



PCRBIOSYSTEMS

simplifying research

## VeriFi™ Hot Start Polymerase

[www.pcrbio.com](http://www.pcrbio.com)

### Product description:

VeriFi™ Hot Start Polymerase is a robust and versatile proofreading enzyme with AptaLock™ hot start technology for highly precise PCR. The enzyme is designed for all PCR applications where greater sequence accuracy is required, together with improved PCR success rates of long and challenging templates.

VeriFi™ Hot Start Polymerase is derived from Pfu DNA polymerase for its 3'-5' exonuclease (proofreading) activity. The enzyme is engineered with proprietary mutations that significantly increase processivity, resulting in shorter extension times (30 s/kb), higher yields and the ability to amplify longer and more difficult targets, including eukaryotic genomic templates in excess of 17.5 kb.

PCRBio's innovative AptaLock™ technology uses a proprietary aptamer-like molecule that reversibly inhibits both the 3'-5' exonuclease activity and 5'-3' polymerase activity of the enzyme at ambient temperatures. This unique hot start molecule prevents primer dimer formation and non-specific amplification to maximise the sensitivity and specificity of your PCR. This feature makes the enzyme highly suitable for multiplexing and enables reactions to be set up at room temperature.

The enhanced accuracy of VeriFi™ Hot Start Polymerase results in fidelity that is approximately 100 times higher than Taq DNA polymerase. The enzyme is ideal for applications where greater accuracy is needed, such as cloning, site-directed mutagenesis and sequencing. PCR products generated with this range of products are blunt ended.

Component	100 units	500 units
VeriFi™ Hot Start Polymerase (2 u/μL)	1 x 50 μL	1 x 250 μL
5x VeriFi™ Buffer	1 x 1.7 mL	3 x 1.7 mL
10x VeriMax Enhancer	1 x 1.7 mL	2 x 1.7 mL

VeriFi™ Hot Start Polymerase is provided with an advanced buffer system including dNTPs, Mg and enhancers, enabling high fidelity PCR of a wide range of targets and fragment sizes regardless of GC content.

### Shipping and storage

On arrival the kit should be stored between -30 °C and -15 °C. If stored correctly the kit will retain full activity for 12 months. The kit can be stored at 4 °C for 1 month.

### Limitations of product use

The product may be used for in vitro research purposes only.

### Technical support

Help and support is available on our website at <https://pcrbio.com/resources/> including answers to frequently asked technical questions. For technical support and troubleshooting you can submit a technical enquiry online, or alternatively email [technical@pcrbio.com](mailto:technical@pcrbio.com) with the following information:

- Amplicon size
- Reaction setup
- Cycling conditions
- Screen grabs of gel images

## Important considerations

**5x VeriFi™ Buffer:** The 5x buffer contains 15 mM MgCl<sub>2</sub>, 5 mM dNTPs, enhancers and stabilizers. It is not recommended to add further MgCl<sub>2</sub> to the reaction. The buffer composition has been optimised to maximise PCR success rates.

**Reaction Enhancer:** In situations where no amplification is observed, we recommend adding the 10x VeriMax Enhancer to the reaction mix. This enhancer can improve the performance of VeriFi™ Hot Start Polymerase on some difficult or long templates, for example GC-rich templates or those with complex secondary structures.

**Primers:** Primers should have a predicted melting temperature of around 60 °C, using default Primer 3 settings (<http://bioinfo.ut.ee/primer3/>). The final primer concentration in the reaction should be between 0.2 µM and 0.6 µM.

**Denaturation:** Denaturation should be performed at 95 °C. However, if the presence of high GC regions results in low yields, increasing the melting temperature to 98-100 °C can improve the amount of product.

**Annealing:** We recommend performing a temperature gradient to experimentally determine the optimal annealing temperature. Alternatively, we recommend a 60 °C annealing temperature then increase in 2 °C increments if non-specific products are present.

**Extension:** Optimal extension is achieved at 72 °C. The optimal extension time is dependent on amplicon length and complexity of template. 30 seconds per kilobase (kb) is recommended for most applications. Two-step cycling protocols may also be used with combined annealing and extension at 68-75 °C.

**Multiplex PCR:** The optimal extension time for multiplex reactions will be dependent on the complexity of template, the length of amplicons, and the number of targets. We recommend starting with the extension time of the longest fragment, and then increasing in increments of between 10 and 30 seconds if necessary.

## Reaction setup

1. Allow 5x VeriFi™ Buffer (and 10x VeriMax Enhancer, if used) to reach room temperature, then briefly vortex.
2. Prepare a master mix based on the following table:

Reagent	25 µL reaction	50 µL reaction	Final concentration	Notes
5x VeriFi™ Buffer	5.0 µL	10.0 µL	1x	
10x VeriMax Enhancer (optional)	2.5 µL	5.0 µL	1x	See above for use of enhancer
Forward primer (10 µM)	1.0 µL	2.0 µL	400 nM	See above for optimal primer design
Reverse primer (10 µM)	1.0 µL	2.0 µL	400 nM	See above for optimal primer design
Template DNA	<100 ng genomic DNA <5 ng less complex DNA	<200 ng genomic DNA <10 ng less complex DNA	variable	
VeriFi™ Hot Start Polymerase (2 u/µL)	0.25 µL	0.5 µL		
PCR grade dH <sub>2</sub> O	Up to 25 µL final volume	Up to 50 µL final volume		

3. Cycle using conditions based on the following table:

Cycles	Temperature	Time	Notes
1	95 °C	1 min	Initial denaturation
25-35	95 °C	15 seconds	Denaturation (see above for high GC templates)
	60 °C to 75 °C	15 seconds	Anneal
	72 °C	30 seconds/kb	Extension (see above for multiplex PCR)