



Your Choice for NGS Kits



Watchmaker DNA Library Prep Kit with Fragmentation

Highly efficient conversion of DNA of variable quality into Illumina®-compatible sequencing-ready libraries for highly sensitive clinical and translational applications

Equinox Library Amplification Kit

Ultimate library amplification kit specifically address the stringent demands of applications



KEY BENEFITS: Watchmaker DNA Library Prep Kit with Fragmentation

- Verv high efficiency
- Compatible with ultra-low and low input DNA (1 ng 500 ng) and FFPE samples
- Highly tunable and consistent fragmentation for application-specific needs
- Low bias an ultra-high-fidelity amplification
- Equinox Polymerase (included in the kit) reduces misincorporation events by up to 40% to improve overall assay sensitivity
- Support the coverage for >75% of all read families (including high GC regions) within 3X of the mean family depth
- High sensitivity and accuracy with uniform sequence coverage
- Improves assay accuracy and sensitivity for variant and rare mutation detection
- Get up to a 90% reduction in sequence artifacts (e.g., false chimeric reads and false SNVs)
- Uniform sequence coverage even for complex genomes with extreme (15 85%) GC content



APPLICATIONS

- Somatic mutation calling and other lowfrequency variant detection NGS assays, including those utilizing challenging samples such as FFPE and cfDNA
- Human and microbial whole-genome sequencing (WGS)
- Whole exome sequencing (WES)
- · Single-cell analysis
- Metagenomic analysis
- Bulk RNA sequencing (using cDNA as input)
- · Viral genome sequencing
- · Inherited disease sequencing
- · Amplicon sequencing
- ChIP-seq, ATAC-seq, and associated epigenetic applications †
- Suitable for Illumina and non-Illumina sample preparation workflows

Note:

† Workflow can incorporate Equinox Library Amplification Kit

Incorporate the stand-alone Equinox Library Amplification Kit for template amplification and still experience the robust performance of Watchmaker solutions

MORE BENEFITS: Equinox Library Amplification Kit (with Equinox Polymerase)

- Ultra-high-fidelity library amplification from a wide range of template amounts (0.1 pg 500 ng) up to 1 kb in length
- Effective hot start formulation inhibits both polymerase and 3' → 5' exonuclease activities, for low-input applications and automated workflows
- · Compatibility with paramagnetic beads ensures robust performance in hybridization capture workflows