



Red-Taq Polymerase 2x Master Mix

2x Master Mix Kit (1.5 mM MgCl₂)

Cat. No.	Reactions	R-Taq DNA Polymerase	MgCl ₂
AO180303	500	2x Master Mix	1.5 mM
AO180306	2,500	2x Master Mix	1.5 mM
AO180307	5,000	2x Master Mix	1.5 mM
AO180308	10,000	2x Master Mix	1.5 mM
AO180309	20,000	2x Master Mix	1.5 mM

Store at -20°C.

For in vitro laboratory use only

General Description

AS ONE R-Taq DNA Polymerase Mix is a ready-to-use 2x reaction mix. Simply add primers, template, and water to successfully carry out primer extensions and other molecular biology applications.

AS ONE Taq polymerase, the NH₄⁺ buffer system, dNTPs, inert red dye, stabilizer, and magnesium chloride are conveniently present in the R-Taq DNA Polymerase Mix.

AS ONE R-Taq DNA Polymerase Mix offers several advantages. Set up time is significantly reduced. The chance of contaminating component stocks is eliminated. Reduction of reagent handling steps leads to better reproducibility. Standard tests can be set up with the confidence that results will be consistent every time.

There is no need to use a separate loading dye. Simply load a portion of the reaction product onto an agarose gel for electrophoresis and subsequent visualization. The red dye front runs at 1000-2000 bp on a 0.5-1.5% agarose gel.

Composition of 2x R-Taq Master Mix

- 150 mM Tris-HCl pH 8.5, 40 mM (NH₄)₂SO₄, 3 mM MgCl₂, 0.2% Tween 20®
- 0.4 mM dNTPs
- 0.2 units/μL AS ONE Red-Taq polymerase
- Inert red dye and stabilizer

Suggested Protocol using R-Taq 2x Mix

This protocol serves as a guideline for primer extensions. Optimal reaction conditions such as incubation times, temperatures, and amount of template DNA may vary and must be individually determined.

Notes:

- Set up reaction mixtures in an area separate from that used for DNA preparation or product analysis. Work on ice at all times.
- The final MgCl₂ concentration of this 2x Master Mix is 1.5 mM. If more than 1.5 mM MgCl₂ is required, use 25 mM MgCl₂ (may be purchased separately) to adjust the Mg²⁺ concentration.

Additional volume (μl) of MgCl₂ per 50 μl reaction

Final MgCl ₂ conc. in reaction (mM)	1.5	2.0	2.5	3.0	3.5	4.0	4.5
Additional volume of 25 mM MgCl ₂ per reaction (μL):	0	1	2	3	4	5	6

- Thaw R-Taq 2 Master Mix and primers. It is important to thaw the solutions completely and mix thoroughly before use to avoid localized concentrations of salt. Keep all components on ice.
- The table below shows the reaction set up for a final volume of 50 μL. If desired, the reaction size may be scaled down. Use 10 μl of the R-Taq 2x Master Mix in a final volume of 20 μl.
- **Important:** Spin R-Taq Master Mix vials briefly before use.

1. Set up each reaction as follows:

Component	Vol./reaction	Final Conc.
R-Taq 2x Master Mix	25 μL	1X
25 mM MgCl ₂	0 μl (0-7 μl)	1.5 mM (1.5-5 mM)
Primer A	1 μL (0.5-5 μl)	0.2 μM (0.1–1.0 μM)
Primer B	1 μL (0.5-5 μl)	0.2 μM (0.1–1.0 μM)
Distilled Water	Variable	----
Template DNA	Variable	Genomic DNA 50 ng (10-500 ng) Plasmid DNA 0.5 ng (0.1-1 ng) Bacterial DNA 5 ng (1-10 ng)
TOTAL volume	50 μL	----

2. Mix gently by pipetting the solution up and down a few times.
3. Program the thermal cycler according to the manufacturer's instructions. For maximum yield and specificity, temperatures and cycling times should be optimized for each new template target or primer pair.
4. Place the tubes in the thermal cycler and start the reaction.



Three- step PCR Program:

Cycles	Duration of Cycles	Temperature
1	2-5 minutes ^a	95°C
25-35	20-30 seconds ^b	95°C
	20-40 seconds ^c	50-65°C
	30 seconds ^d	72°C
1	5 minutes ^e	72°C

^a Initial denaturation step. Optional, but recommended for genomic DNA

^b Denaturation step: Heating the DNA template disrupts the hydrogen bonds yielding ssDNA.

^c Annealing step: Primers anneal to ssDNA template. Typically annealing temperature is 3-5°C below the T_m of the primers.

^d Extension/elongation step: R-Taq DNA polymerase synthesizes a new DNA strand complementary to the DNA template. Extension time depends on length of DNA fragment to be amplified. At optimum temperature the DNA polymerase will polymerize 1000 bases per minute.

^e Final elongation step: After the last PCR cycle to ensure that any remaining ssDNA is fully extended.

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