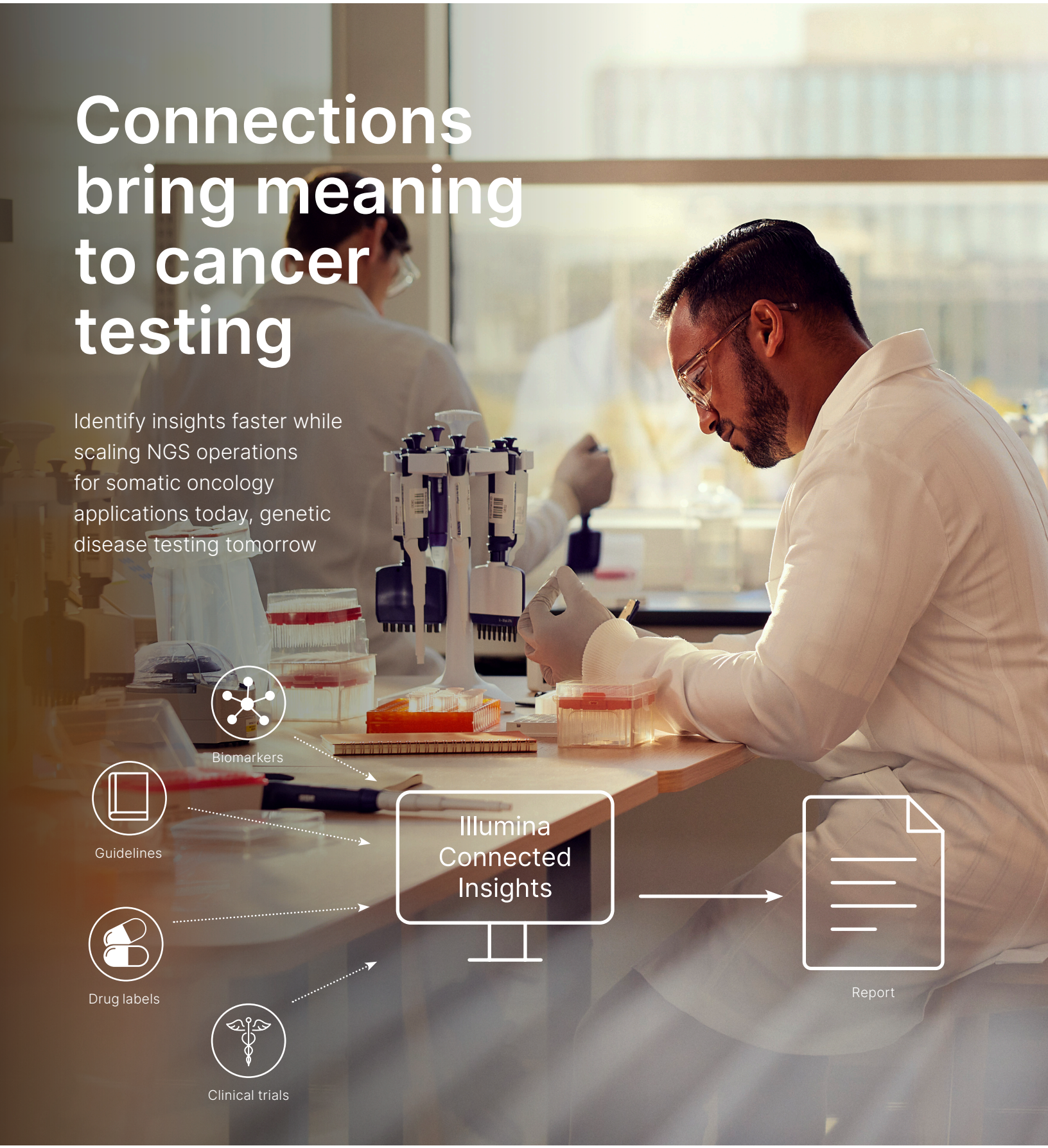
illumina® | Connected Insights

Connections bring meaning to cancer testing

Identify insights faster while scaling NGS operations for somatic oncology applications today, genetic disease testing tomorrow



Available in select countries.

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 touch points and manual data movement by connecting to sequencing systems to simplify and secure assay workflows

Powered for growth

Scale the volume and throughput of your NGS workflows without increasing headcount to keep place 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

Flexible regionality

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

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 (TM)), a trusted knowledgebase of over 100,000 clinicians worldwide

Regular updates

Access the most up-to-date, relevant content, with Connected Insights updates made on a monthly basis

Available in select countries.

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

Solid tumor testing (tissue)

Assess comprehensive gene panels across multiple variant types, including TMB, MSI, and more

Liquid biopsy

Detect and analyze cancer with high sensitivity and specificity using low levels of ctDNA 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

Broad portfolio of tests and variant types

Evolve with confidence, broadening 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

User-defined SOPs and advanced filters

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

Lab-specific curation

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

Streamlined workflow and collaboration

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

Automated custom reporting

Customize, edit, and automatically populate reports; minimal manual interaction

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

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

Learn more at **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
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

Available in select countries.

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