | |
| IA | EGFR exon 19 deletion | | | | | | |



The first page of the report can be customized to show the relevant biomarkers and variant details sections.

Templates can be customized with your logo, with custom text, and in one of eleven languages.

The report builder allows sections to be included and excluded as needed, enabling easy customization.

| Sample data | | | |
|-------------|--------------|--|--|
| Gene | Finding | | |
| BRAF | BRAF V600E | | |
| KIT | Not detected | | |
| NTRK1 | Not detected | | |
| NTRK2 | Not detected | | |
| NTRK3 | Not detected | | |

Oncomine Reporter software

- Oncomine Reporter software produces clear and concise reports
- Includes relevant biomarker data with relevant evidence
- Optimized for Oncomine assays across applications including FFPE tissue testing, liquid biopsy, TMB, and hemato-oncology, and cytogenetics



* Sakai K, et al. (2019) A comparative study of curated contents by knowledge-based curation system in cancer clinical sequencing. *Sci Rep.* 9(1):11340. Published 2019 Aug 5. doi:10.1038/s41598-019-47673-9

Rely on the data

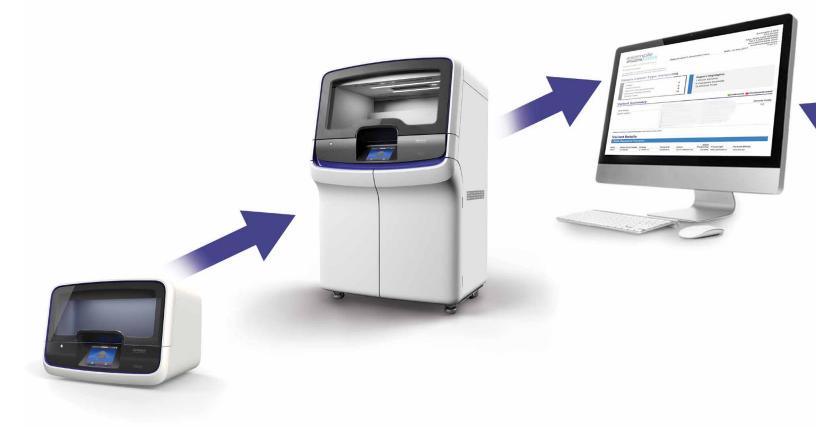
Monthly updated data

 Ion Torrent[™] Oncomine[™] Knowledgebase data, based on which filtration and annotation of the sequencing data are done, are updated monthly.

Meticulous curation and QC

• After data are collected from various global data sources, a team of experienced curation scientists manually reviews all candidate evidence. Two independent reviewers examine each piece of candidate evidence for context and standardization. The process has QC steps built in at various stages.

Also, when it comes to the NGS workflow, one size does not fit all



Genexus System

The new Genexus System is the first turnkey NGS solution that automates the specimen-to-report workflow and delivers results in a single day with just two user touchpoints.* The simplicity and practicality of the Genexus System means every lab can bring NGS in-house, regardless of the level of NGS expertise.

The Genexus System is compatible with the Oncomine Precision Assay, Oncomine Comprehensive Assay v3, Oncomine TCR Beta-LR Assay, and the Oncomine Myeloid Research Assay.

Get results for your routine samples with one-day TAT* and minimum operational resources with the Genexus System and the Oncomine Precision Assay



Ion GeneStudio S5 System

The Ion GeneStudio[™] S5 System combines with the Ion Chef[™] Instrument for library preparation and templating, providing a highly automated NGS workflow with scalability. A single Ion GeneStudio S5 instrument can process several different chip sizes, allowing the instrument to handle multiple levels of throughput.

The Ion GeneStudio S5 System is compatible with all assays from the Oncomine portfolio except the Oncomine Precision Assay.

Utilize the capacity of the Ion GeneStudio S5 instrument for projects requiring the full breadth of the Oncomine portfolio



Enable your oncology research with trusted Oncomine Solutions

Explore Oncomine Solutions at **oncomine.com**

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