# Datasheet



#### **Description:**

SNPase is Taq DNA Polymerase with unique N-terminal deletion and proprietary amino acids substitutions introduced into the active center of the enzyme. This modification causes dramatic increase of sensitivity of the enzyme to mismatches at 3'-end of the primer. Consequently , non-perfect annealing of the primers does not result in unspecific amplicons formation. This enzyme has only 5'-3' polymerase activity and is recommended for SNP genotyping by allele-specific PCR (AS-PCR), allele-specific primer extension (AS-PEX) and minisequencing procedures.

#### **Features:**

- 10-15 fold lower mutation rate than Taq DNA Polymerase
- high fidelity allele-specific amplification of DNA fragments
- high specificity with lowest background AS-PEX and AS-PCR
- Hot-Start activity for less primer dimers
- only 5'-3' polymerase activity, lack of 5'-exonuclease activity

#### **Applications:**

- High specific PCR
- Single Nucleotide Polymorphism (SNP)
- Multiplex PCR
- Real-Time PCR with intercalation dyes
- high fidelity dNTPs and ddNTPs
- Mini-Sequencing, SNP-genotyping

#### **Unit definition:**

One unit is defined as the amount of enzyme that incorporates 10nmoles of dNTPs into acid-insoluble form in 30 minutes at 72°C.

Concentration: 20-25 u/µl

#### **Reaction Buffer supplied:**

5X Reaction buffer without MgCl<sub>2</sub> MgCl<sub>2</sub> 100 mM

#### Note:

- optimal MgCl<sub>2</sub> concentration: 3.0 -3.5 mM in the 1X reaction mixture
- higher MgCl<sub>2</sub> concentrations results in higher yield (up to 4.5 mM)
- lower MgCl<sub>2</sub> (2.5 mM) results in higher specificity
- DNA fragments up to 400 bp from Human genomic DNA and 500 bp from Phage-DNA



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Components	Volume per reaction
5X reaction buffer	5 μΙ
MgCl <sub>2</sub>	2.5 - 4 mM
dNTP-Mix	0.2 mM each
primer mix (5 µM stock)	0,9-1,1 μl (5 pmol)
Template DNA	75-125 ng/25 µl genomic DNA
SNpase (Single Nucleotide Polymorphism)	0.2 - 0.5 μl (5-12 units)
Sterile dest. Water (molecular grade)	up to 25 μl total reaction volume

## **General Thermo-Cycler protocol:**

Step	Time	Temperature
Initial denaturation	1-2 min	94-95°C
30-35 Cycles:		
Denaturation	20-30 sec	94-95°C
Annealing	15-30 sec	59-68°C
Extension	30-40 sec	68-72°C per 1kb
Final extension	5 min	72°C

### **Order Information**

Prod. No.	Description	Quantity
S9500	SNPase Hot Start Polymerase	500 Units
S9505	SNPase Hot Start Polymerase	2500 Units

