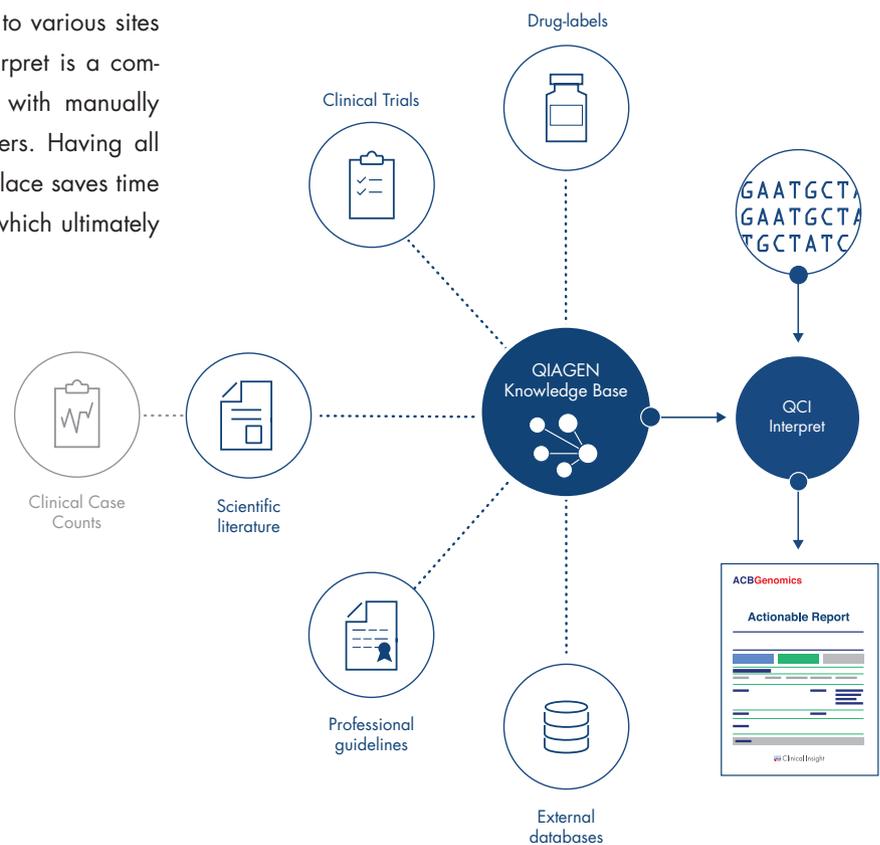




Excellence in content, & growing in data

QCI™ Interpret for Hereditary Cancer

In hereditary cancers, clear insight into genomic risk factors helps navigate better patient care. With QCI Interpret for Hereditary Cancer you no longer need to go to various sites and sources to score NGS variants. QCI Interpret is a comprehensive clinical decision support solution with manually curated scientific evidence for germline cancers. Having all the information you need aggregated in one place saves time and improves interpretation of NGS variants which ultimately helps increase clinical decision accuracy.



Growth powered by scalable insights

QCI Interpret replaces your complex and manual clinical NGS variant research with a scalable interpretation workflow alternative that helps grow your indication menu and volume.

Key Benefits:

- Increase operational efficiency associated with scoring NGS variants by reducing time, costs, and complexity to classify variants
- Focus interpretation time on high-risk variants that better inform genetic-profile specific patient care plans
- Build your own experience-based database with variants assessed, increasing speed and accuracy with subsequent interpretations

Reliable curation with superior depth

QCI Interpret is an efficiency tool that helps you identify pathogenic variants fast with results you can trust, all in one secure web application.

Key Features:

- Computed classifications that are based on rules published in ACMG professional medical guidelines for sequence variant interpretation
- Manually curated clinical case counts with clickable hyperlinks direct to source materials for easy retrieval and review
- Simple, clear, and easy-to-use configurable reports that also include bibliographic reference citations

Speak to us about QCI Interpret or request a demo

Find out if QCI interpret for Hereditary Cancer fits your NGS interpretation and reporting needs. Speak to a representative from QIAGEN Bioinformatics to take a closer look. Request your free, no obligation, demonstration of QCI™ Interpret for Hereditary Cancer today.

“Clinical labs rolling out NGS-based tests are confronted with two key challenges: the complexity of turning molecular profiling information into precise medical recommendations, and the time and effort it takes to generate actionable reports. QIAGEN Clinical Insight provides a rich and detailed, yet very clear and concise, report that suggests management and treatment options based on the patient’s gene variations that profile their disease and outline causal links. It is this kind of interpretation that gives clinical value to the data, and what enables the actual insights into a patient’s specific disease and treatment options.

Madhuri Hegde, PhD, FACMG

Professor of Human Genetics at the Emory University School of Medicine and Executive Director of the Emory Genetics Laboratory

Contact us now at: QCI-Info@qiagen.com

Learn more online at: qiagenbioinformatics.com/QCI

QCI Interpret is an evidence-based decision support software intended as an aid in the interpretation of variants observed in genomic next-generation sequencing data. The software evaluates genomic variants in the context of published biomedical literature, professional association guidelines, publicly available databases, annotations, drug labels, and clinical-trials. Based on this evaluation, the software proposes a classification and bibliographic references to aid in the interpretation of observed variants. The software is NOT intended as a primary diagnostic tool by physicians or to be used as a substitute for professional healthcare advice. Each laboratory is responsible for ensuring compliance with applicable international, national, and local clinical laboratory regulations and other specific accreditations requirements.

