

# Connections bring meaning to cancer testing

Identify insights faster while  
scaling NGS operations  
for somatic oncology  
applications







# Move precision medicine forward

## Streamlined

Accelerate time to report by harnessing the power of automation, user-defined workflow configuration, and > 45 knowledge sources

## Integrated

Remove unnecessary touchpoints and manual data movement by connecting data analysis with upstream sequencing to simplify and secure the comprehensive workflow

## Powered for growth

Scale the volume and throughput of your NGS workflows without increasing headcount to keep pace with evolving science, technology, and demand

## Connected Insights harnesses > 45 knowledge sources to bring insights to diverse applications

### Comprehensive knowledge in one place

Apply the power of API integrations to connect LIMS, storage, pipelines, and third-party knowledge bases; integrate variant classifications, relevant drug labels, and pertinent clinical trials within a single view

### Evidence-based clinical content

Leverage comprehensive solid and hematological cancer content, somatic gene variant annotations, and related content powered by The Jackson Laboratory Clinical Knowledgebase (JAX-CKB™), a trusted knowledge base of over 100,000 users worldwide

### Flexible regionality

Incorporate relevant lab- and region-specific practices within user-defined workflow; support customized variant interpretation and reporting based on regional-tiering guidelines

### Regular updates

Access up-to-date, relevant variant annotations, insights, guidelines, and more, with Connected Insights-Research and integral regular, automated content updates (as frequently as monthly for some knowledge sources)

# Unlock meaning for a wide range of applications within a single, customizable platform



## Solid tumor testing (tissue)

Assess comprehensive gene panels covering multiple variant types, including TMB, MSI, and GIS to assess HRD



## Liquid biopsy

Detect and analyze cancer variants with high analytical sensitivity and specificity using low levels of ctDNA found in the bloodstream



## Hematological cancer

Streamline insight generation for myeloid leukemia, lymphoma, and other hematologic malignancies



## Hereditary disease\*

Uncover insights by interpreting genome-wide disease biomarker signatures (STR, paralog) with increasing relevance to precision medicine

\*Capabilities expected in future roadmap.

## Powerful features to streamline integration and adoption of a single-vendor workflow

### Enterprise-level security and privacy standards

Protect the privacy of your genomic data with industry-leading global and local security standards

### User-defined SOPs and advanced filters

Implement SOPs on the platform by creating sets of predefined custom filters or using the advanced filtering system

### Streamlined workflow and collaboration

Facilitate teamwork; save time and remove manual data movement steps with autoingestion of variant data and autolaunch of Connected Insights

### Broad portfolio of tests and variant types

Evolve your analysis to comprehensive panels, exomes, or genomes; analyze across a range of DNA and RNA variant and biomarker types, including TMB, MSI, and GIS to assess HRD

### Lab-specific curation

Maintain a private knowledge base of your organization's curated data, including information about past variant interpretations and reporting

### Automated custom reporting

Customize, edit, and automatically populate reports with minimal manual interaction; generate reports in 18+ different languages as needed

Available in select countries.

# Integrate and streamline your workflows from library prep, sequencing, and data analysis

White paper



## [GenomeWeb KOL white paper](#)

Learn the thoughts of key opinion leaders on the current challenges and promise of NGS interpretation and reporting in clinical oncology

Data sheet



## [Connected Insights data sheet](#)

Read how Connected Insights streamlines, integrates, and powers laboratories for scale and growth

Video



## [Connected Insights animated video](#)

Understand how Connected Insights can connect various knowledge sources to streamline operations for powerful insights

Learn more at [illumina.com/connected-insights](https://www.illumina.com/connected-insights)

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## Abbreviations

API application programming interface

CNV copy number variant

ctDNA circulating tumor DNA

GIS genomic instability score

HRD homologous recombination deficiency

Indels insertion/deletion

KOL key opinion leader

LIMS laboratory information management system

MSI microsatellite instability

NGS next-generation sequencing

SNV single nucleotide variant

SOP standard operating procedure

SV structural variant

TMB tumor mutational burden

WES whole-exome sequencing

WGS whole-genome sequencing