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## New England Biolabs Product Specification

| Product Name:          | AfeI  |
|------------------------|---|
| Catalog #:             | R0652S/L  |
| Concentration:         | 10,000 units/ml   |
| Unit Definition:       | One unit is defined as the amount of enzyme required to digest 1 μg of pXba DNA in rCutSmart Buffer in 1 hour at 37°C in<br>a total reaction volume of 50 μl. |
| Shelf Life:            | 24 months   |
| Storage Temp:          | -20°C   |
| Storage Conditions:    | 10 mM Tris-HCl, 300 mM NaCl, 1 mM DTT, 0.1 mM EDTA, 50% Glycerol, 500 μg/ml rAlbumin (pH 7.4 @<br>25°C)   |
| Specification Version: | PS-R0652S/L v2.0  |
| Effective Date:        | 02 Mar 2023   |

Assay Name/Specification (minimum release criteria)

Endonuclease Activity (Nicking) - A 50  $\mu$ l reaction in rCutSmart<sup>TM</sup> Buffer containing 1  $\mu$ g of supercoiled PhiX174 DNA and a minimum of 10 units of AfeI incubated for 4 hours at 37°C results in <20% conversion to the nicked form as determined by agarose gel electrophoresis.

**Exonuclease Activity (Radioactivity Release)** - A 50  $\mu$ l reaction in rCutSmart<sup>TM</sup> Buffer containing 1  $\mu$ g of a mixture of single and double-stranded [<sup>3</sup>H] *E. coli* DNA and a minimum of 100 units of AfeI incubated for 4 hours at 37°C releases <0.1% of the total radioactivity.

**Ligation and Recutting (Terminal Integrity)** - After a 10-fold over-digestion of pXba DNA with AfeI, >95% of the DNA fragments can be ligated with T4 DNA ligase in 16 hours at 16°C. Of these ligated fragments, >95% can be recut with AfeI.

Non-Specific DNase Activity (16 Hour) - A 50  $\mu$ l reaction in rCutSmart<sup>TM</sup> Buffer containing 1  $\mu$ g of pXba DNA and a minimum of 10 units of AfeI incubated for 16 hours at 37°C results in a DNA pattern free of detectable nuclease degradation as determined by agarose gel electrophoresis.

qPCR DNA Contamination (*E. coli* Genomic) - A minimum of 10 units of AfeI is screened for the presence of *E. coli* genomic DNA using SYBR® Green qPCR with primers specific for the *E. coli* 16S rRNA locus. Results are quantified using a standard curve generated from purified *E. coli* genomic DNA. The measured level of *E. coli* genomic DNA contamination is  $\leq 1$  *E. coli* genome.

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