Understanding how biodiversity is distributed in nature and how it is impacted by human activities has become crucial for nature conservation. When analyzing biodiversity, it is essential to comprehend the dynamics and geographical distributions of genetic variability in natural populations. Angiosperms (or flowering plants) exhibit different modes of inheritance and patterns of genetic information among individuals. The main mechanisms generating and exploiting genetic variation are based on diverse reproductive strategies used by plants, from imposed cross-pollination to asexual reproduction via seed (apomixis) throughout different intermediate syndromes. Unlike sexually reproducing species, facultative apomicts can produce both sexual and asexual derived seeds through the formation of unreduced female gametophytes.

Until a few decades ago, the use of time-consuming microsectioning techniques was the only way to learn with certainty about these reproductive processes in plants, and by then, most of the different reproductive developmental patterns currently known had already been established. Nonetheless, the arrival of new technologies has allowed us to more simply and quickly unravel the main reproductive syndromes of plants (sexual vs asexual reproduction; see Figure 1) by using special methodologies such as flow cytometry (FC).

Although the first patent on a prototype was issued in 1953, the first flow cytometry device based on fluorescence was developed in 1968 by Wolfgang Göhde from the University of Muenster (Germany) and was first commercialized in 1968–69 by Partec GmbH (Muenster, Germany) through PHYWE AG (Goettingen, Germany). Development of flow cytometry methods

Flow cytometry was initially developed for medical applications, but later it became useful in different fields for such applications as cell counting, sorting, and biomarker detection. Although the first paper on plant nuclei analysis using flow cytometry was published in 1973, the breakthrough in plant DNA analysis by FC came during the late 1980s, when researchers started to persistently apply this technology in plants. Basically, botanists use flow cytometry to measure DNA content in plant nuclei. For a detailed description of the principles and applications of flow cytometric cell analysis in plants, see Ref. 9.

Figure 2a shows an overview of how the flow cytometer lines up stained or targeted particles (sized between 0.2 and 150 µm) by suspending them in a hydrodynamically focused stream of fluid through electronic detection equipment. In plant research, nuclei (i.e., particles) are stained with a fluorescence marker (e.g., DAPI [4’,6-diamidino-2-phenylindole] or propidium iodide). Then, each suspended nucleus is exposed to a beam of light (usually laser or UV light), after which the light path is scattered and perceived by detectors that analyze the fluorescence intensity of every individual particle, providing information about the DNA content of the nucleus. The pooled data of thousands of intact nuclei give information about the DNA content of the respective tissues (see peaks in Figure 1). Since the physical and/or chemical characteristics of up to thousands of particles per second are simultaneously analyzed, the quality and purity of solutions used for sample preparation and sample examination are crucial.

When a low-quality or inappropriate water supply system is used, the presence of contaminating particles that do not belong to the sample can exhibit fluorescence and create “noise,” interfering with the results and causing inaccurate evaluations. Hence, the use of high-quality ultrapure water is mandatory.

Ultrapure water system for production of ultrapure ASTM Class I quality water

Ultrapure water systems that produce ultrapure ASTM Class I quality water are offered by Sartorius (Goettingen, Germany). To evaluate the effects
FLOW CYTOMETRY continued

Figure 2 – a) Schematic overview of a typical flow cytometer technology setup (courtesy of M. Steinberg). Information from every plant nuclei is collected by different detectors, one aligned to the light beam (forward scatter), and several perpendicular to it (side scatters). b) CyFlow device (Partec GmbH) located at the Department of Systematic Botany, Albrecht-von-Haller Institute for Plant Sciences, Georg-August University of Goettingen.

Figure 3 – arium pro VF ultrapure water system (photo courtesy of Sartorius).

Figure 4 – Schematic flow diagram for the arium pro VF ultrapure water system (for enhanced clarity, the valves and their controllers are omitted).

The arium pro VF system (Figure 3) produces ultrapure water from pretreated drinking water by removing all contaminant particles. Ultrapure water production requires continuous recirculation and a constant water flow rate, which is achieved using a pump system with controlled pressure. The conductivity of the water is measured at the feed water inlet and at the downstream port, or product water outlet.

The authors examined the reproductive pathways used by a facultative apomictic angiosperm to produce seeds. This was done by employing the ArUP\textsubscript{H2O} to run the samples and comparing the results to those obtained using the standard sheath fluid solution (0.04% sodium azide, 0.01% detergent) (Partec GmbH).

Materials and methods

Seed samples from hexaploid *Ranunculus carpathica* plants were collected under open pollination conditions. Individual seeds were chopped in a
plastic petri dish with 300 µL of extraction buffer (CyStain UV Precise P, Partec GmbH) and incubated for 10 min at room temperature, after which nuclei were filtered (30 µm mash, CellTric®, Partec GmbH) into a 5-mL plastic tube. Then, 1.2 mL of staining buffer (CyStain UV Precise P) was added, and after 60 sec, samples were analyzed in a flow cytometer (CyFlow Space, Partec GmbH; see Figure 2b) in the blue fluorescence channel.

Setting parameters were adjusted using known diploid (2x) or tetraploid (4x) Ranunculus sp. genotypes. The relative fluorescence of at least 2000 individual nuclei per sample was quantified and represented in a histogram. The relative positions of the peaks were used to determine ploidy levels of embryo and endosperm tissues. Reproductive pathways followed during seed formation were reconstructed according to Ref. 10.

To flow the samples, two different fluids were used: A) sheath flow for flow systems (Partec GmbH), and B) ultrapure water from anrium pro VF. Statistical analyses for trial comparisons were conducted with an Excel spreadsheet (Microsoft Corp., Redmond, WA).

Results
The reproductive pathway was reconstructed for a total of 61 individual seeds. Thirty seeds were flowed using sheath fluid, and 31 seeds were flowed using ArUP$_{H_{2}O}$ (conductivity: 0.055 µS/cm or 18.2 MΩ × cm resistivity compensated to 25 °C) following the standard recommended procedure. An average of 2329 nuclei were measured per sample, representing 73% of the total particles counted, while the other 27% was represented by nuclei in the G$_1$ phase of cell cycle and background signals. Embryo and endosperm mean peak positions were calculated for all analyzed seeds based on an averaged total number of nuclei gathered for every peak.

Relative positions of nuclei on an x-axis histogram plot are based on its relative ploidy according to the tissue of origin (see Figure 1). Hence, a tetraploid nucleus (4n) will have approximately twice as much DNA as a diploid one (2n), emitting a twofold intense fluorescence signal and therefore being placed at twice the relative distance on the x-axis (Figure 5). The reproductive origin of each seed and proportion of sexuality and apomixis were established after index peak determination (Table 1).

Different statistical parameters were taken for each trial and reproductive pathway to compare treatments (Table 2). Absolute values of the coefficient of variation (CV%) were obtained for every sample, and ranges per trial and reproductive pathway, as well as the skewness values of each embryo and endosperm-tissue peak for all apomictic and sexual samples, are presented in Table 2. The Pearson product-moment correlation was calculated for mean values of embryo and endosperm peaks measured per sample and fluid treatment (Table 2).

Discussion
The hexaploid cytotypes of Ranunculus carpaticola are facultative apomictic plants that use a variety of reproductive strategies to generate new individuals in natural populations. Flow cytometric seed screening is a useful, easily applied, and timeless technique for the evaluation of reproductive strategies in seed plants. The two independent treatments employed here (sheath fluid vs ArUP$_{H_{2}O}$) do not show significant differences in peak quality, peak position, or index peak ranges.

The coefficient of variation is a normalized measure used to quantify how dispersed or clustered a set of observed events is compared to a standard probability distribution or frequency distribution (an assumed statistical model). The absolute values of the CV or relative standard deviation (RSD or %RSD) correspond to CV values expressed as a percentage. In this case, though maximum CV% values were slightly higher when using ArUP$_{H_{2}O}$, the difference to the sheath fluid treatment was always smaller than the minimum CV% values, pointing to a similar level of dispersion of counts between treatments.

In addition, even though the bell-shaped peaks tend to have longer right tails (positive skewness), the values are close to zero, indicating the peak had a nearly symmetric shape around the mean. Moreover, the high correlation observed between values of embryo and endosperm peaks indicates that the use of ArUP$_{H_{2}O}$ does not affect the mean relative intensity of measured fluorescence when evaluating nuclei populations, thus preventing distorted correlations and maintaining a dispersion of values within expected ratios for embryo-to-endosperm peaks.
Table 2 – Statistical analyses of flow cytometry histograms from 61 R. carpaticola seeds

<table>
<thead>
<tr>
<th>Seed type</th>
<th>Treatment</th>
<th>Tissue</th>
<th>Mean</th>
<th>SD</th>
<th>Skewness</th>
<th>CV% range</th>
<th>r (p&lt;0.05)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Apo</td>
<td>A</td>
<td>Em</td>
<td>112.75</td>
<td>24.00</td>
<td>2.07</td>
<td>3.74–8.13</td>
<td>0.82</td>
</tr>
<tr>
<td></td>
<td>En</td>
<td>339.16</td>
<td>46.08</td>
<td>0.77</td>
<td>2.8–10.02</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>B</td>
<td>Em</td>
<td>110.00</td>
<td>18.78</td>
<td>1.58</td>
<td>3.74–10.34</td>
<td>0.76</td>
</tr>
<tr>
<td></td>
<td>En</td>
<td>345.78</td>
<td>58.93</td>
<td>−0.07</td>
<td>3.32–11.92</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Sex</td>
<td>A</td>
<td>Em</td>
<td>190.04</td>
<td>43.00</td>
<td>−0.61</td>
<td>4.04–7.48</td>
<td>0.85</td>
</tr>
<tr>
<td></td>
<td>En</td>
<td>297.64</td>
<td>69.07</td>
<td>0.72</td>
<td>3.93–8.22</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>B</td>
<td>Em</td>
<td>170.78</td>
<td>46.97</td>
<td>0.07</td>
<td>3.91–9.69</td>
<td>0.86</td>
</tr>
<tr>
<td></td>
<td>En</td>
<td>299.46</td>
<td>86.94</td>
<td>0.41</td>
<td>4.13–9.23</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

*Apo: apomictic derived seeds; sex: sexually derived seeds; Em: embryo; En: endosperm; Mean: mean peak value; SD: standard deviation; r: Pearson product-moment correlation coefficient.

However, proportions of sexual to asexual seeds were similar in both treatments, as expected when randomly selected samples are used. These results indicate that the determination of reproductive origins is not biased by unproportional shifting of peak positions between treatments.

Overall, comparatively few discrepancies in the obtained results prove the arium system is a time-effective and economical alternative for the analysis of relative ploidies in plant materials. The high quality of arium particle-free water proved to be effective in providing reproducible results in flow cytometric seed screenings. However, since additives like antibiotics and detergents have positive short- and long-term effects on the reliable operation of the flow cytometry system by preventing biofilm formation and improving wetting of the inner surfaces of containers, tubing, valves, and flow cells, the supplier recommends the addition of such substances to a self-made sheath fluid.

To summarize, ArUP

\(_{\text{H}_2\text{O}}\)

can be readily used for flow cytometry of plant cells. Because flow cytometric technology is gaining importance in other applications, such as detection of tumor cells, quantitative estimation and morphological differentiation of cells, cell cycle analysis, DNA–RNA content estimations, and apoptosis measurements, the suitability of ArUP

\(_{\text{H}_2\text{O}}\)

is opening up new opportunities for arium ultrapure water in a variety of applications using the emergent technology of flow cytometry.

References


Dr. Diego H. Hojsgaard is with the Albrecht-von-Haller Institute for Plant Sciences, Department of Systematic Botany, Georg-August University, Untere Karlsrupeule 2, 37073 Goettingen, Germany; tel.: +49 (0) 551/39 5731; fax: +49 (0) 551/39 22329; e-mail: Diego.Hojsgaard@biologie.uni-goettingen.de. Dr. Elmar Herbig is with *Sartorius AG*, Goettingen, Germany. The authors acknowledge Matthias Steinberg (Partec GmbH, Muenster, Germany) for his helpful discussions and suggestions, and for providing the flow cytometry setup image (Figure 2a). The authors also thank Prof. Dr. Elvira Hörandl and Dr. Simone Klatt (Department of Systematic Botany, Georg-August University, Goettingen, Germany) for their critical reading of the manuscript.