

# Resource Summary Report

Generated by [dkNET](#) on Aug 6, 2026

## Sage Science: BluePippin system

RRID:SCR\_020505

Type: Tool

### Proper Citation

Sage Science: BluePippin system (RRID:SCR\_020505)

### Resource Information

**URL:** <https://sagescience.com/products/bluepippin/>

**Proper Citation:** Sage Science: BluePippin system (RRID:SCR\_020505)

**Description:** Used for DNA Size Selection for Next-Gen Sequencing, with Pulsed-Field (100bp ? 50kb).BluePippin has the DNA capabilities of the Pippin Prep, plus the extra benefit of pulsed-field electrophoresis for resolving and collecting high molecular weight DNA. For long-range genomic applications, high-pass filtering allows users to collect all fragments above a size threshold set by the user. Target sizes or ranges of sizes are entered in software, and fractions are collected in buffer. Up to 5 samples/gel cassette may be run, with no possibility of cross contamination.

**Resource Type:** instrument resource

**Keywords:** Sage, Automated DNA Size Selector, Instrument Equipment, USEDit

**Availability:** Commercially available

**Resource Name:** Sage Science: BluePippin system

**Resource ID:** SCR\_020505

**Alternate IDs:** Model\_Number\_BluePippin

**Record Creation Time:** 20220129T080350+0000

**Record Last Update:** 20260801T190652+0000

### Ratings and Alerts

No rating or validation information has been found for Sage Science: BluePippin system.

No alerts have been found for Sage Science: BluePippin system.

## Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We found 5 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [dkNET](#) .

Khan N, et al. (2025) Human Papillomavirus Integration Induces Oncogenic Host Gene Fusions in Oropharyngeal Cancers. *Cancer discovery*, 15(9), 1927.

Guang X, et al. (2025) Telomere-to-telomere African wild rice (*Oryza longistaminata*) reference genome reveals segmental and structural variation. *GigaScience*, 14.

Akagi K, et al. (2023) Intratumoral Heterogeneity and Clonal Evolution Induced by HPV Integration. *Cancer discovery*, 13(4), 910.

Wang Z, et al. (2022) A chromosome-level reference genome of *Ensete glaucum* gives insight into diversity and chromosomal and repetitive sequence evolution in the Musaceae. *GigaScience*, 11.

Smolander OP, et al. (2022) Improved chromosome-level genome assembly of the Glanville fritillary butterfly (*Melitaea cinxia*) integrating Pacific Biosciences long reads and a high-density linkage map. *GigaScience*, 11(1).