

Feb 03, 2026

Version 2

Saturation Genome Editing V.2

DOI

<https://dx.doi.org/10.17504/protocols.io.rm7vzjyd4lx1/v2>

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Protocol status: Working

We use this collection and it's working

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Abstract

This collection details the saturation genome editing workflow.

Guidelines

Data Analysis

- Demultiplex samples from Illumina output directory using BCLConvert software.

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