## Advanced Expression Vector Systems: New Weapons for Plant Research and Biotechnology

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Scientific discoveries often coincide with the development of new and robust methodologies. Modern plant biology and biotechnology are of no exception to this rule, especially when it comes to production of new vector systems for gene expression. Thus, for example, the progress in plant genetic engineering could not have been as productive as it is today without the development of small, easy-to-manipulate, and simple-to-use Agrobacterium binary vectors (e.g. Komari et al., 2006; Komori et al., 2007), and studies of protein subcellular localization in plant cells have been greatly simplified and advanced with the introduction of vectors that express GFP fusions (Goodin et al., 2007). Indeed, the ability to transiently and stably express foreign genes, to genetically interfere with the expression of native genes, and to functionally study the expression, interaction, localization, and modification of proteins in cells, tissues, and whole plants are fundamental to modern plant basic research and biotechnology.

More than two decades had passed since the introduction of the first generation of plant transformation binary vectors (e.g. Bevan, 1984). Although revolutionary at their time, these vectors were rather simply designed, lacking cloning and expression versatility, and offered very little flexibility for their manipulation for specific research or application purposes. Vector technology has evolved throughout the years, and during this time plant transformation vectors have been a subject of constant improvements (e.g. Becker et al., 1992; Datla et al., 1992; Hajdukiewicz et al., 1994). Newer generations of plant transformation vectors provided plant biologists and biotechnologists with improved strategies for cloning and delivering their genes of interest into plant cells, typically using Agrobacterium as vehicle for the transformation process. Some of these vectors were developed as families of plasmids, and others represented single constructs designed for specific purposes. For example, the pCB mini-binary vector series of plasmids (Xiang et al., 1999) provided an excellent alternative for the relatively large first-generation binary plasmids, whereas the pBI121 plasmid—only recently sequenced (Chen et al., 2003) but no longer available commercially was specifically designed to foster the use of the *GUS* gene as a reporter in genetic transformation experiments.

More recently, we have witnessed an impressive increase in the "introduction" of new and novel vectors suitable for performing various tasks for plant research and biotechnology (Fig. 1). These days, it seems that one can find a plasmid for every task, including such relatively unique applications as activation tagging (e.g. the pSKI015 and pSKI074 binary vectors; Weigel et al., 2000) or dexamethasone-inducible expression (e.g. the pOp/LhGR transcription activation system; Samalova et al., 2005). Also, vectors have been constructed to allow plant biologists to take advantage of radically new cloning methodologies, such as recombinase-mediated gene cloning (Gateway; e.g. the pMDC plasmid collection; Curtis and Grossniklaus, 2003), or of new approaches to modulate gene expression, such as RNA interference-mediated gene silencing (Meyer et al., 2004) and virus-induced gene silencing (Burch-Smith et al., 2006) for knocking out/down gene expression, and the use of viral RNA silencing suppressors to enhance expression of genes of interest (Voinnet et al., 2003). Overall, it appears that virtually every new gene expression technology developed for non-plant systems very quickly finds its application in plant biology via new vector systems. Some recent examples of such vector systems are those that introduce into plant biology the use of the bimolecular florescent complementation assay for protein-protein interaction (Bracha-Drori et al., 2004; Walter et al., 2004; Citovsky et al., 2006), C- and N-terminal protein tagging with various autofluorescent markers (Tzfira et al., 2005; Chakrabarty et al., 2007), CRE/loxP recombination to produce single-copy T-DNA inserts (De Buck et al., 2007), and many others. Besides adapting novel technologies for studies of gene expression and protein interactions, new vector systems are being produced to utilize transgenic technologies in an ever-expanding range of plant species, such as forest trees and transformation-recalcitrant crops (e.g. Meyer et al., 2004; Coutu et al., 2007). Furthermore, vectors for systemic gene expression without permanent genetic modification of the plant are

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Figure 1. From vectors to applications to cellular functions. Introduction of genetic information into target plant cells and acquisition of new data as a result of transgene expression may require a network of modular vectors, flexible gene cloning and expression systems, and specialized plasmids that result in different modes of transgene expression. Modular vectors may represent a starting point for assembly of custom-made expression vectors, multigene expression vectors, and other types of plant transformation vectors. These vectors in turn provide the users with the abilities to overexpress and down-regulate genes, as well as with the capacity for specific, and often unique, applications, useful for obtaining novel traits and functional data, protein imaging in living plant cells, and generating transgenic plants for plant research and biotechnology.

being developed based on different plant viruses (e.g. Gleba et al., 2005; Marillonnet et al., 2005).

It is difficult to overestimate the effect a truly versatile yet simple-to-use expression vector system can have on many fields of plant research and biotechnology. As more and more vectors and vector systems for gene expression in plants become available, it also becomes essential to make their existence and the scope of their use known to the diverse community of plant researchers, basic and applied. This Focus Issue is a small step in this direction. It presents a collection of original articles describing the development of new vector systems useful for plant research and biotechnology, as well as a compilation of short review articles that highlight some of the major developments in vector-assisted plant research technologies. To name a few, the reader will find papers describing an extensive collection of MultiSite Gateway-based plant expression vectors (Karimi et al., 2007), updates on the use of bimolecular florescent complementation for analyses of protein-protein interactions in living plant cells (Ohad et al., 2007) and on the introduction of multiple genes into plant cells (Dafny-Yelin and Tzfira, 2007), a guide to vectors for chloroplast transformation (Lutz et al., 2007), and descriptions of a yeast-plant coupled system for detection of functional nuclear localization signals (Zaltsman et al., 2007), a virus-induced gene silencing system for reverse genetics of floral scent (Spitzer et al., 2007), a system of transformation vectors with the superpromoter (Lee et al., 2007), and a new cloning strategy for recombinase-mediated cassette exchange (Louwerse et al., 2007).

## ACKNOWLEDGMENTS

We thank all the authors who chose to publish their findings and updates in this *Focus Issue* and all the reviewers whose comments and constructive



criticisms made the publication of this collection of manuscripts possible. We hope that the readers will find the reports and updates not only interesting, but also immediately useful for enhancing their own work in plant basic research and biotechnology. The work in our labs was supported by grants from the National Institutes of Health, the National Science Foundation, the U.S. Department of Agriculture, the United States-Israel Binational Agricultural Research and Development Fund, and the United States-Israel Binational Science Foundation to V.C., and by the Biotechnology Research and Development Corporation and University of Michigan start-up funds to T.T.

Received October 25, 2007; accepted October 26, 2007; published December 6, 2007.

## LITERATURE CITED

- Becker D, Kemper E, Schell J, Masterson R (1992) New plant binary vectors with selectable markers located proximal to the left T-DNA border. Plant Mol Biol 20: 1195–1197
- Bevan MW (1984) Binary Agrobacterium vectors for plant transformation. Nucleic Acids Res 12: 1811–1821
- Bracha-Drori K, Shichrur K, Katz A, Oliva M, Angelovici R, Yalovsky S, Ohad N (2004) Detection of protein-protein interactions in plants using bimolecular fluorescence complementation. Plant J 40: 419–427
- Burch-Smith TM, Schiff M, Liu Y, Dinesh-Kumar SP (2006) Efficient virusinduced gene silencing in Arabidopsis. Plant Physiol 142: 21–27
- Chakrabarty R, Banerjee R, Chung SM, Farman M, Citovsky V, Hogenhout SA, Tzfira T, Goodin MM (2007) pSITE vectors for stable integration or transient expression of autofluorescent protein fusions in plants: probing *Nicotiana benthamiana*-virus interactions. Mol Plant Microbe Interact 20: 740–750
- Chen PY, Wang CK, Soong SC, To KY (2003) Complete sequence of the binary vector pBI121 and its application in cloning T-DNA insertion from transgenic plants. Mol Breed 11: 287–293
- Citovsky V, Lee LY, Vyas S, Glick E, Chen MH, Vainstein A, Gafni Y, Gelvin SB, Tzfira T (2006) Subcellular localization of interacting proteins by bimolecular fluorescence complementation *in planta*. J Mol Biol 362: 1120–1131
- Coutu C, Brandle J, Brown D, Brown K, Miki B, Simmonds J, Hegedus DD (2007) pORE: a modular binary vector series suited for both monocot and dicot plant transformation. Transgenic Res 16: 771–781
- Curtis MD, Grossniklaus U (2003) A Gateway cloning vector set for highthroughput functional analysis of genes in planta. Plant Physiol 133: 462–469

- Dafny-Yelin M, Tzfira T (2007) Delivery of multiple transgenes to plant cells. Plant Physiol 145: 1118–1128
- Datla RS, Hammerlindl JK, Panchuk B, Pelcher LE, Keller W (1992) Modified binary plant transformation vectors with the wild-type gene encoding NPTII. Gene 122: 383–384
- De Buck S, Peck I, De Wilde C, Marjanac G, Nolf J, De Paepe A, Depicker A (2007) Generation of single-copy T-DNA transformants in Arabidopsis by the CRE/*loxP* recombination-mediated resolution system. Plant Physiol **145**: 1171–1182
- Gleba Y, Klimyuk V, Marillonnet S (2005) Magnifection—a new platform for expressing recombinant vaccines in plants. Vaccine 23: 2042–2048
- Goodin MM, Chakrabarty R, Banerjee R, Yelton S, DeBolt S (2007) New gateways to discovery. Plant Physiol 145: 1100–1109
- Hajdukiewicz P, Svab Z, Maliga P (1994) The small, versatile pPZP family of Agrobacterium binary vectors for plant transformation. Plant Mol Biol 25: 989–994
- Karimi M, Bleys A, Vanderhaeghen R, Hilson P (2007) Building blocks for plant gene assembly. Plant Physiol 145: 1183–1191
- Komari T, Takakura Y, Ueki J, Kato N, Ishida Y, Hiei Y (2006) Binary vectors and super-binary vectors. Methods Mol Biol 343: 15–42
- Komori T, Imayama T, Kato N, Ishida Y, Ueki J, Komari T (2007) Current status of binary vectors and superbinary vectors. Plant Physiol 145: 1155–1160
- Lee L-Y, Kononov ME, Bassuner B, Frame BR, Wang K, Gelvin SB (2007) Novel plant transformation vectors containing the superpromoter. Plant Physiol 145: 1294–1300
- Louwerse JD, van Lier MCM, van der Steen DM, de Vlaam CMT, Hooykaas PJJ, Vergunst AC (2007) Stable recombinase-mediated cassette exchange in Arabidopsis using *Agrobacterium tumefaciens*. Plant Physiol **145**: 1282–1293
- Lutz KA, Azhagiri AK, Tungsuchat-Huang T, Maliga P (2007) A guide to choosing vectors for transformation of the plastid genome of higher plants. Plant Physiol **145**: 1201–1210
- Marillonnet S, Thoeringer C, Kandzia R, Klimyuk V, Gleba Y (2005) Systemic Agrobacterium tumefaciens-mediated transfection of viral replicons for efficient transient expression in plants. Nat Biotechnol 23: 718–723

- Meyer S, Nowak K, Sharma VK, Schulze J, Mendel RR, Hansch R (2004) Vectors for RNAi technology in poplar. Plant Biol (Stuttg) 6: 100–103
- **Ohad N, Shichrur K, Yalovsky S** (2007) The analysis of protein-protein interactions in plants by bimolecular fluorescence complementation. Plant Physiol **145**: 1090–1099
- Samalova M, Brzobohaty B, Moore I (2005) pOp6/LhGR: a stringently regulated and highly responsive dexamethasone-inducible gene expression system for tobacco. Plant J 41: 919–935
- Spitzer B, Ben Zvi MM, Ovadis M, Marhevka E, Barkai O, Edelbaum O, Marton I, Masci T, Alon M, Morin S, et al (2007) Reverse genetics of floral scent: application of tobacco rattle virus-based gene silencing in petunia. Plant Physiol 145: 1241–1250
- Tzfira T, Tian GW, Lacroix B, Vyas S, Li J, Leitner-Dagan Y, Krichevsky A, Taylor T, Vainstein A, Citovsky V (2005) pSAT vectors: a modular series of plasmids for fluorescent protein tagging and expression of multiple genes in plants. Plant Mol Biol 57: 503–516
- Voinnet O, Rivas S, Mestre P, Baulcombe DC (2003) An enhanced transient expression system in plants based on suppression of gene silencing by the p19 protein of tomato bushy stunt virus. Plant J **33**: 949–956
- Walter M, Chaban C, Schütze K, Batistic O, Weckermann K, Näke C, Blazevic D, Grefen C, Schumacher K, Oecking C, et al (2004) Visualization of protein interactions in living plant cells using bimolecular fluorescence complementation. Plant J 40: 428–438
- Weigel D, Ahn JH, Blazquez MA, Borevitz JO, Christensen SK, Fankhauser C, Ferrandiz C, Kardailsky I, Malancharuvil EJ, Neff MM, et al (2000) Activation tagging in Arabidopsis. Plant Physiol 122: 1003–1013
- Xiang C, Han P, Lutziger I, Wang K, Oliver DJ (1999) A mini binary vector series for plant transformation. Plant Mol Biol 40: 711–717
- Zaltsman A, Yi B-Y, Krichevsky A, Gafni Y, Citovsky V (2007) Yeast-plant coupled vector system for identification of nuclear proteins. Plant Physiol 145: 1264–1271