Advancement Method and Process in Plant Synthetic Biology

Rayees Afzal Mir, Syed Aasif Hussain Andrabi

Glocal School of agricultural sciences Glocal University, Saharanpur (UP). India 247121 raies.afzal@gmail.com

Abstract

Our Research Paper "Advancement Method and Process in Plant Synthetic Biology" is an Engineered science is an arising field joining researchers from all disciplines determined to plan or updating organic cycles. At first, manufactured science forward leaps came from microbial science, science, physical science, software engineering, materials science, arithmetic, and designing disciplines. A progress to multicellular frameworks is the following consistent advance for engineered researcher and plants will give an optimal stage to this new period of examination. This gathering report features a portion of the intriguing plant manufactured science ventures, and devices and assets, introduced and examined at the 2013 GARNet studio on plant engineered science.

Key words: Methods, plant science, resources, science policy, synthetic biology, tools.

Introduction

Engineered science is another methodology in logical exploration however, in numerous ways, it is the normal movement of the manner in which present day science has been embraced for the beyond twenty years. It is as a matter of first importance driven by the explorative and innovative nature of essential examination, and has the potential for colossal business sway. Also, manufactured science is innately between disciplinary, cutting across the logical fields of science, physical science, science, math, designing, sociology, and software engineering, just as incorporating plan, financial aspects, and showcasing. These two fundamental standards of manufactured

biology are in keeping both with the ebb and flow need to boost gets back from public financing, and with the cross-cutting nature of organic examination since the beginning of the genomics era.

Synthetic science hit the UK government's radar in 2012. The Department of Business Innovation and Skills (BIS) actuated a guide for UK manufactured science (UK Synthetic Biology Roadmap Coordination Group, 2012), a £20 million speculation responsibility around here, and the Synthetic Biology Leadership Council to evaluate progress, update proposals, and shape needs. In 2013, this venture converted into two freedoms for synergistic manufactured science projects: the Research Councils UK (RCUK) call for proposition for Multi-disciplinary Research Centers and the ERASynBio call for transnational examination projects across Europe and the US; and the UK empowering devices and advancements improvement reserve from the UK government financing bodies: the Technology Strategy Board (TSB), the Biotechnology and Biological Sciences ResearchChamber (BBSRC), and the Engineering and Physical Sciences Research Council (EPSRC). Also, TSB, BBSRC, and EPSRC granted £10 million to another Innovation and Knowledge Center based at Imperial College London. This Center was opened in July 2013 to work with the change between manufactured science research and modern application.

In this specific circumstance, GARNet, the Arabidopsis research organization, held the second UK studio on engineered science for the plant sciences at the University of Nottingham on 21–22 May 2013. This

followed on from the fourth New Phytologist studio in May 2012 (Osbourn et al., 2012), which additionally expected to acquaint engineered science with UK plant researchers.

Synthetic biology

The most widely recognized meaning of engineered science is the plan and development of new organic parts, gadgets, and frameworks and the re-plan of existing natural frameworks (Arkin et al., 2009). For some scientists, this implies reconstructing frameworks from the DNA level up, while for different gatherings, including the UK's Synthetic Biology Leadership Council, it should likewise incorporate a business end-point (Arkin et al., 2009). According to Garnet's viewpoint, it is key that crucial examination supports all of manufactured science and that imaginative items are an outcome of this.At when Western economies are battling, engineered science is seen as an expected distinct advantage for advancement and industry. Subsequently, numerous engineered science. funders overall intently screen the bioethics of manufactured science rules that should be viewed as when drafting recommendations and stringently clung to all through the undertaking (The National Science Federation, 2012; The European Group on Ethics in Science and New Technologies to the European Commission, 2009).

As displayed at the GARNet studio, projects inside the circle of 'manufactured science' fluctuate extraordinarily in scope. They start at the sub-atomic scale, for instance, the exquisite anew protein development unit of looped curl peptides (Motiveless and Wolfson, 2009; Fletcher et al., 2012) that can be utilized to make a developing number of constructions including self-collecting confines (Fletcher et al., 2013). At the supramolecular levels, DNA and the plan and age of negligible manufactured genomes are significant objectives in engineered science. The driven Yeast 2.0 task intends to answer a portion of the essential inquiries basic genomics, like the job of introns and the significance of genome association (Diamond et al., 2011), and to give an anticipated host cell to future manufactured biologists.

Methods created by engineered science, specifically cloning and genome altering methods, have extended the potential outcomes of customary hereditary and metabolic designing (Neumann and Neumann-Stibitz, 2010). The development of high worth results of intricacies not already conceivable is presently a reality. Artemisinin is an enemy of malarial that normally happens in the plant Artemisia annua however, until this year, has been inadmissible for large scale manufacturing. Nonetheless, the biosynthetic pathway of its forerunnerartemisinic acid has been re-constructed in *Saccharomyces cerevisiae* and is now close to being marketed (Paddon *et al.*, 2013). Further, as described below, nitrogen-fixing wheat is a probable goal rather than an unattainable dream.

Plant synthetic biology

Engineered science is probably going to be an important methodology in the plant scientist's tool stash to assist with meeting the test of giving food, energy, and different materials from restricted normal assets. Plant manufactured science is critical to this application. It is additionally the following consistent advance in the movement of engineered science after the atom, genome, and separated cell level methodologies that have been tested to date: a change to multicellular frameworks. As featured by studio participants in the conversation gatherings, plants are the conspicuous decision for this 'next section' of manufactured science. Plant manufactured researcher have a fantastic information base to work with, and their subjects are sessile, self-fixing, and not encompassed by the very moral issues that are frequently experienced in creature research.

One plant engineered science project that embodies the capability of plants as stages for manufactured science is the nitrogen-fixing cereals project drove by Giles Holroyd at the John Innes Center. Nitrogen is a basic restricting component for plant development and improvement, however utilization of nitrogen composts is costly and can have negative ecological effects. Holroyd and partners have recently described the pathways that permit a few plants, like vegetables, to fix their nitrogen by means of a harmonious relationship with rhizobium microorganisms (Capoen et al., 2011; Oldroyd et al., 2011; Xie et al., 2012). They are currently endeavoring to integrate these pathways in wheat to foster a 'self-treating' cereal.

An illustration of an effectively executed engineered pathway in plants is the sign transduction framework worked by June Medford and her gathering at Colorado State University. The 'sentinel' plant containing the total manufactured pathway goes about as a biosensor that de-colors when it distinguishes a particular molecule(s) like TNT. Improvement of this sentinel included combining and upgrading a bacterial-based sign transduction pathway in plant cells. Significantly, utilization of a prokaryotic system has created a manufactured pathway that is secluded from any endogenous movement, limiting cross-talk inside the plant cell climate and expanding signal identification (Antunes et al., 2006, 2009, 2011).

The two tasks depicted above fit what may be viewed as objective situated engineered science: explicitly updating a plant/crop for a specific point. Other plant researchers arethinking about the master plan: investigating and measuring specific parts of plants to give apparatuses and assets to the more extensive plant science local area to attempt engineered biology.

One potential symmetrical 'tool compartment' for plant manufactured science might come from as of late recognized operon-like groups (Field et al., 2011; Mugford et al., 2013). In microscopic organisms, novel anti-microbials have been biosynthesized by distinguishing proof, portrayal and revamp of quality bunches (Gottelt et al., 2010; Gomez-Escribano et al., 2012). The revelation that some plant optional metabolite qualities, for example, antimicrobial terpenoids in oat and triterpenes in the vegetable Lotus japonicus, are encoded by quality bunches (Krokida et al., 2013; Mugford et al., 2013) presents the chance of rearranging qualities inside a group to create novel mixtures in plants similarly as in bacteria.

Despite the upsides of plant models for manufactured science and their possible uses as illustrated above, it stays a reality that plants are more convoluted than existing engineered science models like Saccharomyces cerevisiae. They have complex flagging pathways, numerous organs, and frequently have extremely enormous genomes. To attempt to beat these restrictions and push the limits of engineered science in plants, studio speaker Jim Hasselhoff and a gathering of global teammates propose the liverwort Merchantmanpolymorph as another model for plant manufactured science, because of its smoothed out genome design (Ohyama et al., 2009) containing less hereditary repetition (Sasaki et al., 2007) than Arabidopsis thaliana, its formative effortlessness and simple development in suspension, agar or on soil. If 'explorative' plant engineered science is to turn into a set up field in the UK, reception of new frameworks like Merchantman might be fundamental.

Tools for synthetic biology in plants

To thrive, plant manufactured science will require a set-up of related instruments and assets, some of which will be plant-explicit, however the larger part are probably going to be adjusted from different fields. Truth be told, one of the basic standards of manufactured science is that a natural 'part' ought to have the option to squeeze into any framework and act in an anticipated manner. In spite of the fact that building a plant or plant pathways from symmetrical parts—an idea regularly compared to a kid's Lego unit—isn't as of now practical, instruments that can create engineered 'parts' in plants do exist and a couple of models are laid out below.

Genome assembly

Gibson Isothermal Assembly, introduced at the GARNet studio by Jim Ajioka (University of Cambridge), is an incredible asset that can be utilized to gather a few hundred kilogausses of DNA into a solitary particle. DNA sections are ready for get together by the ligation of explicit groupings to each end. For this situation, the particular shades that outline the DNA section are integral to the shade of another DNA piece and during gathering, the DNA sorts fit out like a jigsaw puzzle. As verification of the effectiveness and limit of the strategy, Gibson et al. (2009) developed a DNA atom as extensive as 583 kb and cloned items up to 300 kb long in Escherichia coli.

Sylvestre Marillonet's (Leibniz Institute of Plant Biochemistry) Golden Gate modular cloning system (Engler*et* al., 2008) empowers profoundly effective get together of multigene DNA builds in a solitary cylinder utilizing an ordinary benchtop thermocycler. It depends on Type IIS limitation endonucleases, generally BsaI, which are focused on to the right area by BsaI locales that flank the quality or section of interest. The proteins perceive the BsaI destinations and make a slice close to them, leaving a particular four base pair overhang on one or the other strand, as displayed in Fig. 1A. It is feasible for enormous quantities of various parts to be produced and ligated in a solitary cycle; everything necessary is for the BsaI destinations to be planned accurately (Fig. 1B). With legitimate plan, a ligation between two sticky ends left from BsaI absorption will result in a 'scar less' clone.

The Modular Cloning (MoClo; Weber et al., 2011) transformation of Golden Gate is a various leveled cloning framework that empowers any multigene develop to be cloned. It is a unit like technique containing various modules and vectors to work with quality stacking. This takes out the requirement for the perplexing groundwork plan essential for cloning the BsaI locales and explicit flanking successions on to each DNA part in anticipation of gathering. A subsequent variation, GoldenBraid (SarrionPartiones et al., 2011), works on the strategy even further by u2019tilizing just four objective plasmids. This restricts the quantity of parts it is feasible to collect in one run, yet as every objective plasmid has two distinctive Type IIS limitation destinations, they can be re-organized on different occasions (Sarrion-Perdigones et al., 2011).

Gibson Assembly and the Golden Gate cloning strategies, alongside new, quick DNA blend stages, have reformed sub-atomic science, which had recently been restricted by limitation chemicals and ligases requesting particularity. Five years prior, building a whole engineered science pathway would have been viewed as inconceivable because of the time scales included, yet presently it is a possible option.

Genome altering tools

Transcription activator-like effectors (TALEs; and the connected TALE-nucleases), introduced at the studio by coinventor Sebastian Schornack, and bunched routinely interspaced short palindromic rehashes (CRISPR) advancements are

Fig. 1. (A) Schematic diagram of Type II constraint protein activity, with BsaI for example. Unidirectional BsaI areas are arranged near express groupings, S1 and S2. After absorption, the shades are comparing to one another and the BsaI objections are extricated. Right when the parts are ligated together, there is no scar. There is no limitation to the proportion of unequivocal progressions used to top and tail the 'parts,' so this technique can be used to manufacture works with many parts, as in (B), accepting the BsaI regions are arranged correctly.

genome modifying gadgets made for and by designed researchers. They perform site-unequivocal, twofold deserted DNA cleavage to take out characteristics, forsaking no new DNA, and can fortify homologous recombination or non-homologous end joining.

TALEs are used by specific sorts of the pathogenic proteobacteria family Xanthomonas to turn on express host characteristics (Schornack et al., 2008). They incorporate nuclear imprisonment hails, a transcriptional inception space, and a movement a few repeats. Two variable stores in each repeat conclude the particular

DNA grouping the TALE will target (Boch et al., 2009). Story nucleases (TALENs) are TALEs intertwined to the synergist area of a fokl nuclease (Christian et al., 2010). They cut DNA strands at profoundly explicit areas and can perform erasures (Fig. 2A) or then again, assuming a contributor format is given, trigger consistent inclusion by homologous recombination. In 2012, TALEN innovation was utilized to eliminate a segment of a TALE restricting site in a rice sucrose efflux carrier captured by X. oryzae during disease. This unpretentious hereditary adjustment, which presented no unfamiliar DNA by any means, delivered rice plants impervious to X. oryzae (Li et al., 2012).

CRISPR/Cas frameworks normally happen in microscopic organisms and archaea, giving guard memory against attacking phages: CRISPR-related endonuclease Cas9 is directed to target genomic successions, generally key contamination qualities, by two RNA atoms. In programmable CRISPR/Cas genome altering innovation, explicit arrangements on a CRISPR RNA particle are intended to focus on a particular area on the genome (Jinek et al., 2012; Fig. 2B). Utilization of the CRISPR/Cas framework for genome altering was at first shown in prokaryotes (Gasiunas et al., 2012; Jiang et al., 2013) and creature cells (Hwang et al., 2012; Chang et al., 2013; Cong et al., 2013). Since the GARNet studio on Synthetic Biology, CRISPR/Cas innovation has likewise been exhibited in plants (Li et al., 2013; Nekrasov et al., 2013; Shan et al., 2013).

An engineered science articulation system

A incredible articulation framework for manufactured science applications is the CPMV-HT transient articulation framework created at the John Innes Center by George Lomonossoff. The framework is itself an accomplishment of manufactured science designing: the creators changed cow-pea mosaic infection (CPMV) RNA-2 to transform it into a non-viral articulation vector that conveys as high an articulation as viral frameworks (Sainsbury and Lomonossoff, 2008). In contrast to customary change, which requires a timescale of weeks before the ideal protein is communicated in a youthful plant, very significant level articulation is seen only five days after invasion with the vector. This framework empowers undeniable level transient articulation of unfamiliar proteins in tobacco (Vardakou et al., 2012) and, assuming that it tends to be applied to different species, may turn into a helpful device for manufactured science. For instance, specialists have utilized this way to deal with produce void CPMV-like vessels equipped for conveying weighty metals or other unfamiliar particles like medications, introducing energizing prospects in drug conveyance (Aljabali et al., 2010).

Synthetic biology resources

As mentioned in several presentations at the GARNet workshop, many synthetic biology techniques including Gibson Assembly, Brilliant Gate Cloning, and TALE innovation, are available for scholarly use; widely portrayed in papers, sites, and online conventions. The open ethos of the worldwide manufactured science local area might have been cultivated, and was unquestionably fortified, by the iGEM Foundation and its related Registry of Parts. iGEM (global Genetically Engineered Machine) began in 2003 as a yearly rivalry for college understudies. Groups work on a manufactured science project and the subsequent items or apparatuses are added to the Parts Registry. Passages in ensuing years expand on existing work utilizing the persistently extending apparatuses in the iGEM Registry as opposed to beginning from scratching. The opposition presently has divisions for schools and business visionaries just as students. Anybody can utilize the Registry to store and demand DNA parts including advertisers, preliminaries, ribosome restricting destinations, and protein spaces, for the expense of postage. The Registry in this manner gives great assets to those working in the manufactured science circle. Truth be told, as indicated by an overview of engineered researcher (Kahl and Endy, 2013), iGEM's Registry of Standard Biological Parts is the most broadly utilized manufactured science storehouse. Addgene, additionally available to establish researchers, is the third most

generally utilized. Like the Parts Registry it is local area based, yet little handling charges guarantee the quality and dependability of its parts.

As well as the assets preparing the manufactured science local area with actual organic parts, there are numerous sites giving open source programming. One model is the Infobiotics Workbench, the co-maker of which Natalio

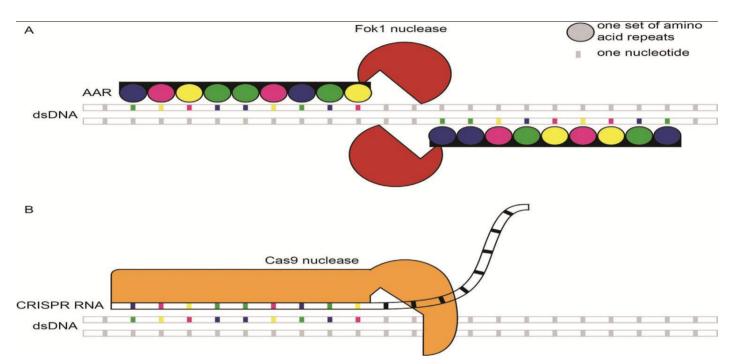


Fig. 2. Schematic diagram comparing (A) TALEN and (B) CRISPR/Cas9 genome editing. TALEN specificity is determined by a series of 36 amino acid repeats (AAR), each containing two hypervariable amino acids in positions 13 and 14. The nuclease makes a single-stranded break, necessitating the design and synthesis of two TALENs for one double-stranded break. CRISPR RNA, including a specific sequence, guides the Cas9 nuclease to the right target, where it makes a double-stranded break.

Krasnagor presented at the GARNet conference; a framework for designing, implementing, and analysing *in silico* experiments, including model checking (Blakes*et al.*, 2011).

Software engineers are also building programs for specific synthetic biology applications. Information extracted from time-lapse images of *Coleochaetescutata* cells was used to build CellModeller, a tool to model cell division in plant tissues and bacterial provinces (Dupuy et al., 2010). Close by functional trials, it is currently being tried as a model for the impact of unfamiliar record factors on plant morphogenesis.

In request to normalize the many bits of programming produced for and by engineered researcher, a global group including Guy-Bart Stan (Imperial College London) fostered the manufactured science open language (SBOL). SBOL is an open-source standard for in silico portrayal of hereditary plans—not a programming language and not language-explicit, but rather a graphical portrayal of a plan

Conclusions

Manufactured science is a set up approach in logical examination, especially in the microbial and substance fields. There are remarkable lead plant-based manufactured science projects and, as the current apparatuses and assets from different circles are either straightforwardly applied, or adjusted for use in plant science, it is

normal that the plant manufactured science local area will develop and take advantage of the capability of this methodology for understanding plant frameworks and designing organic solutions.

Like the field of frameworks science that went previously, it will set aside effort for engineered science to turn into an 'regular' research approach in plant science. Financing openings for specialists to investigate and construct assets around here (sandpits), close by a library of plant parts and the advancement and age of guidelines to empower casual or formal sharing of parts, will all assist with building a durable plant manufactured science community.

We would firmly urge plant researchers to plunge their toe into the pool of engineered science and exploit the expanded subsidizing designated to this space. There are numerous chances to engage in engineered science; for instance, UK college groups routinely enter the iGEM rivalry and have won acknowledgment previously. This is a demonstrated method of finding more with regards to the engineered science mentality and making important associations for future work.

The 'manufactured plants' being planned and worked by the studio speakers and referenced in this article are only the beginning: later on, plant manufactured researcher might accomplish more prominent things even than plant sentinels and mineral-fixing crops. The expected effect of manufactured science in plants is certainly tremendous and ought not be disregarded.

References

- 1. Aljabali AAA, Sainsbury F, Lomonossoff GP, Evans DJ. 2010. Cowpea mosaic virus unmodified empty viruslike particles loaded with metal and metal oxide. *Small*6, 818–821.
- Antunes MS, Ha SB, Tewari-Singh N, *et al.* 2006. A synthetic de-greening gene circuit provides a reporting system that is remotely detectable and has a re-set capacity. *Plant Biotechnology Journal*4, 605–622. Antunes MS, Morey KJ, Smith JJ, *et al.* 2011. Programmable ligand detection system in plants through a synthetic signal transduction pathway. *PLoS One*6, e16292.
- 3. Antunes MS, Morey KJ, Tewari-Singh N, *et al.* 2009. Engineering key components in a synthetic eukaryotic signal transduction pathway. *Molecular Systems Biology***5**, 270.
- 4. Arkin A, Berry D, Church G, et al. 2009. What's in a name? Nature Biotechnology27, 1071–1073.
- 5. Blakes J, Twycross J, Romero-Campero FJ, Krasnogor N. 2011. The Infobiotics Workbench: an integrated *in silico* modelling platform for systems and synthetic biology. *Bioinformatics*27, 3323–3324.
- Boch J, Scholze H, Schornack S, Landgraf A, Hahn S, Kay S, Lahaye T, Nickstadt A, Bonas U. 2009. Breaking the code of DNA binding specificity of TAL-Type III effectors. *Science*326, 1509– 1512.
- 7. Capoen W, Sun J, Wysham D, *et al.* 2011. Nuclear membranes control symbiotic calcium signaling of legumes. *Proceedings of the National Academy of Sciences, USA*108, 14348–14353.
- 8. Chang N, Sun C, Gao L, Zhu D, Xu X, Xhu X, Xiong J-W, Xi JJ. 2013.
- 9. Genome editing with RNA-guided Cas9 nuclease in Zebrafish embryos. *Cell Research*23, 465–472.
- Christian M, Cermak T, Doyle EL, Schmidt C, Zhang F, Hummel A, Bogdanove AJ, Voytas DF. 2010. Targeting DNA double-strand breaks with TAL effector nucleases. *Genetics*186, 757–761. Cong L, Ran FA, Cox D, *et al.* 2013. Multiplex genome engineering using CRISPR-Cas systems. *Science*339, 819–823.
- 11. **Dupuy L, Mackenzie J, Haseloff J.** 2010. Coordination of plant cell division and expansion in a simple morphogenetic system. *Proceedings of the National Academy of Sciences, USA***107,** 2711–2716.
- 12. Dymond JS, Richardson SM, Coombes CE, *et al.* 2011. Synthetic chromosome arms function in yeast and generate phenotypic diversity by design. *Nature***477**, 471–476.

- 13. Engler C, Kandzia R, Marillonnet S. 2008. A one pot, one step, precision cloning method with high throughput capability. *PLoS One*3, e3647.
- 14. Field B, Fiston-Lavier A-S, Kemen A, Geisler K, Quesneville H, Osbourn AE. 2011. Formation of plant metabolic gene clusters within dynamic chromosomal regions. *Proceedings of the National Academy of Sciences, USA*108, 16116–16121.
- 15. Fletcher JM, Boyle AL, Bruning M, et al. 2012. A basis set of *de novo* coiled-coil peptide oligomers for rational protein design and synthetic biology. ACS Synthetic Biology6, 240–250.
- 16. Fletcher JM, Harniman RL, Barnes FRH, et al. 2013. Self-assembling cages from coiled-coil peptide modules. *Science* **3**, 595–599.
- 17. Gasiunas G, Barrangou R, Horvath P, Siksnys V. 2012. Cas9-crRNA ribonucleoprotein complex mediates specific DNA cleavage for adaptive immunity in bacteria. *Proceedings of the National Academy of Sciences, USA*109, E2579–E2586.
- 18. Gibson DG, Young L, Chuang R-Y, Venter JC, Hutchison III, CA, Smith HO. 2009. Enzymatic assembly of DNA molecules up to several hundred kilobases. *Nature Methods*6, 343–345.
- 19. Gomez-Escribano JP, Song L, Fox DJ, Yeo V, Bibb MJ, Challis GL. 2012. Structure and biosynthesis of the unusual polyketide alkaloid coelimycin P1, a metabolic product of the *cpk* gene cluster of *Streptomyces coelicolor* M145. *Chemical Science***3**, 2716–2720.
- 20. Gottelt M, Kol S, Gomez-Escribano JP, Bibb M, Takano E. 2010. Deletion of a regulatory gene within the *cpk* gene cluster reveals novel antibacterial activity in *Streptomyces coelicolor* A3. *Microbiology*156, 2343–2353.
- Hwang WY, Fu Y, Reyon D, Maeder ML, Tsai SQ, Sander JD, Peterson RT, Yeh JR, Joung JK. 2013. Efficient genome editing in zebrafish using a CRISPR-Cas system. *Nature Biotechnology* 31, 227–229.
- 22. Jiang W, Bikard D, Cox D, Zhang F, Marraffini LA. 2013. RNAguided editing of bacterial genomes using CRISPR-Cas systems. *Nature Biotechnology***31**, 233–239
- 23. Jinek M, Chylinski K, Fonfara I, Hauer M, Doudna JA, Charpentier E. 2012. A programmable dual-RNA-guided DNA endonuclease in adaptive bacterial immunity. *Science***337**, 816–821.
- 24. Kahl L, Endy D. 2013. A survey of enabling technologies in synthetic biology. *Journal of Biological Engineering***7**, 13.
- Krokida A, Delis C, Geisler K, Garagounis C, Tsikou D, PeñaRodríguez LM, Katsarou D, Field B, Osbourn AE, Papadopoulou KK. 2013. A metabolic gene cluster in *Lotus japonicus* discloses novel enzyme functions and products in triterpene biosynthesis. *New Phytologist*200, 675–690.
- 26. Li J-F, Norville J, Aach J, McCormack M, Zhang D, Bush J, Church G, Sheen J. 2013. Multiplex and homologous recombination-mediated genome editing in *Arabidopsis* and *Nicotianabenthamiana* using guide RNA and Cas9. *Nature Biotechnology***31**, 688–691. Li T, Liu B, Spalding M, Weeks D, Yang B. 2012. High-efficiency TALEN-based gene editing produces diseaseresistant rice. *Nature Biotechnology***30**, 390–392.