

Instructions for Testing

Important Note:

PanCanSeek® serves as an auxiliary diagnostic tool and should not be used as the sole basis for clinical decisions. All results require professional interpretation by professional clinicians.

Who should take this test?

- The PanCanSeek® test is indicated for patients with suspected or diagnosed hematologic malignancies

Understanding your results:

■ The results consist of the following parts: structural variations (SVs), copy number aberrations (CNAs) and copy number neutral loss of heterozygosity (CN-LOHs), and single nucleotide variations (SNVs, InDels). These abnormalities are classified into three levels:

Tier.1 variation: Variants with clear clinical significance (Diagnostic, prognostic, and/or therapeutic).

Tier.2 variation: Variants with potential clinical significance (Diagnostic, prognostic, and/or therapeutic).

Tier.3 variation: Variants with unclear clinical significance.

Important reminders:

- Blood sample collection, transportation, storage, and non-standard processing may impact the performance of PanCanSeek®.
- The content that can be detected by PanCanSeek® includes SVs, whole-genome aneuploidy abnormalities, as well as CNAs (≥ 1 Mb) and CN-LOHs (≥ 5 Mb), and SNVs.
- PanCanSeek® cannot detect chromosomal duplications or deletions up to 1Mb.
- Never rely on this test alone. The result must be confirmed by professional clinicians following clinical standardized procedures.

Questions?

Contact your doctor or our support team: info@seekincancer.com