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PCR Amplification of Desired Gene

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Protocol status: Working

We use this protocol and it's working

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Materials

water

Green HF buffer

dNTPS

F Primer

R Primer

Template DNA

Phusion DNA Polymerase

1% agarose gel

10000x GelRed nucleic acid stain

1x TAE buffer

Load DNA ladder



1 Pipette PCR components in the following order. Always set up at least two replicates.

2

# React ions	[μΙ]	Ther mocy cler Condi tions	Roun ds
Water	32.5	1. 98°C for 0:30 min	
Gree n HF Buffe r	10.0	2. 98°C for 0:10 min	5x
dNTP s (10 mM)	1.0	3. X°C for 0:20 min	
F Prime r (10 μΜ)	2.5	4. 72°C for X min	
R Prime r (10 µM)	2.5	5. 98°C for 0:10 min	30x
Temp late DNA (10 ng/ µL)	1.0	6. 72°C for X min	
Phusi on DNA Poly mera se	0.5	7. 72°C for 5:00 min	
Total	50.0	8. 4°C until use	



- 3 For step 3, use the annealing temperature of the gene-specific sequences of your primers.
 - For step 4 and 6, adjust the extension time based on the amplicon size (30 s / kb).
- 4 Prepare a 1 % agarose gel by dissolving agarose in 1x TAE buffer by microwaving (~ 3 min, swirling every min).
 - Add 2.5 µL of 10000x GelRed nucleic acid stain per 50 mL gel and mix well by swirling. When the conical flask is almost cool enough to touch (3-5 min after microwaving), pour slowly in gel tray and then insert the combs. Use wide-wells for large PCR reaction volumes.
 - Gel will be solidified after 45-60 min. Remove combs and add enough 1x TAE buffer to cover the gel.
- 5 Load 2-3 µl DNA ladder and whole PCR samplesinto the gel pockets. Connect lid (DNA will migrate to the + electrode). Turn on power supply and run at 80-110 V depending on the thickness and size of the gel. Examine gel after 30-60 min of electrophoresis and run until the marker reached the desired height.
 - Extract bands of the expected height from the gel with and clean up the fragments with an extraction kit.
- 6 Thermocycler Step 3: 5 cycles with annealing temperature that is equal to the forward or reverse primer with lower Tm. Has to be adjusted for every gene.
- 7 Termocycler Step 4: Adjust the elongation time with 30 s for each kb.
- 8 Termocycler Step 6: 30 cylces with 72 °C for elongation and adjusted extention time. Change time with 30 s per kb.
- 9 If not enough DNA was produced run another PCR after DNA extraction with the purified amplicon as a template and skip the 5 initial rounds.
- 10 Other optimisation: Increase time to 40 sec/kb