

Determining Genome Size from Spores of Seedless Vascular Plants

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[Abstract] Seedless vascular plants, including ferns and lycophytes, produce spores to initiate the gametophyte stage and to complete sexual reproduction. Approximately 10% of them are apomictic through the production of genomic unreduced spores. Being able to measure the spore nuclear DNA content is therefore important to infer their reproduction mode. Here we present a protocol of spore flow cytometry that allows an efficient determination of the reproductive modes of seedless vascular plants. **Keywords:** Apomixis, Bead-vortex, Fern, Flow cytometry, Lycophyte, Spore

[Background] In seedless vascular plants, sporogenesis features, such as meiotic chromosome counts, were traditionally used to infer nuclear DNA content as well as reproductive modes. However, these approaches are time-consuming, or can only provide indirect evidence. An efficient and reliable method to estimate spore nuclear DNA content of these plants had not been established until Kuo *et al.* (2017). Herein, we describe a protocol using flow cytometry to evaluate spore genome sizes of these plants based on the work of Kuo *et al.* (2017).

Materials and Reagents

- 1. Pipette tips (10, 100, and 1,000 µl)
- 2. 50-ml tube
- 3. 1.7-ml tubes with caps
- 4. 2.0-ml tubes with caps
- 5. 2.3-mm stainless steel beads (Bio Spec Products, catalog number: 11079123ss)
- 6. 30-µmnylon meshes (Sysmex, CellTrics[®], catalog number: 04-0042-2316)
- 7. 20-µmnylon meshes (Sysmex, CellTrics[®], catalog number: 04-0042-2315)
- 8. 10-µm nylon meshes (Sysmex, CellTrics[®], catalog number: 04-0042-2314)
- 9. Glass Petri dish (Corning, PYREX[®], catalog number: 423790)
- 10. Leaf tissue of C-value standard (*e.g.*, *Nicotiana tabacum* L. 'Xanthi'; 2C = 10.04 pg, Johnston *et al.*, 1999)
- 11. Spores of ferns or lycophytes (kept by dry storage and under < 4 °C)
- 12. PVP-40 (Sigma-Aldrich, catalog number: PVP40)
- 13. 2-mercaptoethanol (Sigma-Aldrich, catalog number: M3148)
- 14. RNaseA solution (10 mg/ml in ddH₂O) (Sigma-Aldrich, catalog number: R5000-100MG)

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- 15. Triton X-100
- 16. Sodium sulfite (Na₂SO₃)
- 17. Tris-HCI (pH 7.5)
- 18. Propidium iodide
- 19. Backmen stock buffer (see Recipes)*
 *Note: LB01 buffer (Doležel et al., 2007) or GPB buffer (Loureiro et al., 2007) can be alternatively used depending on plant material properties.
- 20. PI solution (see Recipes)

Equipment

- 1. Pipette (10, 100, and 1,000 µl)
- 2. Vortex (Scientific Industries, model: Vortex-Genie 2)
- 3. Razors and razor pen
- Flow cytometer (BD, BD Biosciences, model: FACScan)*
 *Note: FACScan with a 15 microW blue argon ion laser of an emission wave length of 488 nm.

Software

1. BD FACSCan system (BD Biosciences, Franklin Lake, NJ, USA)

Procedure

- 1. Prepare buffer for use
 - a. Allocate appropriate amount of Backmen stock buffer to a 50-ml tube based on an estimation of 1-1.5 ml per sample.
 - b. Add 0.04 g PVP-40, 5 µl 2-mercaptoethanol, 1 µl RNase per ml of buffer.
- 2. Extract spore nuclei by bead-vortexing
 - a. For each sample, weigh ~0.007 g spores into a 1.7-ml tube. Green spores that usually have thin spore walls, 4 times the amount of spores are recommended (Kuo *et al.*, 2017)*.
 *Note: For bead-vortexing, the detailed process can be seen in the video supplied in Kuo et al. (2017): <u>http://onlinelibrary.wiley.com/store/10.1111/nph.14291/asset/supinfo/nph14291-sup-0002-VideoS1.mov?v=1&s=dae6f0590d33413f1444bc0add857d285fb73cd6</u>
 - b. Add 16 stainless steel beads into each 1.7-ml tube.
 - c. Add 250 µl of buffer into each 1.7-ml tube.
 - d. Vortex these tubes at a speed of 1,900 rpm for 1 min. For green spores, a speed of 3,200 rpm and a vortex duration of 0.5 min are recommended (Kuo *et al.*, 2017).
 - e. Filter bead-vortexed samples into 2.0-ml tubes through nylon meshes. The size of nylon mesh is selected based on spore sizes to prevent spore being filtered through the mesh.

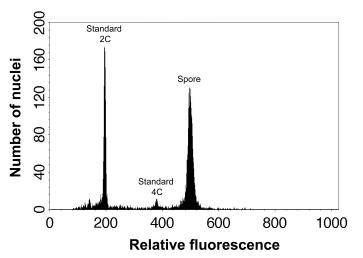
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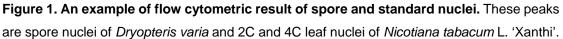
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- f. Add additional buffer to the samples, and ensure each of filtered spore nuclei solutions is greater than 500 μl in volume.
- 3. Extract standard nuclei by chopping leaf tissue
 - a. Add 500 µl of buffer to a glass Petri dish.
 - b. Add a (~400 mm²) piece of young leaf to the Petri dish, and chop it with a razor on ice until most tissue slices are less than 1 mm in size.
 - c. Filter the chopped sample into a 2.0-ml tube through a 30-µm nylon mesh.
 - d. Add additional buffer to the sample, and ensure that the filtered leaf nuclei solution is greater than 500 μ l in volume or more depending on need.
- 4. Staining nuclei solutions
 - a. Mix spore nuclei and standard leaf nuclei solutions into a 500-µl volume in 2.0-ml tubes.
 - b. Add 10 µl Pl solution into each of mixed nuclei solutions.
 - c. Incubate in the dark at 4 °C for 1 h for staining.

Data analysis

- 1. Set up a histogram plot of particle count vs. linear value of relative fluorescence in BD FACSCan system.
- 2. After PI staining, measure nuclear DNA content of the samples in BD FACSCan system, and adjust the fluorescence laser voltage to visualize the nuclei peaks on the histogram plot.
- 3. Measure > 1,300 particles for each peak, and the coefficient variation for each peak should be lower 5% as the quality criteria suggested by Greilhuber *et al.* (2007).
- 4. Genome size of spore nuclei (pg or Mbp) = $\frac{Sporenuclei mean peak position}{Standard 2C leaf nuclei mean peak position}$ x standard 2C value (Figure 1).







<u>Notes</u>

 To dry fertile leaves to release and collect spores, air-drying process under room temperature for 2 to 3 days is recommended. For long-term storage, spore material is better stored in tubes without any solution at a temperature lower than 4 °C.

Recipes

- Backmen stock buffer (Ebihara *et al.*, 2005)
 1.0% Triton X-100
 50 mM Na₂SO₃
 50 mM Tris-HCl (pH 7.5)
 ddH₂O (the solvent)
 Note: Store at 4 °C up to 1 year.
- PI solution
 2.04 mg/ml propidium iodide ddH₂O (the solvent)
 Note: Store in the dark at 4 °C for long-term storage.

Acknowledgments

The bead-vortexing condition to extract spore nuclei of seedless vascular plants is accessed and constructed by Kuo *et al.* (2017). We thank Fay-Wei Li and three anonymous reviewers for providing comments on the draft of this manuscript.

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