Invitrogen™ Platinum™ SuperFi™ II PCR Master Mix

invitrogen

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Print Options

Package contents

Catalog number	Size
12368010	100 reactions
12368050	500 reactions
12368250	5 × 500 reactions

Kit contents

Storage conditions

- Store all contents at –20°C.

Required materials

Click here for required materials

Product description

- Platinum™ SuperFi™ II DNA Polymerase is a proofreading DNA polymerase that combines high fidelity with Platinum™ hot-start technology and universal primer annealing. It is ideally suited for cloning, mutagenesis, and other applications.
- Platinum™ SuperFi™ II PCR Master Mix is a ready-to-use mixture of DNA polymerase, salts, magnesium, and dNTPs for efficient PCR amplification, which retains all the features of the Platinum™ SuperFi™ II DNA Polymerase.
- The annealing temperature with Platinum™ SuperFi™ II DNA Polymerase is 60°C. Proprietary additives in the reaction buffer stabilize primer-template duplexes during the annealing step, and contribute to increased specificity without the need to optimize annealing temperature for each primer pair.
- Due to proprietary additives in the reaction buffer, Platinum™ SuperFi™ II DNA Polymerase shows efficient amplification of both AT and GC rich targets. Additional DNA melting agents are not required for GC-rich PCR (up to 75% GC).
- Platinum™ hot-start technology inhibits DNA polymerase activity at ambient temperatures, allowing room temperature reaction setup and storage of pre-assembled PCR reactions. Enzyme activity is restored after the initial denaturation step.
- Platinum™ SuperFi™ II DNA Polymerase has 5' to 3' polymerase and 3' to 5' exonuclease activities, but lacks 5' to 3' exonuclease activity. It produces blunt end DNA products.

Selection guide

PCR Enzymes and Master Mixes

Go online to view related products.

Online resources

Visit our product page for additional information and protocols.

For support, visit thermofisher.com/support.

Enzyme characteristics

Hot-start:	Antibody
Length:	Up to 20 kb
Fidelity vs. *Taq*:	>300X
Timing:	Varies depending on amplicon length
Format:	Master Mix

PCR setup

Component	Final concentration	20-µL rxn	50-µL rxn
2X Platinum™ SuperFi™ II PCR Master Mix [1]	1X	10 µL	25 µL
Forward primer	0.5 µM [2]	X µL	X µL
Reverse primer	0.5 µM [2]	X µL	X µL
Template DNA	0.1–10 ng plasmid (5–100 ng genomic DNA)	X µL	X µL
Water, nuclease-free	—	to 20 µL	to 50 µL

[1] Provides 1.5 mM $MgCl_2$ in 1X concentration.

[2] Reduce the primer concentration to 0.2 µM final for amplification of >5 kb targets from genomic DNA and for multiplex reactions.

PCR protocol

See page 2 to prepare and run your PCR experiment.

Important guidelines

Click here for important PCR guidelines.

Optimization strategies and troubleshooting

Click here for guidelines to optimize your PCR experiment.

Click here for guidelines to troubleshoot your PCR experiment.

Purchaser notification

Click here for Limited Warranty, Disclaimer, and Licensing information.

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Prepare and run PCR

The example PCR procedure below shows appropriate volumes for a single **20-µL** or **50-µL** reaction. For multiple reactions, prepare a master mix of components common to all reactions to minimize pipetting error, then dispense appropriate volumes into each 0.2-mL or 0.5-mL PCR tube before adding template DNA and primers.

<table>
<tr><th>Steps</th><th>Action</th><th>Procedure details</th></tr>
<tr><td>1  </td><td>Thaw reagents</td><td>Thaw, mix, and briefly centrifuge each component before use.</td></tr>
<tr><td>2  </td><td>Prepare reaction mix with template DNA and primers</td><td>Add the following components to each PCR tube.
Note: Consider the volumes for all components in reaction mix to determine the correct amount of water required to reach your final reaction volume.
<table>
<tr><th>Component</th><th>Final concentration</th><th>20-µL rxn</th><th>50-µL rxn</th></tr>
<tr><td>2X Platinum™ SuperFi™ II PCR Master Mix [1]</td><td>1X</td><td>10 µL</td><td>25 µL</td></tr>
<tr><td>Forward primer</td><td>0.5 µM [2]</td><td>X µL</td><td>X µL</td></tr>
<tr><td>Reverse primer</td><td>0.5 µM [2]</td><td>X µL</td><td>X µL</td></tr>
<tr><td>Template DNA</td><td>0.1–10 ng plasmid
(5–100 ng genomic DNA)</td><td>X µL</td><td>X µL</td></tr>
<tr><td>Water, nuclease-free</td><td>—</td><td>to 20 µL</td><td>to 50 µL</td></tr>
</table>
[1] Provides 1.5 mM $MgCl_2$ in 1X concentration.
[2] Reduce the primer concentration to 0.2 µM final for amplification of >5 kb targets from genomic DNA and for multiplex reactions.
Cap each tube, mix, and then briefly centrifuge the contents.</td></tr>
<tr><td>3  </td><td>Incubate reactions in a thermal cycler
(3-step protocol)</td><td>3-step protocol
<table>
<tr><th>Cycle step</th><th>Temperature</th><th>Time</th><th>Cycles</th></tr>
<tr><td>Initial denaturation</td><td>98°C</td><td>30 seconds</td><td>1</td></tr>
<tr><td>Denaturation
Annealing
Extension</td><td>98°C
60°C
72°C</td><td>5–10 seconds
10 seconds
15–30 seconds per 1 kb</td><td>25–35</td></tr>
<tr><td>Final extension</td><td>72°C
4°C</td><td>5 minutes
Hold</td><td>1
—</td></tr>
</table>
Note: Refer to “Optimization strategies”, page 1, for guidelines to optimize cycling conditions.</td></tr>
</table>

<table>
<tr><th>Steps</th><th>Action</th><th>Procedure details</th></tr>
<tr>
<td>3  </td>
<td>Incubate reactions in a thermal cycler
(2-step protocol)</td>
<td>
2-step protocol (for primers >30 nt long)[1]
<table>
<tr><th>Cycle step</th><th>Temperature</th><th>Time</th><th>Cycles</th></tr>
<tr><td>Initial denaturation</td><td>98°C</td><td>30 seconds</td><td>1</td></tr>
<tr><td>Denaturation
Annealing/Extension</td><td>98°C
72°C</td><td>5–10 seconds
15–30 seconds per 1 kb</td><td>25–35</td></tr>
<tr><td>Final extension</td><td>72°C
4°C</td><td>5 minutes
hold</td><td>1
—</td></tr>
</table>
[1] Without non-complementary parts (e.g. restriction tags).
Note: Refer to “Optimization strategies”, page 1, for guidelines to optimize cycling conditions.
</td>
</tr>
<tr>
<td>4  </td>
<td>Add gel loading buffer and analyze with gel electrophoresis</td>
<td>
Add gel loading buffer to 10 µL of PCR product, mix, and briefly centrifuge the contents.
Note: For optimal separation using E-Gel™ agarose gels, dilute the sample 2- to 20-fold.
Analyze the sample using agarose gel electrophoresis.
Use your PCR product immediately in down-stream applications, or store it at –20°C.
</td>
</tr>
</table>