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# In vitro digestion of DNA with Cas9 Nuclease, *S. pyogenes* (M0386) V.4

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[dx.doi.org/10.17504/protocols.io.be6fjhbn](https://dx.doi.org/10.17504/protocols.io.be6fjhbn)**New England Biolabs (NEB)**Tech. support phone: +1(800)632-7799 email: [info@neb.com](mailto:info@neb.com)**New England Biolabs**  
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Cas9 Nuclease, *S. pyogenes*, (Cas9) is a double-stranded DNA endonuclease that is guided to its target by sequence complementarity of a small RNA loaded into the protein. This protocol describes how to digest double-stranded DNA *in vitro* using Cas9 and a single guide RNA (sgRNA).

DOI

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Breton Hornblower

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## REFERENCES:

1. Jinek et al. (2012) Science 337 (6096) 816-821.
2. Larson et al. (2013) Nature Protocol 8 (2180-2196).
3. Mali et al. (2013) Science 339 (6121): 823-826.

## MATERIALS

 [Cas9 Nuclease, \*S. pyogenes\* - 70 pmol](#) **New England**

**Biolabs Catalog #M0386S**

 [Proteinase K, Molecular Biology Grade - 2 ml](#) **New England**

**Biolabs Catalog #P8107S**

 [HiScribe T7 Quick High Yield RNA Synthesis Kit - 50 rxns](#) **New England**

**Biolabs Catalog #E2050S**

 [Nuclease-free Water](#) **New England**

**Biolabs Catalog #E7764**

## REQUIRED MATERIALS:

- Cas9 Nuclease, *S. pyogenes* (NEB [#M0386](#))
- NEBuffer 3.1
- Nuclease-free water
- Proteinase K, Molecular Biology Grade (NEB [#P8107S](#))
- sgRNA containing the targeting sequence in the region of interest
- *sgRNAs can be generated by in vitro transcription using the HiScribe T7 Quick High-Yield RNA synthesis Kit (NEB [#E2050](#)) using linearized plasmid, PCR products, or oligonucleotides as templates*
- *sgRNAs must contain sequence complementary to the target DNA (1,2)*
- *For information on design of sgRNA transcription templates please visit [Addgene](#)*
- DNA substrate containing the target sequence
- The substrate DNA can be circular or linearized plasmid, PCR products, or synthesized oligonucleotides

## OPTIONAL MATERIALS:

Apparatus and reagents for DNA fragment analysis

- E. g. Agarose gel electrophoresis apparatus
- DNA Loading Dye (e.g. Gel Loading Dye, Purple (6X) NEB [#B7024S](#))
- E.g. Agilent Bioanalyzer or similar

Please refer to the Safety Data Sheets (SDS) for health and environmental hazards.

- We strongly recommend wearing gloves and using nuclease-free tubes and reagents to avoid RNase contamination. Further recommendations for avoiding ribonuclease contamination can be found [here](#).
- Reactions are typically 30 µl but can be scaled up as needed. Reactions should be assembled in nuclease-free microfuge tubes or PCR strip tubes.
- It is essential to keep the molar ratio of Cas9 and sgRNA per target site at 10:10:1 or higher to obtain the best cleavage efficiency. A calculator can be found [here](#).
- If planning to use higher concentration Cas9 Nuclease, *S. pyogenes* (NEB #M0386T and NEB #M0386M) for *in vitro* digestion of DNA, the enzyme can be diluted to

**1** **1 Micromolar (μM)** in **1 X Buffer 3.1** and used immediately. If the 1 μM dilution will be stored at **-20 °C**, it should be diluted using Diluent B (NEB #B8002S):

**300 Milimolar (mM) NaCl**, **10 Milimolar (mM) Tris-HCl**,

**0.1 Milimolar (mM) EDTA**, **1 Milimolar (mM) DTT**, **500 μg/ml BSA** and

**50 % glycerol** (**pH 7.4**) @ **25 °C** prior to the reaction assembly.

**1** Prepare **300 Nanomolar (nM) sgRNA** by diluting the stock with nuclease-free water **On ice**.

**2** Prepare **30 Nanomolar (nM) substrate DNA** with a single target sequence by diluting the stock with nuclease-free water **On ice**.

**3** Assemble the reaction at **Room temperature** in the following order:

A	B
COMPONENT	VOLUME (for 30 μl reaction)
Nuclease-free water	20 μl
NEBuffer 3.1	3 μl
300 nM sgRNA	3 μl (30 nM final)
1 μM Cas9 Nuclease, S.pyogenes (M0386S)	1 μl (~30 nM final)
<i>Reaction volume</i>	<i>27 μl</i>

\*The sgRNA and nuclease-free water are not included.

**4** 

Pre-incubate for **00:10:00** at **25 °C**.

**5** 

Add **3 μL 30 nM substrate DNA** (3 nM final).


**6** 

Mix thoroughly and pulse-spin in a microfuge.

**7** 

Incubate at **37 °C** for **00:15:00**.

8 

Add  **1 µL Proteinase K** to each sample. Mix thoroughly and pulse-spin in a microfuge.

9 

Incubate at  **Room temperature** for  **00:10:00**.

10 Proceed with fragment analysis.