Protocol

# PanGreen<sup>™</sup> Universal SYBR<sup>®</sup> Green Master Mix

#### 08 DEC 2020

Catalog Number	Size	Concentration
QSD01-0100	100 reactions (20 $\mu$ l vol)	2X

Advion Interchim

#### **Storage Conditions**

Stable for up to 3 months at 4°C. Stable for up to 24 months at -20°C.

#### Description

PanGreen<sup>™</sup> Universal SYBR<sup>®</sup> Green Master Mix is a 2x concentrated, ready for use Master Mix reaction enhanced for dye-based quantitative PCR (qPCR) and compatible with the majority of commercially available real-time PCR systems (ROX-independent and ROX-dependent). It contains NanoTaq hot-start DNA polymerase, dNTPs, MgCl<sub>2</sub>, SYBR<sup>®</sup> Green I dye, enhancers, stabilizers and essentials for a success PCR reaction.

#### Kit Content(s)

2X universal SYBR<sup>®</sup> Green Master Mix

1ml x 1 vial

#### Required materials but not provided

- A compatible real-time PCR instrument
- Vortex or equivalent
- Microcentrifuge
- Plates and seals for your instruments

#### **Instrument Compatibility**

This Master Mix is compatible with the majority of commercially available real-time PCR systems.

#### **Reaction Setup**

- Thaw PanGreen<sup>™</sup> Universal SYBR<sup>®</sup> Green Master Mix and the rest of frozen reaction components to a temperature of 4°C. In order to entirely collect solutions, combine thoroughly and centrifuge briefly, then store at 4°C and avoid from light.
- Prepare (on ice or at room temperature) enough assay Master Mix for all reactions by adding all necessary components, except the DNA template, according to the recommendations in Table 1 (below).





Components	Volume per 20 µl Reaction	Volume per 10 µl Reaction	Final Concentration
PanGreen™ Universal SYBR <sup>®</sup> Green Master Mix (2x)	10 µl	5 µl	1x
Forward and reverse primers	Variable	Variable	300–500 nM each primer
DNA template (add at step 4)	Variable	Variable	cDNA: 1pg–10ng Genomic DNA: 50ng-250ng
Nuclease-free H <sub>2</sub> O	Variable	Variable	
Total reaction mix volume	20 µl	10 µl	1500-01 III (

- 3. Combine the assay Master Mix thoroughly to ensure consistency and equally dispense the solution into each qPCR tube or into the wells of a qPCR plate. Employ good pipetting practice to ensure assay precision and accuracy.
- Add DNA samples (and DNase-free H<sub>2</sub>O if needed) to the PCR tubes or wells containing assay Master Mix (Table 1), seal the tubes or wells with flat caps or optically transparent film. Note: to ensure thorough mixing of reaction components, vortex for approximately 30 seconds (or more).
- 5. Spin the tubes or plate to remove any air bubbles and collect the reaction mixture in the vessel bottom.
- 6. Setup the thermal cycling protocol on a real-time PCR instrument according to Table 2. **Note:** optimization may be needed for better performance.
- 7. Load the PCR tubes or plate into the real-time PCR instrument and commence the run.
- 8. Perform data analysis according to the instrument-specific instructions.
- Process in the thermal cycler for 35~45 cycles as follows:

Table 2. Thermal Cycling Protocol		
Initial Denaturation	3-5 minutes at 95°C (5 mins for GC rich or complex templates)	
Denaturation	15 seconds at 95°C	
Annealing & Extension	60 seconds at 60°C and Plate Read	
Melting curve	Refer to specific guidelines for instrument used	

Note: Optimal conditions for amplification will vary depending on the primers and thermal cycler used. It may be necessary to optimize the system for individual primers, template, and thermal cycler.



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## Template

Purified high quality DNA is needed for a success PCR reaction. The final concentration of cDNA template please refer to table 1.

### Important notes

- 1. Shake gently before use to avoid foaming and low-speed centrifugation.
- 2. During operation, always wear a lab coat, disposable gloves, and protective equipment.

### Troubleshooting

Refer to the table 3 below to troubleshoot problems that you may encounter when quantifying of nucleic acid targets with the kit.

Table 3. Troubleshooting			
Trouble	Cause	Solution	
Poor Signal or No Signal	Inhibitor Present	<ol> <li>Perform a dilution series of the PCR template to determine whether the effect of the inhibitory agent can be reduced.</li> <li>Take extra care with the nucleic acid extraction steps to minimize carryover of PCR inhibitors.</li> </ol>	
	Degraded Template Material	<ol> <li>Do not store diluted template in water or at low concentrations.</li> <li>Check the integrity of template material by automated or manual gel electrophoresis.</li> </ol>	
	Inadequate Thermal Cycling Conditions	<ol> <li>Try using a minimum extension time of 30 sec for genomic DNA and 15 sec for cDNA.</li> </ol>	
Signal in Negative Control	Contamination of Reaction Components with Target Sequence	<ol> <li>To minimize the possibility of contamination of PCR components by PCR product or other template, designate a work area exclusively for PCR assay setup.</li> <li>Use a solution of 10% bleach instead of ethanol to prepare the workstation area for PCR assay setup. Ethanol will only induce precipitation of DNA in your work area, while the 10% bleach solution will hydrolyze, as well as dissolve, any residual DNA.</li> </ol>	
Poor Reproducibility Across Replicate Samples	Inhibitor Present	<ol> <li>Perform a dilution series of the PCR template to determine whether the effect of the inhibitory agent can be reduced.</li> <li>Take extra care with the nucleic acid extraction steps to minimize carryover of PCR inhibitors.</li> </ol>	
	Primer Design	1. Verify primers design at different annealing temperatures.	



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		1. Reduce primer concentration.
		2. Evaluate primer sequences for complementarity and secondary
Low or High	Primer- Dimer	structure. Redesign primers if necessary.
Reaction		3. Perform melt-curve analysis to determine if primer- dimers are
Efficiency		present.
	Insufficient	1. Use a thermal gradient to identify the optimal thermal cycling
	Optimization	conditions for a specific primer set.





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