

Quick Guide for the Adept[™] Rapid PCR-Plus Protocol

Element Adept Library Compatibility Workflow

Introduction

This quick guide provides concise instructions for the Adept PCR-Plus Protocol, which uses the Adept Rapid PCR-Plus Kit. For comprehensive information and detailed instructions, see the *Element Adept Library Compatibility Workflow User Guide for the Rapid PCR-Plus Protocol (MA-00040)*.

Amplify Library

- 1. Prepare 0.05–0.5 pmol linear library in 0.1 mM EDTA and 10 mM Tris, pH 8.0.
- 2. Combine the following reagents, allowing 10–15% overage.

Reagent	Volume per Reaction (μl)
Adept PCR+ Primer Mix	5
Adept PCR+ Master Mix	25

- 3. Add 20 µl library to a new PCR plate.
- 4. Add 30 μl master mix.
- 5. Run the following program:

Temperature	Time	Number of Cycles
Volume set to 50 μl		
Lid set to 105°C		
98°C	1.5 minutes	1
98°C	30 seconds	5
60°C	30 seconds	
72°C	30 seconds	
72°C	1 minute	1
4°C	Hold	1

- Clean Up Library
- 1. Add 50 μl beads (1x).
- 2. Mix beads and library:
 - » For a plate, shake at 1500–1800 rpm for 2 minutes.
 - » For tubes, pipette 10 times.

- 3. Incubate at room temperature for 5 minutes.
- 4. Place on the magnet and wait until supernatant clears. *Keep on the magnet.*
- 5. Remove the entire volume of supernatant.
- 6. Wash the content of each well or tube:
 - a. Add 200 µl 80% ethanol and incubate 30–60 seconds.
 - b. Remove and discard ethanol.
 - c. Repeat steps a-b one time.
 - Remove residual ethanol.
- 7. Air-dry for 3–5 minutes.
- 8. Add 32 μ l low TE buffer (10 mM Tris, pH 8.0 and 0.1 mM EDTA).
- 9. Resuspend the beads:
 - » For a plate, shake at 1500–1800 rpm for 2 minutes.
 - » For tubes, pipette 10 times.
- 10. Incubate at room temperature for 2 minutes.
- 11. Place on the magnet and wait until supernatant clears.
- 12. Transfer 30 μl *supernatant* to a DNA LoBind tube.
- 13. Proceed to <u>Quantify Library</u> and sequencing or store at -25° C to -15° C for ≤ 15 days.

Quantify Library

- 1. Quantify 2 μl library using Qubit.
- 2. Size 2 μl library using TapeStation.
- 3. Calculate the molar concentration:

 $nM = (ng/\mu l)/(660 * average library size) * 1,000,000$

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