

Flow Cytometry Protocol

Bethany Zumwalde and Jessie Pelosi

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REAGENTS:

Woody Plant Buffer* (WPB; Loureiro et al., 2007)

- 0.2M Tris, pH 7.5
- 4mM MgCl₂
- 2mM EDTA
- 86 mM NaCl
- 10 mM Sodium Metabisulfite
- 1% (w/v) PVP-10
- 1% (v/v) Triton X-100

*Final concentrations are given. Buffers should be stored in aliquots at 4-8°C and remain stable for multiple months.

For a final volume of 1000 mL of WPB:

Reagent	Volume/Mass	Final Concentration
1M Tris-HCl, pH 7.5	200mL	0.2M
MgCl ₂ 6H ₂ O	0.81g	4mM
0.5 M EDTA Na ₂ H ₂ O	4mL	2mM
NaCl	5.03g	86mM
Sodium metabisulfite	1.901g	10mM
PVP-10	10g	1%
Triton X-100, pH 7.5	10mL	1% (v/v)
H ₂ O	Top up to 1000mL (~700-746mL)	

P.I. Stock Solution

- 1 mg propidium iodide in 1 mL of DI water

Note: Store as 1 mL aliquots at -20°C. Once thawed, stable at room temperature but should be protected from light.

Other supplies

- RNase A (optional, for fresh material)
- BD Falcon tubes with 5uL mesh cell strainer cap, 5 mL polystyrene round-bottom
 - o Ref 352253, 25/pack: pore size 15-50 um
- Petri dishes on ice
- Single-edged razor blades
- Standards

Standard	2C Genome Size (pg)	Reference
<i>Raphanus sativus</i> cv. Saxa	1.11	Doležel et al. 1992
<i>Solanum lycopersicum</i> cv. Stupicke	1.96	Doležel et al. 1992
<i>Glycine max</i> cv. Polanka	2.50	Doležel et al. 1994
<i>Zea mays</i> CE-777	5.43	Lysák et al. 1998
<i>Pisum sativum</i> cv. Citrad	9.09	Doležel et al. 1998
<i>Secale cereale</i> cv. Dankovske	16.19	Doležel et al. 1998
<i>Vicia faba</i> cv. Inovec	26.90	Doležel et al. 1992
<i>Allium cepa</i> cv. Alice	34.98	Doležel et al. 1998

METHODS

Note: In advance, book the flow cytometry machine via iLab and prepare the buffer and staining solutions (above).

1. Place the petri dish with the samples on ice.
2. Place leaf material in half of a petri dish. The amount of tissue is group-specific and can be rather subjective. Estimate by eyeballing the size of a pinky nail – aim for 250 mg fresh weight. Include a standard tissue to be co-chopped. A smaller amount of the standard relative to the standard may be used to prevent the standard signal from swamping out the sample.
3. Add 1000 uL of ice-cold WPB to the petri dish and chop tissue with a razor blade for 30-60 seconds. Retain the pipette tip from the step to mix the sample later. Note: the amount of chopping may be adjusted depending on the tissue. Finer chopping may result in more noise (more degraded nuclei). Use a new razor blade for each sample. Make sure to not over-chop!
4. Swirl the chopped material in the WPB for 20-30 seconds until a green tint appears in the liquid.
5. Remove the end of the retained pipette tip with the razor blade and use the larger opening to mix the material by pipetting up and down a couple of times (optional).
6. Filter the suspension through the filtered tubes and keep on ice. Transfer 500 uL of the filtrate to a new 1.5 mL Eppendorf tube.
7. Repeat steps 2-6 for each sample.
8. After finishing all samples, add 2.5uL of RNaseA (1mg/mL to 10mg/mL stock) and incubate for 10 minutes. This step is optional.
9. Add 5 uL of P.I. to each tube containing filtrate. Cover tubes with tin foil to protect them from light. Incubate on ice for at least 10 minutes. Dry tissue may require longer incubation.
10. Run on Accuri C6 flow cytometer (see Galbraith, 2009). Note: the BD Accuri C6 cytometer uses a nozzle of 150-200 um. Speed: start slow. If it looks good, can increase to medium or fast. Count to at least 2,000 events (or many more).