

# BioArk® 2×SYBR Green qPCR Master Mix (Low ROX)

## Cat. #: BSY3320LR

#### **Product Information**

Product Name	Cat. No.	Spec.
	BSY3320LR-01	1 mL
2×SYBR Green qPCR Master Mix (Low ROX)	BSY3320LR-05	5×1 mL
	BSY3320LR-15	15×1 mL

# **Product Description/Introduction**

This product is a special 2× premixed solution for qPCR reaction using SYBR Green I chimeric fluorescence method. It contains all qPCR components except primers and DNA templates, which can reduce the operation steps, shorten the sample addition time and decrease the probability of contamination. The core component is the genetically engineered heat-activated Taq DNA Polymerase, which effectively blocks DNA Polymerase activity and prevents nonspecific amplification at low temperatures by efficiently binding monoclonal antibodies to the Taq DNA Polymerase. The heat-activated Taq DNA Polymerase has many advantages such as high specificity and detection sensitivity. With the reaction buffer optimized for qPCR, it is very suitable for high-specificity and high-sensitivity qPCR reactions. This product is a 2 × premixed reagent containing the optimal concentration of SYBR Green I for qPCR reaction, which can obtain a good standard curve in a wide quantitative area. The quantification of target genes is accurately, reproducible and reliable.

#### **Storage and Handling Conditions**

Transport with wet ice. Store at -20 ° C without light, valid for 12 months. Avoid freeze/thaw cycle. After thawing, it can be stably stored at 4°C for one month without light.

#### Component

Component	BSY3320LR-01	BSY3320LR-05	BSY3320LR-15
2×SYBR Green qPCR Master Mix (Low ROX)	1 mL	5×1 mL	15×1 mL
Manual	One copy		

## Assay Protocol / Procedures

#### Preparation before experiment

- 1. Real Time PCR amplification apparatus;
- 2. Special reaction tube or reaction plate for experiment;
- 3. PCR primers (reference primer design principles);
- 4. Micropipette and pipette head (autoclaving);

# Procedures

#### 1. Recommend the qPCR reaction system:

Component	20 μL rxn	50 μL rxn	Final Concentration
2×SYBR Green qPCR Master Mix (Low ROX)	10 μL	25 μL	1×
Forward Primer (10 μM) <sup>a</sup>	0.4 μL	1 μL	0.2 μΜ
Reverse Primer (10 μM) <sup>a</sup>	0.4 μL	1 μL	0.2 μΜ
Template <sup>b</sup>	Variable	Variable	as required
Nuclease-Free Water	Add to 20 μL	Add to 50 µL	



- a. Usually, a good amplification effect can be obtained with the final concentration of 0.2  $\mu$  M. When the reaction performance is poor, the primer concentration can be adjusted in the range of 0.2-1.0  $\mu$ M.
- b. The amount of template addition varies with the copy number of the target gene in the template solution, and the appropriate amount of template addition is discussed by gradient dilution. The best addition amount of template DNA in the 20  $\mu$ L reaction system was less than 100 ng. When the cDNA (RT reaction solution) of RT-PCR reaction was used as template, the addition amount should not exceed 10% of the total volume of PCR reaction solution.

2. PCR reaction procedure (can be adjusted according to the instruments):

A. Two-step method *			B. Three-step method *						
Stag	Step	Cycl	Temper	Time	Stag	Step	Cycles	Temperatu	Time
е	٥٠٥٦	es	ature		е	3 3 p	0,0.00	re	
Stag	Predegenerat	1	95℃	30 sec	Stag	Predegenera	1	95℃	30
e 1	ion	1	30 0	30 300	e 1	tion		33 C	sec
	Degeneration		95°€	95℃ 15 sec		Degeneratio		95℃	15
	Degeneration		33 0 13		n		30 0	sec	
Stag		40		Stag	Annealing	40	55-65°C	10	
e 2	Annealing-Ex		60℃	10	60°C 30 sec <sup>a</sup>	e 2	7 time aimig	40	33 03 0
	tension		00 C 30 Se	30 sec	Extension		72°C	30	
				Extension		72 C	sec <sup>a</sup>		
Stag	Stag e 3 Melting curve 1	1	Instrument	nt default	Stag	Melting	1	Instrument	default
e 3		1	Settings		e 3	curve	1	Setting	JS

<sup>\*:</sup> If amplification specificity needs to be improved, two-step procedure or annealing temperature can be used; To improve the amplification efficiency, a three-step procedure or extension time can be used.

#### Note

- 1. Mixed gently upside down before use. Do not swirl and shake to avoid bubbles. Mix the reagents well before using.
- 2. Reagents should be placed on ice when preparing reaction solution.
- 3. The product contains fluorescent dye SYBR Green, so strong light should be avoided when preparing PCR reaction solution.
- 4. Please using new disposable head for the preparation and packaging of the reaction solution to avoid contamination between samples.
- 5. Avoid repeated freeze-thawing of Master Mix and try to use it within one month after thawing.

# Compatibility

Brand of PCR machine	BSY3320NR (None ROX)	BSY3320LR (Low ROX)	BSY3320HR (High ROX)
ABI Thermo life	PikoReal™ Cycler	7500/7500 Fast, ViiA 7™ QuantStudio™ series	5700/7000/7300/7700/7900/ 7900HT/7900 HT Fast, StepOne™, StepOne Plus™
Stratagene		Mx3000P <sup>®</sup> /3005P <sup>™</sup> /4000 <sup>™</sup>	
Bio-Rad	All series		
Eppendorf	Realplex 2s, Mastercycler <sup>®</sup> ep realplex		
Illumina	Eco QPCR		
Cepheid	SmartCycler <sup>®</sup>		

a: For fluorescence signal collection, please set the experimental procedure according to the instruction manual of the instrument.



Qiagen Corbett	Rotor-Gene® series	
Roche	LightCycler™ series	
Takara	Thermal Cycler Dice series	
Analytikjena	qTOWER series	
qTOWER	LineGene series	

# **Primer Design Principles**

- 1. The length of amplification product is recommended to be between 80-300 bp;
- 2. Primer length: 18-25 bp;
- 3. The content of base G+C in primers should be between 40%-60%;
- 4. The Tm value difference between forward primers and reverse primers is less than 2°C, and the Tm value between 58-62°C is the best;
- 5. Randomness of base distribution;
- 6. Primers had better not contain self-complementary sequences, otherwise they will form a secondary hairpin structure;
- 7. There should be no more than 4 complementary or homologous bases between two primers, otherwise primer dimer will be formed, especially complementary overlap at the 3' end;
- 8. The 3' terminal base of the primer is suggested to be G or C;
- 9. No other non-specific products were found in NCBI comparison results.

# **Trouble-Shooting**

Problem description	Possible reasons	Solutions	
	The template concentration is too low	Repeat the experiment to reduce the template dilution multiple, and start from the highest concentration when the sample concentration is unknown	
	Template degradation	The template was prepared again and the experiment was repeated	
At the end of the reaction, no amplification curve	There are PCR inhibitors in the system	Generally, the template is carried in, the dilution ratio of the template is increased or the template with high purity is reprepared and repeated	
appeared or CT value appeared too late	Primers may degrade	Primers that have not been used for a long time should first be tested for integrity by PAGE electrophoresis to rule out the possibility of degradation	
	Low amplification efficiency	Increase the primer concentration, try a three-step amplification procedure, or redesign the primer	
	The amplification product is too long	The amplification product length was controlled in the range of 80-300 bp	
The blank control shows the signal	Reaction system pollution	Firstly, the blank control water should be replaced. If the same situation still occurs, the primers, aspirators and PCR tubes should be replaced or a	



		new Master Mix should be started. The reaction system is prepared in a super clean table to reduce aerosol pollution		
	Non-specific amplification such as primer dimers appears	Generally, it is normal for the amplification products to appear in blank control after 35 cycles, which should be analyzed with the melting curve.  Redesign primer, adjust primer concentration or optimize PCR reaction procedure		
	Primer design is poor	The new primer was re-designed according to the primer design principles		
The melting curve has	Primer concentration is too high	Reduce primer concentration appropriately		
multiple peaks	There is genomic contamination in cDNA template	The extracted RNA solution is digested using DNA enzymes, such as dsDNase, to remove genomic contamination, or to design transintron primers		
	The error of adding sample is large	The use of accurate pipette, with high quality suction head accurate pipette; High dilution template, adding large volume template to reduce sampling error; The reaction volume of qPCR was enlarged		
Poor reproducibility of experiments	The template concentration is too low	Repeat the experiment to reduce the dilution times of the template		
	Temperature deviation at different locations of the qPCR instrument	Calibrate the qPCR instrument regularly		
The amplification curve is not smooth	Fluorescence signal is too weak, produced after system correction	Ensure that the dyes premixed in the Master Mix are not degraded; Replace fluorescent signal to collect better qPCR consumables		
Amplification curve breaks or slips	The template concentration was higher and the baseline endpoint value was greater than the CT value	The baseline endpoint (Ct value -3) was reduced and the data were reanalyzed		
Amplification curves of individual Wells suddenly dropped sharply	There are bubbles in the reaction tube	Ensure that MIX is completely dissolved, and do not swirl and oscillate evenly; After the sample is added, the bubbles are removed by centrifugation with light elastic. The pre-denaturation time was extended to 10 min to remove the bubbles		

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