

KAPA EvoPlus Boost Kit

Boost your performance, advance your research

Enhance your NGS sample prep workflow and increase sequencing performance with **KAPA EvoPlus Boost Kits**. This advanced enzymatic DNA library prep solution rivals mechanical shearing methods by delivering exceptional fragmentation, higher conversion efficiency, and fewer sequencing artifacts*. Now with an optimized amplification mix, the KAPA EvoPlus Boost Kits deliver unparalleled performance and confidence for your WGS, WES, and targeted sequencing applications.

- **Increase efficiency** and convenience with **all-in-one enzyme & buffer ReadyMixes**—available in either **automation-friendly** reagent **plates** or tubes
- **Recover more information** with enhanced library conversion by the engineered KAPA EvoT4 DNA Ligase
- **Maximize sequencing efficiency** by ensuring high uniformity and fewer dropouts in difficult regions with an optimized low-bias PCR mix (KAPA EvoAmp ReadyMix)
- Skip mechanical fragmentation – **without the bias** often associated with enzymatic fragmentation



2 hrs
Workflow Time

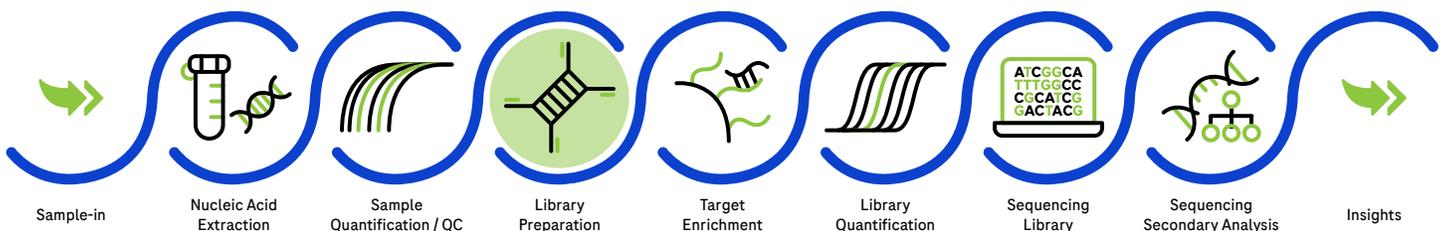


30 min
Hands-on Time



0.1-500 ng
Input Amount

Constantly evolving, efficient, and complete solutions



*Compared to existing KAPA DNA library prep chemistries.

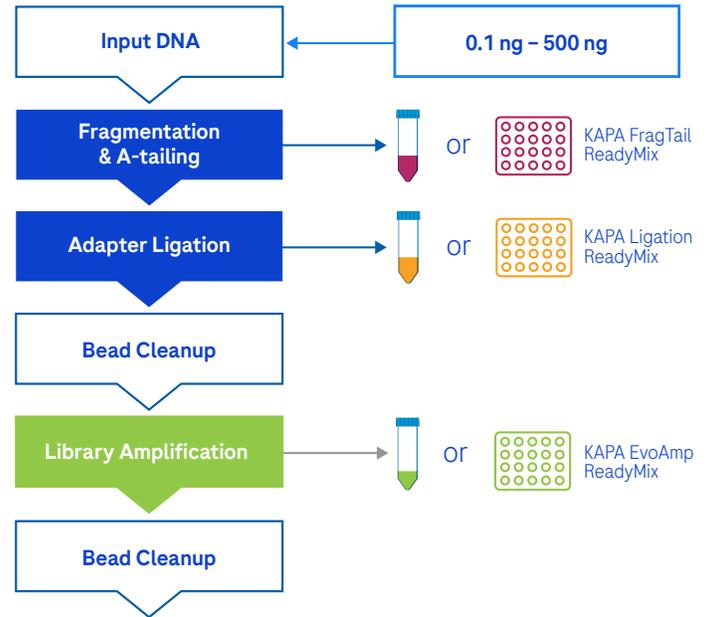
Increase efficiency and convenience

- Simplify library prep and reduce the risk of human error with a **simplified workflow**, including one-step Fragmentation & A-tailing
- Decrease hands-on-time and the number of reagents with **ReadyMix formulations** available in tubes or automation-friendly plates
- Produce sequencing-ready or target-capture-ready libraries in **2 hr**

Figure 1: KAPA EvoPlus Boost Kits are ready-to-use; no master mix preparation is required.

NOTES: KAPA full-length UDI adapters are only available in 96-well plates. KAPA Universal Adapters, which are used in combination with KAPA UDI Primer Mixes and require amplification, are available in plates or tubes. KAPA HyperPure Beads, UDI Adapter Kits & KAPA Library Amplification Primer Mix (10X) are sold separately.

KAPA EvoPlus Boost Kit



Exceptional coverage uniformity, with high fidelity and minimal GC bias

- Increase **sequencing efficiency** by enhancing **coverage uniformity and accuracy**
- Deliver high-quality **sequencing results** with **low error rates** and **minimal artifact** generation for reliable outputs

A. Coverage uniformity

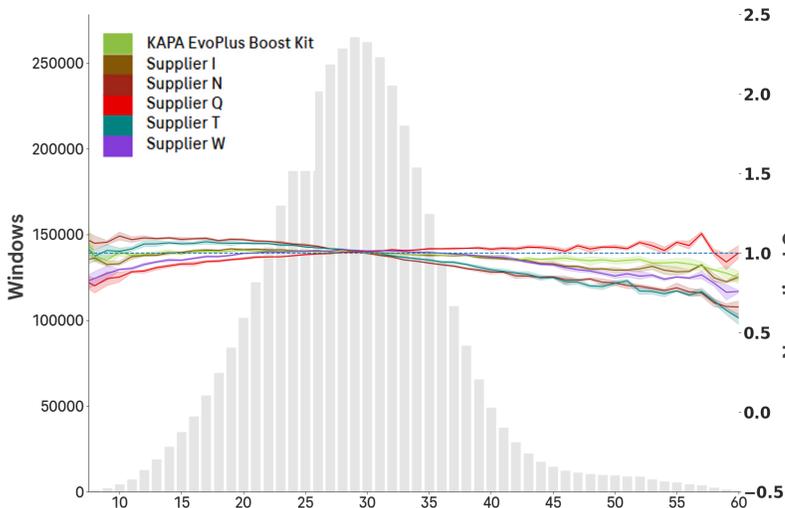
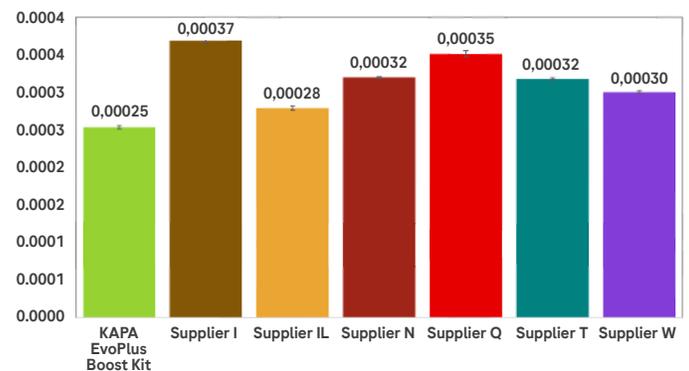
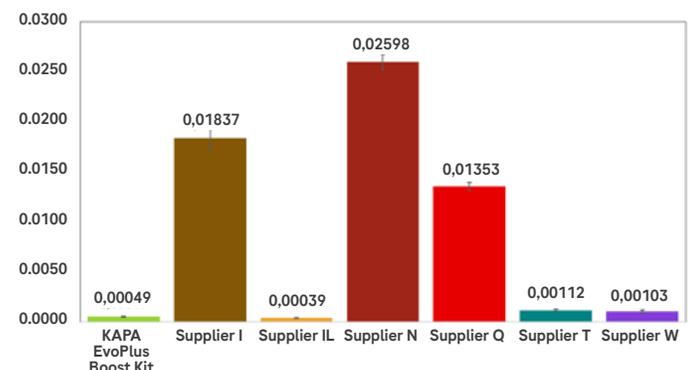


Figure 2: KAPA EvoPlus Boost Kits deliver low-bias amplification, exceptional accuracy, and minimal artifacts for superior sequencing quality. Whole genome sequencing libraries were prepared using 0.1 ng of DNA (*C. difficile*) with the KAPA EvoPlus Boost Kits or Supplier I, Supplier IL, Supplier N, Supplier Q, Supplier T and Supplier W, following each supplier's instructions for use. (A) Lower amplification bias with the KAPA EvoPlus Boost Kits, resulting in uniform coverage across the *C. difficile* genome. Best balance of highest sequencing economy and result confidence observed with the KAPA EvoPlus Boost Kits by (B) Lowest error rate and (C) ultra-low artifact rate compared to suppliers.

B. Error rate



C. Artifact rate



Skip mechanical fragmentation – without the bias

- Achieve **higher sequencing efficiency** with exceptional specificity and duplex molecule recovery
- **Minimize sequencing artifacts** and **boost result confidence** compared to mechanical fragmentation methods

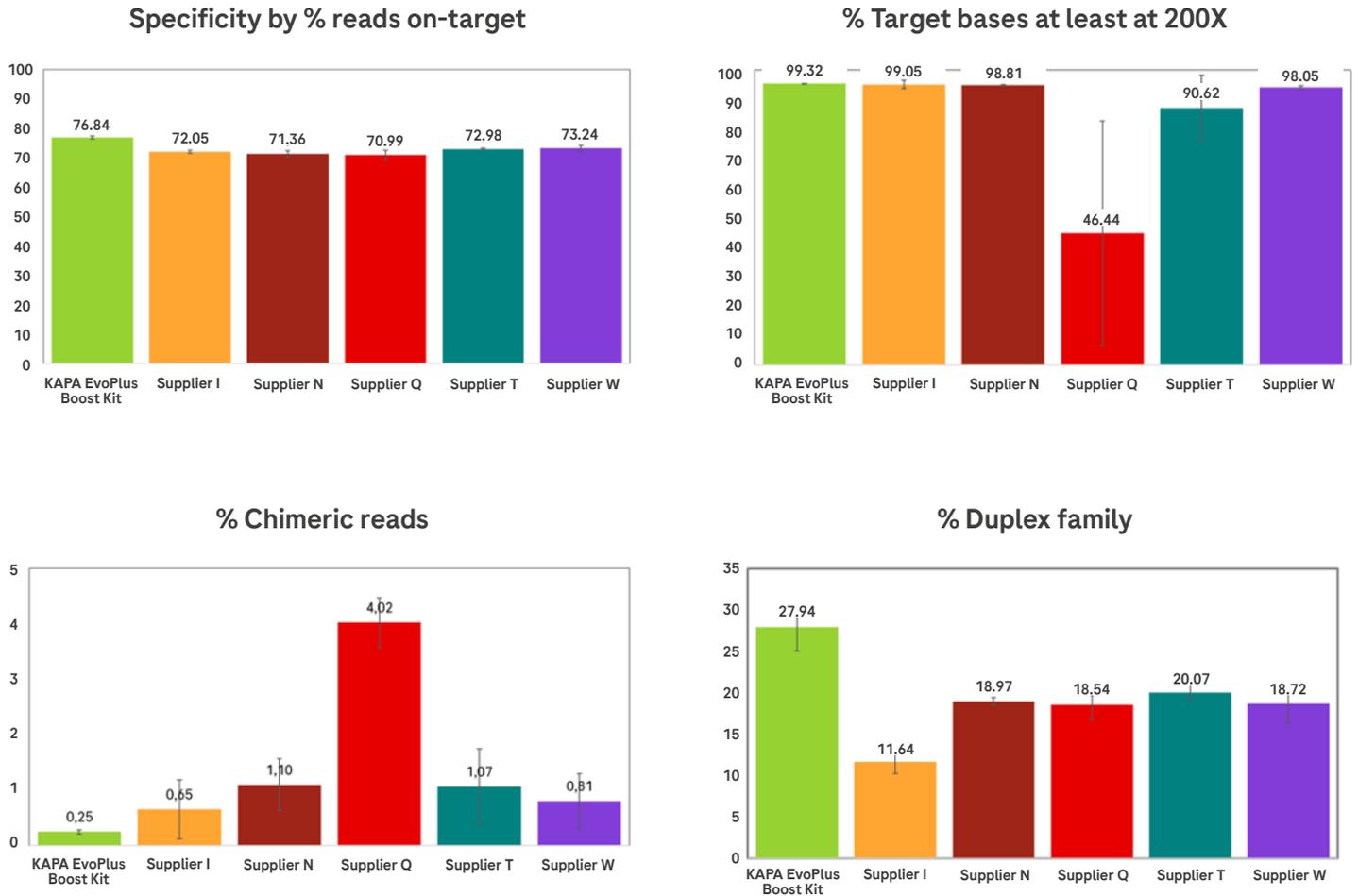


Figure 3. Improved sequencing performance in target capture workflow utilizing FFPET DNA. 50 ng of low quality FFPET DNA was used to prepare triplicate libraries with the KAPA EvoPlus Boost Kit and mechanical fragmentation kits of Supplier I, Supplier N, Supplier Q, Supplier T and Supplier W, following each supplier's instructions for use. Libraries were enriched with the KAPA HyperCap Oncology Panel (214 Kb), following the KAPA HyperCap FFPET Evolved workflow instructions¹. (A) The KAPA EvoPlus Boost Kit had one of the highest percentage of reads on-target compared to mechanical fragmentation suppliers and (B) very high percentage of target bases covered by at least 200X compared to mechanical fragmentation suppliers, thereby showcasing the optimal utilization of sequencing throughput. (C) The KAPA EvoPlus Boost Kit had the lowest percentage of Chimeras present and (D) the highest duplex recovery, resulting in higher data confidence² compared to mechanical fragmentation suppliers (Supplier I, Supplier N, Supplier Q, Supplier T and Supplier W).

1. KAPA EvoPlus Boost Kit data generated in 2025, separate from other supplier data generated in 2024.

2. Chen, *et al.* (2024). Characterization and mitigation of artifacts derived from NGS library preparation due to structure-specific sequences in the human genome. *BMC Genomics* 25:227.

Ordering Information

Roche Cat. No	Description	Kit size	Format
10613696001	KAPA EvoPlus Boost (24rxn)*	24 reactions	Tube
10613700001	KAPA EvoPlus Boost (96rxn)*	96 reactions	Tube/Bottle
10613718001	KAPA EvoPlus Boost (384rxn)*	384 reactions	Bottle
10613726001	KAPA EvoPlus Boost, plated format (96rxn)*	96 reactions	Plate
07958994001	KAPA Library Amplification Primer Mix (250rxn)	250 reactions	Tube
09420410001	KAPA Library Amplification Primer Mix (384 rxn)	384 reactions	Bottle
09420479001	KAPA Library Amplification Primer Mix 96-well plate (96rxn)	96 reactions	Plate

* KAPA Library Amplification Primer Mix (10X) not included, sold separately

For additional pack sizes, please contact your local Roche representative.



Published by:

Roche Sequencing Solutions, Inc.
4300 Hacienda Drive
Pleasanton, CA 94588

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