



5581

Blend Taq

Code: TYB-BTQ-101 (250 units)

Storage: Store at -20° C

| Concentration: | 2.5 units/ul | | | |
|----------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------|--|--|--|
| Unit Definition: | One unit of enzyme is defined as the amount of enzyme that will incorporate 10 nmoles of dNTPs into acid insoluble materials in 30 minutes at $75^{\circ}C$. | | | |
| Assay Condition: | $\begin{array}{llllllllllllllllllllllllllllllllllll$ | | | |
| Storage Buffer: | 20 mM Tris-HCI (pH 8.0) 100 mM KCI 0.1 mM EDTA 1 mM DTT 0.5 % Tween-20 0.5 % Nonidet P-40 50 % Glycerol | | | |
| Materials Provided: | 10×Buffer for Blend Taq (Mg ²⁺ concentration: 20mM) dNTPs : 2mM dATP, dGTP, dCTP, dTTP each | | | |
| Quality Control 1. PCR Assay: | The 17.5kbp fragment of human β globin gene could be amplified using human genomic DNA. | | | |

Purchase of this product is accompanied by a limited license to use it in the Polymerase Chain Reaction(PCR) process for Research Field in conjunction with a thermal cycler whose use in the automated performance of the PCR process is covered by the up-front license fee, either by payment to Perkin-Elmer or as purchased, i. e., an authorized thermal cycler.

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PCR conditions

Reaction mixture for General PCR Amplification

| 10 $	imes$ Buffer for Blend Taq | 5ul | | |
|---------------------------------|--------------------------------------|--|--|
| Blend Taq (2.5 U/ul) | 0.5ul (1.25U) | | |
| Template | 1-50ng (plasmid) 10-1000ng (Genome) | | |
| Primer | 10-50pmoles (final conc.: 0.2-1.0uM) | | |
| 2mM dNTPs | 5ul (final conc.: 0.2mM) | | |
| Total volume | 50ul | | |

Suggested Cycling parameters

| Segment - | | Number of | | |
|----------------|--------------------------------|---------------------------|------------------------------------------|--------|
| | <1kb | 1kb – 6kb | >6kb | Cycles |
| 1 | | 1 | | |
| 2 Denaturation | | | | |
| 3 Annealing | Primer Tm-5℃ (55-60℃) 30sec | | 68℃ | 30 |
| 4 Extension | 72℃ 1min | 72℃ 1min/kb PCR target | 1min/kb PCR target (see notes #2, #3) | |

<u>Notes</u>

- 1. The denaturation time and temperature may require optimization. Typically denaturation time will range 25 30sec at 94° C or 5 10sec at 98° C.
- 2. Toyobo suggests >70 $^{\circ}$ C melting temperature of primer (Tm) for long target (>6kb).
- 3. A dNTP concentration range of 0.3 0.4mM total is recommended for longer target (>10kb).
- 4. Blend Taq produces amplification products that are ready to clone directly into TA cloning vectors without any special optimization. To get sufficient numbers of transformants and white colony, longer ligation time will be efficient, recommended 2 hours overnight for the amplification products of >1kb.
- 5. Blend Taq-Plus- is designed to be suitable for Hot Start PCR. Non-specific amplification due to mispriming and/or formation of primer dimer before thermal cycling can be prevented.

For research use only; not for use as a diagnostic.

