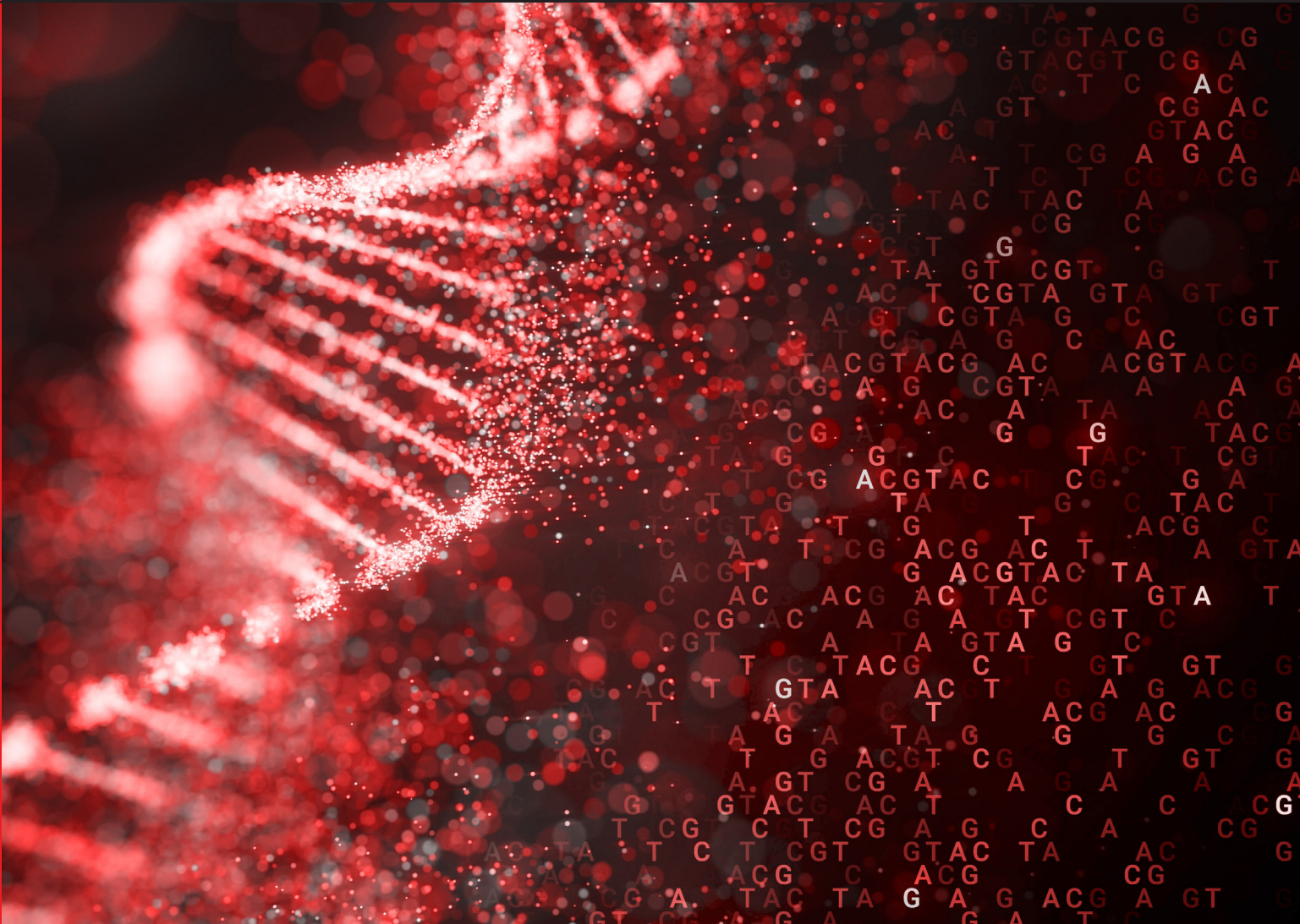


The Foundation for Your Next Generation Sequencing

NGS Library Preparation



analytikjena

An Endress+Hauser Company

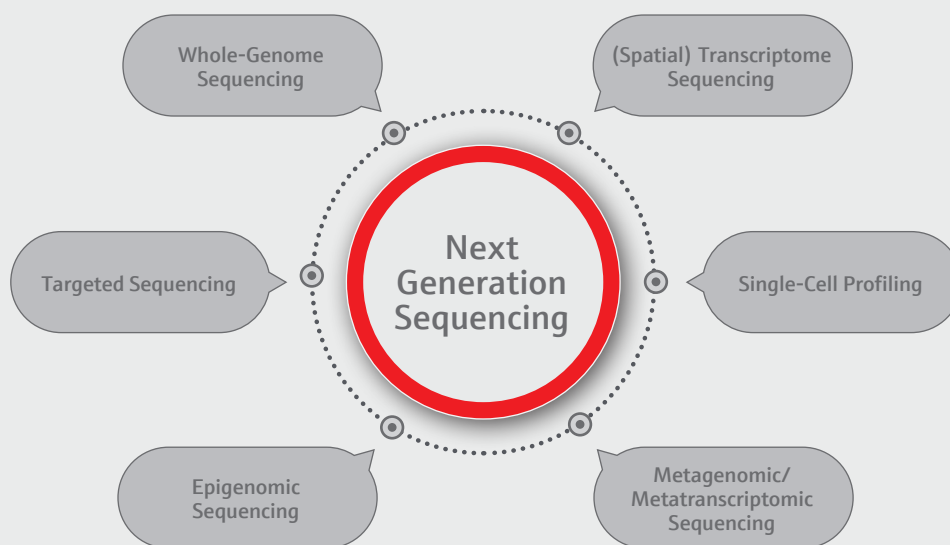
Trust in Your NGS Library Preparation

Next Generation Sequencing (NGS) technologies have revolutionized the field of genomics by performing massive parallel sequencing of millions of DNA fragments.

This powerful technology has paved the way for breakthroughs in several areas of biology, including genomics, personalized medicine and evolutionary biology. NGS enables to study whole genomes or transcriptomes, identify sequence variants or even single cells and to uncover new insights into the structure, function, and variation of genomes.

A critical step for successful NGS results is the sequencing library preparation, which generates a pool of sequencing

fragments from RNA or DNA samples in a format suitable for sequencing. The quality of the library preparation directly impacts the downstream analysis and interpretation of NGS data. By carefully constructing sequencing libraries, researchers can maximize the yield of usable sequencing data, minimize technical biases or artifacts that can occur during the sequencing process and increase sequencing coverage for vulnerable sample material.



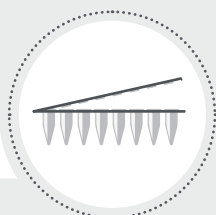
NGS Library Preparation: From Sample to Library

DNA library preparation involves a series of steps with enzymatic incubations, including DNA/RNA fragmentation, cDNA synthesis, adapter ligation, and amplification. In addition, quality control measures such as fragment size assessment and library quantification are integral parts of the workflow.

With Analytik Jena instruments, you can ensure that your DNA libraries fully meet the requirements of your chosen sequencing technology.

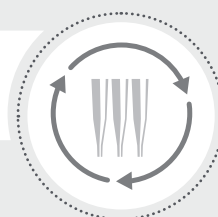


Thermal cyclers



PCR consumables

Automated solutions: Liquid handling systems, robots, automated thermal cyclers



Enhance Your Library Preparation Process

Why should you rely on Analytik Jena instruments?

Minimized bias made possible by:

- Reduced risk of non-specific amplifications
- No temperature over- and undershoots

Optimal yield made possible by:

- Ensured enzyme activity
- Excluded cross contamination

Reproducibility made possible by:

- Temperature homogeneity across the whole block



Biometra TAdvanced



Biometra TRIO 48

Choose your PCR thermal cycler according to your needs

- Biometra TAdvanced for premium performance
- Biometra TRIO for varying applications and throughputs
- Biometra TRobot II: space-saving & robust for optimal automated PCR

... and benefit from

- Consistently high library quality
- Reproducible results across the whole block
- No temperature over- and undershoots to ensure enzyme activity and specific products
- No cross-contamination by reproducible lid pressure
- Compatibility with all common PCR consumables
- High volume capacity with flexible 96 and 384 block format

Leverage from PCR expertise for your automated workflow

- With excellent performance for reliable results
- Accurate temperature control
- Space-saving and robust design
- With advanced software designed for automation

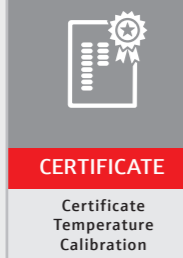


Biometra TRobot II

Intuitive software & reliable service

Seamless thermal cycler management & confidence in performance

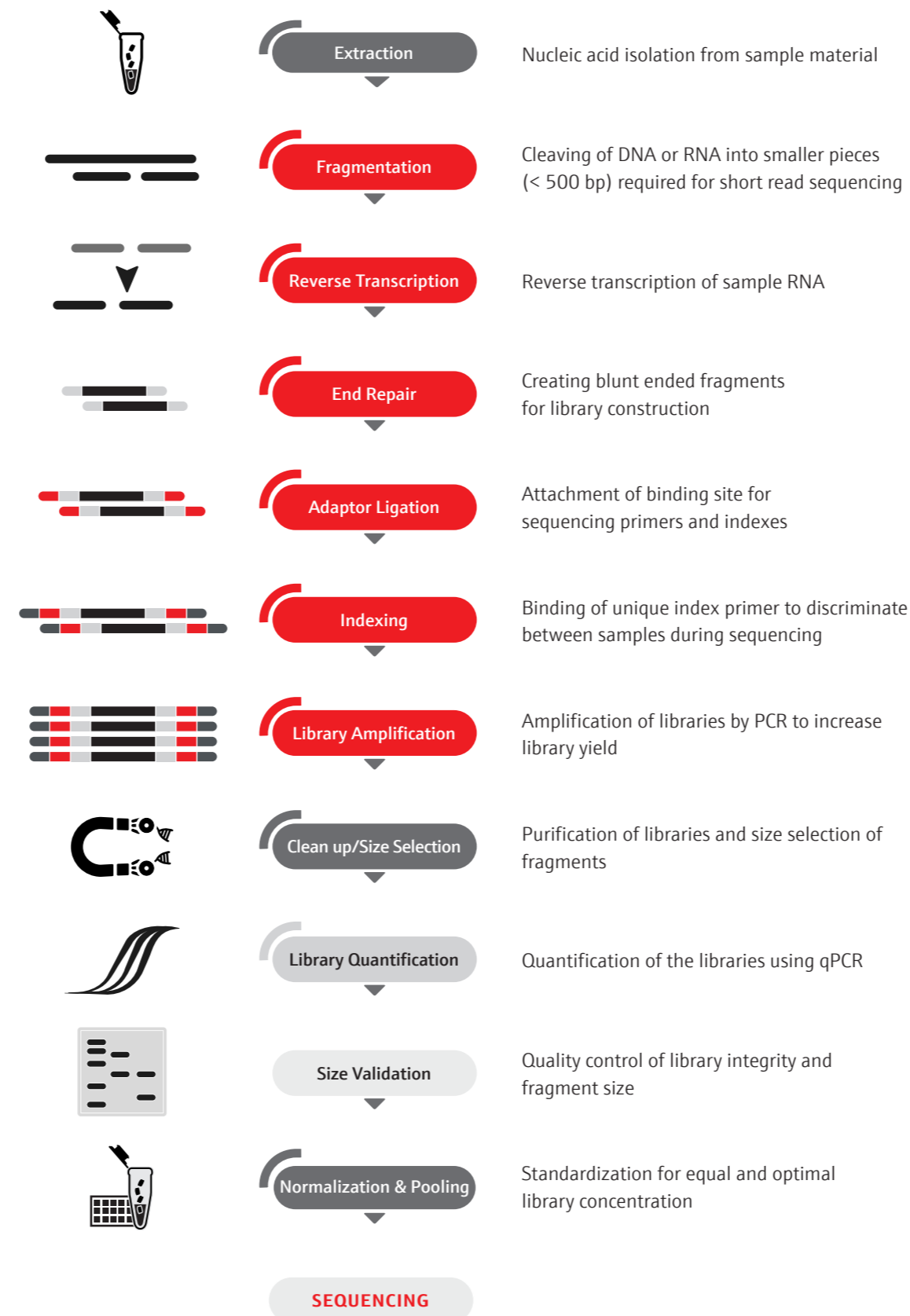
- Comfortable remote control and online monitoring with the intuitive Biometra TSuite software
- Verified thermal cycler performance with the Biometra TMS calibration service, including on-site temperature adjustments if required



CERTIFICATE

Certificate
Temperature
Calibration

Typical Steps of NGS Library Preparation



Want to Automate Your Workflow?

Elevate your NGS library preparation with automation for efficient, precise pipetting to ensure a seamless, error-free workflow.

Rely on proven liquid handling technology

- Save valuable space with the modular benchtop pipetting robot CyBio FeliX
- Enhance reproducibility and throughput
- Expand your personal automation solution any time by connecting automated thermal cyclers and CyBio FeliX via the versatile benchtop robot CyBio Carry



CyBio Carry



CyBio FeliX

Pipetting robot CyBio FeliX seamlessly manages bead-based procedures like **extraction, size selection, and clean-up**, while also excelling in normalization, pooling, and **(q)PCR setups**.

Ensure Library Quality

The determination of library concentrations, fragment size and integrity are essential QC parameters for any NGS sequencing library to ensure precise and adequate input for the sequencing process.

Quantify libraries easy and accurate with our high-precision real-time thermal cyclers

- Highly sensitive detection across a broad range of library concentrations
- Flexible throughput of up to 384 reactions
- Compatibility with all common PCR consumables
- Enhance throughput with qTOWER³ auto: Truly fully automated design for integration into robotic systems



qTOWER³ auto or
qTOWER³ 84 auto



NEW

qTOWER iris

Learn More

Discover your Next Generation Sequencing Library Preparation solutions: From manual workflows to modular automation, achieving excellence for all throughputs.

We recommend the following products from our portfolio.

Biometra TAdvanced 96 S



Order number: 846-x*-070-251

qTOWERiris



Order number: 844-00853-x*

Biometra TRIO 48



Order number: 846-x*-070-723

qTOWERiris 384



Order number: 844-00858-x*

Biometra TRobot II 96 SG



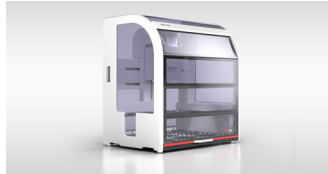
Order number: 846-x*-070-902

qTOWER³ auto



Order number: 844-00604-x*

CyBio FeliX Basic Unit with Enclosure



Order number: OL5015-24-100

* x = 2 for 230 V, x = 4 for 115 V, x = 5 for 100 V

Headquarters

Analytik Jena GmbH+Co. KG
Konrad-Zuse-Str. 1
07745 Jena · Germany

Phone +49 3641 77 70
Fax +49 3641 77 9279
info@analytik-jena.com
www.analytik-jena.com

Pictures: Analytik Jena GmbH+Co. KG
Subjects to changes in design and scope of delivery as well as further technical development.

Version 2.0 - en - 12/2025
886.53001-2-B
© Analytik Jena GmbH+Co. KG