





ENVIS RESOURCE PARTNER ON ENVIRONMENTAL BIOTECHNOLOGY

SUPPORTED BY:

MINISTRY OF ENVIRONMENT, FOREST& CLIMATE CHANGE GOVERNMENT OF INDIA, NEW DELHI

ISSN : 0974 2476

Volume-37

Number-1, 2020

NEWS LETTER

ON

APPLICATION OF CRISPR/CAS9 IN THE ENVIRONMENT



DESKU ENVIS RP, UNIVERSITY OF KALYANI, NADIA, WEST BENGAL Email: desku-envis.nic.in, Phone: +91-33-25828750 Website:http://www.deskuenvis.nic.in

Editors

Prof. Asoke Prasun Chattopadhyay (Coordinator, ENVIS RP) Dr. Subhankar Kumar Sarkar (Deputy Coordinator) Co-editor Dr. (Mrs) Anusaya Mallick (Programme Officer)

ENVIS Staffs

Mr. Sourav Banerjee (Information Officer) Mr. Tanmay Achrjee (IT Officer) Mr. Subham Dutta (Data Entry Operator)

INSTRUCTIONS TO CONTRIBUTORS

ENVIS Resource Partner on Environmental Biotechnology publishes two volumes (4 Nos.) of news letter in a year (**ISSN: 0974 2476**). The articles in the news letter are related to the thematic area of the ENVIS Resource Partner (see the website: http://deskuenvis.nic.in).

The format of the article as follows:

- 1. Font should be Times New Roman and font size to be 12 in 1.5 spacing with maximum of 4-5 typed pages.
- 2. Figures and typed table should be in separate pages and provided with title and serial numbers.
- 3. The exact position for the placement of the figures and tables should be marked in the manuscript.
- 4. The article should be below 10% plagiarized.

Articles should be sent to The Coordinator ENVIS RP University of Kalyani, Kalyani-741235 Nadia, West Bengal

EDITORIAL

The story of CRISPR (Clustered Regularly Interspaced Short Palindromic Repeat) began in late 1980s, when such sequence was discovered in E. coli. Much later, it was realized that this is part of adaptive immune system of bacteria. Using CRISPR associated genes (cas), foreign DNA are cleaved and pasted in DNA known as CRISPR arrays. When a phage DNA invades the bacterial system, CRISPR RNAs (crRNAs) are expressed by some cas proteins, which guide cas nuclease to specific area of invader DNA assisted by protospacer adjacent motif (PAM). The CRISPR-cas complex binds the DNA at a specific location and cleaves it.

Later, it was realized that such DNA cleavage can also be programmed to insert desired sequences in specific sites – using DNA repair. This can happen via nonhomologous end joining (NHEJ), or by homology directed repair (HDR). Since the second decade of the new millennium, things have moved quite fast. High fidelity cas systems, precise editing of DNA and RNA sequences, tagging for visualization and other purposes etc. have become routine. These have repercussions in research on epigenetics, drug research etc.

The articles in the present issue are a brief introduction, followed by application to plant systems and in environmental biotechnology.

> Asoke Prasun Chattopadhyay Subhankar Kumar Sarkar

IN THIS ISSUE:

Introduction to CRISPR/Cas9

• Invited Articles

Email: desku-env@nic.in

- Crispr/Cas9 mediated gene editing in plants for crop improvement and disease registance
- Application of Crispr/Cas-9 in Environmental Biotechnology
- Forthcoming Events
- Query and Feedback Form

Disclamer: Authors of the individual articles are exclusively responsible for the content of their manuscript, including appropriate citation and for obtaining necessary prior permission from the original publisher for reproduction of figures, tables and text (in whole or in part) from previous publications. Publisher and editors do not accept any legal responsibility for errors, omissions and copyright violations of authors or claims of whatsoever. The views and opinions expressed in the articles are of the authors and not reflecting the editors. ENVIS RP on Environmental Biotechnology, University of Kalyani.

Introduction to CRISPR/Cas9

CRISPR-Cas (Genetic Scissors) are adaptive immune response systems that protect prokaryotes from bacteriophages. It stands for Clustered Regularly Interspaced Short Palindromic Repeats. The discovery of the type II prokaryotic CRISPR "immune system" has allowed for the development for an RNA-guided genome editing tool that is simple, easy and quick to implement. CRISPR genome editing has revolutionized the biological sciences, from medicine to agriculture and industry, making it possible to edit the DNA sequence of any living organism.

Genome editing (also called gene editing) is a group of technologies that give scientists the ability to change an organism's DNA. These technologies allow genetic material to be added, removed, or altered at particular locations in the genome. Several approaches to genome editing have been developed. A recent one is known as CRISPR-Cas9, which is short for clustered regularly interspaced short palindromic repeats and CRISPR-associated protein 9. The CRISPR-Cas9 system has generated a lot of excitement in the scientific community because it is faster, cheaper, more accurate, and more efficient than other existing genome editing methods.

Two scientists who pioneered the revolutionary gene-editing technology are the winners of this year's Nobel Prize in Chemistry, who discovered CRISPR-Cas9. Emmanuelle Charpentier and Jennifer A. Doudna share the 2020 Nobel chemistry prize for their discovery of a game-changing gene-editing technique.

Jennifer Doudna is Professor of the Departments of Chemistry and of Molecular and Cell Biology at University of California, Berkeley and an Investigator of the Howard Hughes Medical Institute. Early in her career, she studied the structure and mechanism of ribozymes (enzymatic RNA molecules) and RNA-protein complexes. Now her research focuses on understanding how RNA molecules control gene expression in bacteria and eukaryotic cells, through CRISPR-Cas9 and RNA-mediated

mechanisms, respectively. For her outstanding scientific contributions, she was elected into the American Academy of Arts and Sciences in 2002 and the National Academy of Sciences in 2003, and was awarded the 2015 Breakthrough Prize in the Life Sciences. In 2020, Dr. Doudna received the Nobel Prize in Chemistry for her contribution to the development of CRISPR-Cas technology as a method for genome editing.

Charpentier reported the discovery in 2011 and that year struck up a collaboration with Doudna. In a landmark 2012 paper in Science1, the duo isolated the components of the CRISPR–Cas9 system, adapted them to function in the test tube and showed that the system could be programmed to cut specific sites in isolated DNA. Their programmable gene-editing system has inspired a gold rush of countless application in medicine, agriculture and basic scienceand work continues to tweak and improve CRISPR and to identify other gene-editing tools.

The Nobel Committee's selection of Emmanuelle Charpentier, now at the Max Planck Unit for the Science of Pathogens in Berlin, and Jennifer Doudna, at the University of California, Berkeley, puts an end to years of speculation about who would be recognized for their work developing the CRISPR–Cas9 gene-editing tools. The technology allows precise edits to the genome and has swept through laboratories worldwide since its inception in the 2010s. It has countless applications: researchers hope to use it to alter human genes to eliminate diseases; create hardier plants; wipe out pathogens and more.



Jennifer Doudna & Emmanuelle Charpentier Source:https://www.nature.com/articles/d41586-020-02765-9

CRISPR / CAS9 mediated gene editing in plants for crop improvement and disease resistance

Alakesh Das, Dikshita Deka, Antara Banerjee & Surajit Pathak

Department of Medical Biotechnology, Faculty of Allied Health Sciences, Chettinad Academy of Research and Education (CARE), Chettinad Hospital and Research Institute (CHRI), Chennai- 603103, India.

Abstract

Improvement of agricultural production with the use of modern breeding technologies is needed to enhance the approach for nutritious food around the globe. Among the various new technology developed for enhancement of agricultural production, CRISPR/Cas genome editing enable effective alteration in crops, hence displaying positive outcome. Here, we aim to summarize the advancement in CRISPR/Cas9 and highlight the baseediting tools as an approach for the improved crop production.

Introduction

The modern agricultural techniques that are developed, clearly shows the relation between research and technology to improve crop quality and yield. Although the conventional methods of breeding has been rapid in comparison to what was two decades ago, is likely not able to cope with the increasing food requirement along with the environmental disputes around the globe. Until the breeding of plants remains solely dependent on searching of populace of plants having adequate alternative and on methods of conventional crossing strategies to initiate characteristics into distinct aimed plants, impediment for improvement of crop will still continue. Since the first utilization of CRISPR technology, it has revolutionized research in the field of life sciences. It is thought that CRISPR technologies has the potential to overcome these restrictions and advance the breeding of plants beyond the imagination done earlier (Gao, 2018). It has also been known to improve the shelf life of climacteric fruits which has been

reported by Pathak et al.(2018) (Fig 1). CRISPR/Cas systems are differentiated chiefly into two classes, one possessing multi-subunit effectors, and the other having single protein effectors, and are further differentiated into six types on the basis of their signature proteins. Among them three are chiefly given importance, namely Type I-III which has signature proteins Cas3, Cas9 and Cas10, respectively. CRISPR/Cas9 is considered one of the most studied and extensively utilized CRISPR systems in plants. CRISPR/Cas9 systems operate bv conserving the sequence of DNA from the plasmid or bacteriophage, by which the cell is invaded. Upon re-exposure to some specific plasmid or bacteriophage, the system utilizes the sequences of its transcribed RNA detect and a Cas nuclease assisted to cleave the DNA is Streptococcus pyogenes, bacterium are the main sources of Cas9, and it has been detected as a nuclease having the capability to cleave at two active cleaving sites of the DNA i.e., one of each strand of the DNA. A single Cas9 protein can be remodelled to cleave targeted sites on the bacterium DNA (Esvelt et al., 2013; Jiang et al., 2013). Hence this mechanism of Cas9 has been widely used for the advancement of crop improvement and disease resistance which has been discussed below.



Fig 1: Schematic diagram of the steps involved in ethylene production (Pathak et al., 2018).

CRISPR technology, a promising tool for improved crop production:

The uses of CRISPR technology as promising tool in agriculture has already

been documented in plants. SDN-1 has been documented to be used in wheat, expecting to produce plants with resistance against the destroying powdery mildew. Use of SDN-3 (site-directed nuclease) to the Argos8 gene promoter in maize delivered essential character of the endogenous control gene, showed better results in yield in condition such as drought stress. SDN-1 has also been used in the aim of cause mutations in the modulatory sites of the tomato yield genes, leading to enhancement of genetic variation and yield. As the next generation of CRISPR approach has been initiated for agricultural uses, they can go further DSBbased (double-strand breaks) alteration in increasing the capability of CRISPR approach to particularly aimed sequence of DNA (Gao, 2018) (Fig 2). Removal of negative elements is a known technique in field of genetics enhancement. the Moreover, knocking out genes having unwanted characteristics is very common and a commonly used application of CRISPR/Cas9. Improved characteristics using CRISPR/Cas9 include enhancement in crop yield, enhancement in crop quality, and improved resistance of plants from both biotic and abiotic stress.



Fig 2: Mechanism of rDNA involved in the production of ethylene (Pathak et al., 2018).

• Enhancement in crop yield

The requirement of food security focuses production, the essential aim of gene editing for improving crops quality. Moreover, as production rely on numerous criteria's. knocking out negative modulators proved to have impact in yielddeciding criteria such as grain size (OsGS3), grain count (OsGn1a), panicle size (TaDEP1, OsDEP1), tiller number (OsAAP3) and grain weight (OsGW5, TaGASR7, TaGW2 and OsGLW2), and also generated the aimed phenotypes in with deterioration-of-function crops alterations in these genes, proving the capability of CRISPR/Cas9 in enhancement of crop yield (Chen et al., 2019).

• Enhancement in crop quality

The quality characteristics of crops alter depending on the particular breeding necessities. Till the present time the enhancement of crops by genome editing has influenced nutritional value, fragrance, starch content, and storage quality. Deletion of Waxy via CRISPR/Cas9 has enhanced rice cooking and eating quality. Studies has also shown that using CRISPR/Cas9 knockout waxy can generate corn line with higher production for commercial purpose (Ma et al.,2015; 2016). Using CRISPR/Cas9 Waltz. technology SBEIIb, a starch branching enzyme gene was mutated to generate higher amylose and resistant starch rice. Studies has shown that utilizing TALENaimed severance of OsBADH2, a fragrant rice line was prepared having the same 2acetyl-1-pyrroline content of the naturally altered aromatic rice species. CRISPR/Cas9 alterations encourage a unique possibility to change characteristics monitored by families of large gene with characteristics. superfluous Evidence around the globe shows that knocking out of the required genes using CRISPR/Cas9 led a way for the production of crops with the desired characters. (Chen et al., 2019).

• Resistance of plants from both biotic and abiotic stress

One of the pivotal criteria involved in crop quality and yield is stress. Plants having enhanced biotic stress resistance, comprises bacterial, fungal, insects, and viral disease resistance is possible using the CRISPR/Cas9 knockout technology. Studies has showed that utilizing CRISPR/Cas9 and TALEN technology, knocking out the TaMLO alleles in wheat was possible leading to production of plants having very high contention to powdery mildew fungus (Wang et al., 2014), also studies has showed to produce tomatoes having contention to powdery mildew (Nekrasov V et al., 2017). Blastresistant rice production was also possible by knockout of OsERF922 (Wang et al., case of viral diseases 2016). In CRISPR/Cas9 technology is capable of production of broad potyvirus-resistant eif4e cucumber, eif4g rice with resistant to tungro-disease, and cotton leaf curl disease-contention clcud cotton. In case of abiotic stress, knocking out OsHAK1, OsNramp5, and OsARM1, rancher has produced strains of rice with decreased levels of arsenic, and radioactive cesium (Chen et al., 2019). Recent studies show that pyl1/4/6 triple knockout rice produced by CRISPR/Cas9 editing enhances higher tolerance towards temperature, grain yield, and also reduces pre-harvest of sprouting. (Miao et al., 2018).

Conclusion

The remarkable potential of CRISPR/Cas9 has initiated with rapid advancement in crop breeding and its research. The versatility, robustness, simplicity of this approach have made genome alteration an effective strategy for advancement of crop production. However, challenges have been faced for precisely shifting this approach from the bench site to the field. Hence by the approach of systems and synthetic biology will permit the development of crops with improved features in the fields.

Acknowledgments

The authors are thankful to Chettinad Academy of Research and Education (CARE), Chennai, India. All authors have read and approved the final manuscript.

References

- 1 Chen K, Wang Y, Zhang R, Zhang H, Gao C. CRISPR/Cas genome editing and precision plant breeding in agriculture. Annual review of plant biology. 2019 Apr 29; 70: 667-97.
- 2 Esvelt KM, Mali P, Braff JL, Moosburner M, Yaung SJ, Church GM. Orthogonal Cas9 proteins for RNA-guided gene regulation and editing. Nature methods. 2013 Nov; 10(11): 1116-21.
- 3 Gao C. The future of CRISPR technologies in agriculture. Nature Reviews Molecular Cell Biology. 2018 May; 19(5): 275-6.
- 4 Jiang W, Bikard D, Cox D, Zhang F, Marraffini LA. RNA-guided editing of bacterial genomes using CRISPR-Cas systems. Nature biotechnology. 2013 Mar; 31(3): 233-9.
- 5 Ma X, Zhang Q, Zhu Q, Liu W, Chen Y, Qiu R, Wang B, Yang Z, Li H, Lin Y, Xie Y. A robust CRISPR/Cas9 system for convenient, high-efficiency multiplex genome editing in monocot and dicot plants. Molecular plant. 2015 Aug 3; 8(8): 1274-84.
- 6 Miao C, Xiao L, Hua K, Zou C, Zhao Y, Bressan RA, Zhu JK. Mutations in a subfamily of abscisic acid receptor genes promote rice growth and productivity. Proceedings of the National Academy of Sciences. 2018 Jun 5; 115(23): 6058-63.
- 7 Nekrasov V, Wang C, Win J, Lanz C, Weigel D, Kamoun S. 2017. Rapid generation of a transgene-free powdery mildew resistant tomato by genome deletion. Sci. Rep. 7:482.
- 8 Pathak S, Sriramulu S, Thandavan SP, Jothimani G, Banerjee A, Marotta F. Enhancement of shelf life of the climacteric fruits: a review on application of crispri technology. Trends in Technical and Scientific Research. 2018; 1(2): 555557.
- 9 Waltz E. 2016. CRISPR-edited crops free to enter market, skip regulation. Nat. Biotechnol. 34: 582
- 10 Wang F, Wang C, Liu P, Lei C, Hao W, Gao Y, Liu YG, Zhao K. Enhanced rice blast resistance by CRISPR/Cas9-targeted mutagenesis of the ERF transcription factor gene OsERF922. PloS one. 2016 Apr 26; 11(4): e0154027.
- 11 Wang Y, Cheng X, Shan Q, Zhang Y, Liu J, Gao C, Qiu JL. Simultaneous editing of three homoeoalleles in hexaploid bread wheat confers heritable resistance to powdery mildew. Nature biotechnology. 2014 Sep; 32(9): 947-51.

Application of CRISPR/Cas9 in Environmental Biotechnology

Rasmani Hazra Cold Spring Harbor Laboratory, New York, USA Email: rhazra@cshl.edu

Discoveries of CRISPR/Cas9 technology

The emergence of facile genome editing to manipulate or precisely modify the animal and plant genome using CRISPR/Cas9 technology is an indispensable tool in molecular biology (1). Although various genome engineering systems have been available since the 1970s, CRISPR by far is the most versatile, simple and precise method.

The discovery of the CRISPR editing system was started by a researcher, Francisco Mojica in 1993 to halophilic archaea (haloarchaea). He found an unexpected pattern - multiple copies of partially palindromic 30 bases long repeated sequence at regular interval (2). This regularly spaced repeat sequence had also been shown in bacteria, Escherichia coli (3) and Mycobacterium spp (4) although having no sequence similarity to haloarchaea (2). These repeat elements were termed as short sequence repeats (SSR) (5), spacer interspersed direct repeats (SPIDR) (6) or short regularly spaced repeats (SRSRs) (7). Later, the CRISPR (clustered regularly name interspaced short palindromic repeats) was accepted because of simplicity and it contemplated the major characteristics. (6).Several studies have shown that CRISPR/Cas is an adaptive bacterial immune system that protects against viral genetic material and contains a target specific protospacer that are expressed as CRISPR RNAs (crRNAs). During the CRISPR/Cas-mediated immunity, this crRNAs form a ribonucleoprotein (RNP) complex with Cas protein and cleaves invading viral DNA (or RNA) via base pairing of crRNA and target nucleic acids (8, 9). There are three distinct types of CRISPR/Cas system - type I, type II and

type III. Types I and III are found in both bacteria and archaea and they are distinguished by Cas3 and Cas10 proteins respectively, whereas type II is only found in bacteria and Cas9 protein is the critical component in this system(10).

Mechanism of action of CRISPR/Cas9mediated Genome engineering

In type II CRISPR system, Cas9 protein forms an RNP complex with dual guide RNA made up of crRNA and a transactivating crRNA (tracrRNA) which creates double strand break in the DNA (1, 11). The dual tracrRNA:crRNA was fused into a single guide RNA (sgRNA) to generate a simple two-component system in which, by changing sgRNA sequence toCas9, one can edit any DNA sequence of interest (12, 13, 14). The DNA cleavage by Cas9 enzyme creates two endogenous repair pathways, Homology-Directed Repair (HDR) and Non-Homologous End Joining (NHEJ). By exploiting these two DNA repair pathways, we can generate gene knock-outs or knock-in at the site of cleavage (Figure 1).



Fig 1. CRISPR/Cas9 gene targeting system

This simple CRISPR/Cas9 editing system has enabled a remarkable advancement in genome editing field to precisely edit genomes in a wide array of cells and organisms in a cost-effective manner. These characteristics make CRISPR/Cas9 technology a tremendously powerful tool to transform the field of basic science, biotechnology, and medicine. Remarkably, this invaluable technology is also helping scientists to address the complex environmental challenges to protect the environment and human health (Figure 2).



Fig 2. Implementation of CRISPR/Cas9 technology in agriculture and industrial biotechnology.

Applications and advancements of CRISPR/Cas9

Biofuels: To find sustainable solutions for the constant depletion of fossil fuel, biofuels, produced from biomass, are very promising alternative renewable energy resource and are environment-friendly. Plants, algae material or animal waste naturally produce byproducts such as sugars, fats or alcohols which are all potential alternative fuel sources. Usage of CRISPR techniques in biofuel production has remarkable benefits. For example, recently scientists were able to produce double the amount of biodiesel from phototrophic algae by using CRISPR genome editing technique (15). Yeast plays a critical role in the fermentation process and the pre-treatment chemicals in this process are very harmful to yeast. Current research work showed that CRISPR-modified yeasts are resistant to those chemicals during the biofuel **CRISPR-modified** production (16). acetogenic bacteria has also shown improved efficiency of ethanol production in commercial scale (17).

Bioremediation and Bioplastics: Plastics are non-biodegradable and one of the major sources of environmental pollution. Bioplastics are the alternative source of these toxic pollutants and they are essentially derived from natural biomass, such as cellulose from plants, bacterial cellulose and microalgae. The limitations of these bioplastics are cost and water resistance. However, scientists are using CRISPR/Cas9 technology to modify the carbon chain length to make bioplastics more abundant and cost-effective (18). Further, microorganisms or plants could be engineered using CRISPR for efficient bioremediation and phytoremediation, such as cleaning up oil spills or improving wastewater treatment and taking up heavy metals from soils and water.

Crop improvement: Using CRISPR/Cas9 technologies, we can make our agricultural practices more effective and ethical. Precise genetic manipulation of important agricultural crops generates ideal germplasms, which improve several crops characteristics, including high yield, overall quality, resistance to environmental challenges or microbial infection, which ultimately ameliorate food security while avoiding the introduction of foreign DNA into the food (19). Here are some examples of CRISPR-mediated editing of crops (Table 1).

Healthier forests: Trees are excellent to sequester carbon and tackle the climate crisis. Trees can be vulnerable through viral diseases, like Citrus Tristeza virus (CTV), or bacterial pathogens such as Citrus Greening (also known as Huanglongbing or HLB-disease) and put an entire species at risk. Breeding a tree in a classical method may take decades. On hand. **CRISPR-based** the other technologies can reduce that time to 1-2 years and hold a great promise to improve quality of trees of interest across the world, which eventually will create more sustainable. healthier. climateand disease-resistant tree breeders and healthy forests (20). In addition, CRISPR/Cas may edit mitochondria and chloroplasts genome and trace cell lineages to explain the patterns of plant development and detect internal and external signals for other applications in plant synthetic biology (19). Thus CRISPR/Cas technology has undoubtedly reformed and will continue to refashion both agriculture and plant biotechnology.

Environmental health and toxicology research

Genome editing tools can be used to identify toxic environmental contaminants and their exposure in public health. It advances the researcher's ability to study mechanisms of toxicity in the genomic and epigenomic pathways. For example, using genome-wide CRISPR screening. scientists were able to pinpoint the most important mechanism involved in arsenic trioxide, which is the number one environmental toxic substance according to the 2017 Agency for Toxic Substances and Disease Registry priority list. The genome-wide CRISPR screen identified 100 candidate genes either linked to sensitivity or resistance to arsenic trioxide and its exposure confirmed the increase of reactive oxygen species in cellular content (21).

Limitations and Government regulations

Despite the great potential of CRISPR/Cas technology in the improvement of agriculture and environmental challenges, it has few major concerns of its efficiency and specificity- (i) one possibility of offtarget effects because the 20-bp targeting gRNA sequence and 3 bp PAM motif (3 bp) may be present in other parts in the genome (22).(ii) The second challenge is to increase the efficiency of HDRmediated repair while reducing NHEJ repair path (23).(iii) In addition, a clear understanding of function of Cas9 enzyme is important since different types of Cas9 different bacteria functions from differently, SpyCas9 from e.g., Streptococcus pyogenes finds protospacer by search through billions of base pairs and significantly bend the DNA during this process (24). Therefore, it is absolutely imperative to gain the precise control of Cas9 activity in order to have safe and efficient genome editing in animals and plants.

The incredibly easy and cheap CRISPR editing tool has enormous potential for use in various applications such as clinical treatments to agricultural production to climate change. The expeditious progress in this technology has introduced concerns about its safe, secure, and ethical applications. Surely, genomic engineering of crops and livestock has the potential to escalate food production, but it may also affect ecosystems. Unless we have a clear view of the ecological impact of these genetically modified organisms and plants, we should use the technology in a responsible manner. Further, all the genetic modifications in plants and agriculture must be taken care of by government regulatory policy to protect our environment and human health. Additionally, genetically modified crops should assess the impact in long-term for environment and food safety before it is available commercially. It is also wise to label them as GMO so that they would be traceable and publicly acceptable in the benefits to global food security.

Table 1	l:	Applications	of	CRISPR/Cas9
technolc	οgι	in agricultura	ıl cr	ops

Crop species	Target genes	Target traits	Refe renc es
Rice	GW2, GW5, TGW2	Increased grain size	25
Rice	PYLs	Improved plant growth and production	26
Wheat	Gliadings	Reduced gluten content and exposure	27
Tomato	DMR6, MLO1	Resistance against downy and powderymildew	28, 29
Tomato	Pl, ALC	Long-shelf-life tomato	30, 31
Apple	DIPM1, 2, 4	Resistance against fire blight disease	32
Banana	ORF region of virus	Resistance against banana streak virus	33

References

1. Doudna AJ and Charpentier E (2014) The new frontier of genome engineering with CRISPR-Cas9. Science 346: 1258096-11

- Mojica FJM and Rodriguez-Valera F (2016) The discovery of CRISPR in archaea and bacteria. FEBBS J 283: 3162-9
- Ishino Y, Shinagawa H, Makino K, Amemura M, Nakata A (1987) Nucleotide sequence of the iap gene, responsible for alkaline phosphatase isozyme conversion in Escherichia coli, and identification of the gene product. J Bacteriol 169: 5429-5.
- 4. Hermans PW, Soolingen VD, Bik EM, Haas PE, Dale JW, Embden JD (1991) Insertion element IS987 from Mycobacterium bovis BCG is located in a hot-spot integration region for insertion elements in Mycobacterium tuberculosis complex strains. Infect Immun 59: 2695-10.
- Belkum A, Scherer S, Alphen L, Verbrugh H (1998) Short-sequence DNA repeats in prokaryotic genomes. Microbiol Mol Biol Rev. 62: 275-18.
- Jansen R, Embden JDA, Gaastra W, Schouls L M (2002) Identification of genes that are associated with DNA repeats in prokaryotes. Mol. Microbiol. 43: 1565-10.
- Mojica FJM, Díez-Villaseñor C, Soria E, Uez G (2000) Biological significance of a family of regularly spaced repeats in the genomes of archaea, bacteria and mitochondria. Mol Microbiol. 36:244-2
- Brouns SJ, Jore MM, Lundgren M, Westra ER, Slijkhuis RJ, Snijders AP, Dickman MJ, Makarova KS, Koonin EV, Oost J (2008) Science 321: 960-4
- DBhaya, Davison M, Barrangou R (2011) CRISPR-Cas Systems in Bacteria and Archaea: Versatile Small RNAs for Adaptive Defense and Regulation. Annu. Rev. Genet.45: 273-25
- Makarova KS and Koonin EV (2015) Annotation and Classification of CRISPR-Cas Systems. Methods. Mol. Biol. 1311: 47-75
- Gasiunas G, Barrangou R, Horvath P, Siksnys V (2012) Cas9-crRNA ribonucleoprotein complex mediates specific DNA cleavage for adaptive immunity in bacteria. Proc. Natl Acad. Sci. USA 109: E2579–E2586.
- Cho SW, Kim S, Kim JM, Kim JS (2013) Targeted genome engineering in human cells with the Cas9 RNA-guided endonuclease. Nat. Biotechnol. 31: 230-2
- Cong L, Ran FA, Cox D, Lin S, Barretto R, Habib N, Hsu PD, Wu X, Jiang W, Marraffini LA, Zhang F (2013) Multiplex genome engineering using CRISPR/Cas systems. Science. 339: 819-4
- Jinek M, East A, Cheng A, Lin S, Ma E, Doudna J (2013) RNA-programmed genome editing in human cells. eLife, 2: e00471
- Ozsoz M, Ibrahim AU, Coston PP (2019) Application of Crispr Technology for the Generation of Biofuels: A Review. J Fundam Renewable Energy Appl. 9:2
- 16. Muhammad Javed R, Noman M¹ Shahid M, Muhammad K, Rashid H, Ahmed T, Ismail M, 'SadafM, Khan F (2019) Current situation of biofuel production and its enhancement by CRISPR/Cas9mediated genome engineering of microbial cells. Microbiol. Res. 219: 1-11.
- 17. Jin S, Bae J, Song Y, Pearcy N, Shin J, Kang S, Minton NP, Soucaille P, Cho BK (2020) Synthetic Biology on Acetogenic Bacteria for Highly Efficient Conversion of C1 Gases to Biochemicals. Int. J. Mol. Science. 21: 7639-63.
- Khatami K, Zabaleta M, Owusu I, Cetecioglu Z (2020) Waste to bioplastics: How close are we to

sustainable polyhydroxyalkanoates production? Waste Management. 119: 374-14

- Zhu H, Li C, Gao C (2020) Applications of CRISPR–Cas in agriculture and plant biotechnology. Nat. Rev. Mol. Cell Biol.21: 661–16
- Dort NE, Tanguay P, Hamelin RC (2020) CRISPR/Cas9 Gene Editing: An Unexplored Frontier for Forest Pathology. Front Plant Sci. 11: 1126
- Sobh A, Loguinov A, Yazici NG, Zeidan RS, Tagmount A, Hejazi NS, Hubbard AE, Zhang L, Vulpe CD (2019) Functional Profiling Identifies Determinants of Arsenic Trioxide Cellular Toxicity. Toxicol Sci. 169: 108–121.
- 22. Cho SW, Kim S, Kim Y, Kweon J, Kim HS, Bae S, Kim JS (2014) Analysis of off-target effects of CRISPR/Cas-derived RNA-guided endonucleases and nickases. Genome Res. 1: 132–9.
- Liu M, Rehman S, Tang X, Gu X, Fan Q, Chen Q, Ma W (2018) Methodologies for Improving HDR Efficiency. Front Genet. 9: 691-12.
- 24. Jiang F and Doudna J (2017) CRISPR–Cas9 Structures and Mechanisms. Annu. Rev. Biophys. 46:505-24.
- 25. Xu R, Yang Y, Qin R, Li H, Qiu C, Li L (2016). Rapid improvement of grain weight via highly efficient CRISPR/Cas9-mediated multiplex genome editing in rice. J. Genet. Genomics 43, 529–4.
- Miao C, Xiao L, Hua K, Zoua C, Zhao Y, Bressanb RA (2018). Mutations in a subfamily of abscisic acid receptor genes promote rice growth and productivity. Proc. Natl. Acad. Sci. U.S.A 115, 6058–5.
- 27. Jouanin A, Gilissen LJW, Schaart JG, Leigh FJ, Cockram J, Wallington EJ, Boyd LA, Broeck HC, Meer V, America A, Visser R, Smulders M (2020) CRISPR/Cas9 Gene Editing of Gluten in Wheat to Reduce Gluten Content and Exposure—Reviewing Methods to Screen for Coeliac Safety. Front. Nutr. 7: 1-15
- ThomazellaD, Brail Q, Dahlbeck D,Staskawicz B (2016) CRISPR-Cas9 mediated mutagenesis of a DMR6 ortholog in tomato confers broad-spectrum disease
- resistance. BioRxiv https://doi.org/10.1101/064824.
- NekrasovV, Wang C, Win J, Lanz C, Weigel D, Kamoun S (2017) Rapid generation of a transgenefree powdery mildew resistant tomato by genome deletion. Sci. Rep. 7: 482–4
- Uluisik, S. et al. (2016) Erratum: Corrigendum: genetic improvement of tomato by targeted control of fruit softening. Nat. Biotechnol. 34: 1072-15.
- Yu Q, Wang B, Li N, Tang Y et al. (2017) CRISPR/Cas9-induced targeted mutagenesis and gene replacement to generate long-shelf-life tomato lines. Sci. Rep. 7, 11874-10.
- 32. Malnoy M, Viola R, Jung M, Koo O, Kim S, Kim S, Velasco R, Kanchiswamy NC(2016) DNA-free genetically edited grapevine and apple protoplast using CRISPR/Cas9 ribonucleoproteins. Front Plant Sci. 7: 1904-10.
- 33. Tripathi NJ, Ntui VO, Ron M, Muiruri S, Britt A, Tripathi L (2019)CRISPR/Cas9 editing of endogenous banana streak virus in the B genome of Musa spp. overcomes a major challenge in banana breeding. Commun. Biol. 2: 1-11.

-6220-

Report of

Webinar on 'Promotion of Science'

A webinar on 'Promotion of Science' in the area of Genetic Scissors (CRISPR-Cas9) was organized on 15th October, 2020 by University of Kalyani in collaboration of **ENVIS** RP Environmental on Biotechnology, DST PURSE -II, NASI and INSA under the banner of "Promotion of Science: Recent developments in interdisciplinary Science such as Physical, Biological including Agriculture, Health and Environmental Science".

The welcome address was given by Prof. Rita Ghosh, Coordinator, DST PURSE II, KU. The webinar was inaugurated by Hon'ble Vice Chancellor, University of Kalyani, Prof. Sankar Kumar Ghosh, in the virtual presence of dignitaries and eminent speakers Prof. Indrajit Lahiri, Dean, PG Faculty of Science, Prof. Rita Ghosh, Coordinator, DST PURSE II; Prof. Asoke Prasun Chattopadhyay, Coordinator, ENVIS RP; Prof. Swapan K Datta, Former VC, Visva-Bharati, Ex-Deputy Director General, (Crop Science), ICAR; Prof. Amit Ghosh, Ex-Director, IMTECH, J. C. Bose Distinguished Chair Professor, National Academy of Sciences, India; Prof. Hemanta K. Majumder, Senior Scientist NASI, IICB-CSIR, Kolkata, Conveners, NASI & INSA, Kolkata Chapter and Dr. Anindva Bandyopadhyay, Vice President, Reliance Industries Limited.

This webinar was organized keeping in the mind on the recent Nobel Prize on CRISPR -Cas9, a method of genome editing. The announcement of this year's Nobel Prize in Chemistry to Prof. Jennifer A Doudna and Prof. Emmanuelle Charpentier for their work on "a method of genome editing" has focused attention to the method called CRISPR. Using this technology, commonly termed as "molecular scissors", plasmid or vector DNA can be suitably modified to produce organisms expressing genetic materials of choice. This opens up enormous possibilities for modifying genetic materials and seeing their effects.

In the webinar students, research scholars and faculties were interacted with the eminent experts.

There are 382 participants were participated through google meet and YouTube channel

(https://www.youtube.com/watch?v=RSxU O3UjAHI&feature=youtu.be).

Lastly the concluding remarks and vote of thanks was given by Prof. Asoke Prasun Chattopadhyay, ENVIS coordinator.



ENVIS RP DESKU News Letter on 'Application of CRISPR/Cas9 in the Environment', Vol. 37(1), 2020, ISSN: 0974 2476

FORTHCOMING EVENTS					
Events	Date	Place & Correspondence			
3rd International Virtual Conference "The Ecology Of New Economy Post COVID-19"	23rd to 24th April 2021 Online	Chandigarh, India http://cgc.ac.in/international-conference- 2021			
International Workshop on Technology Innovation of Algae	1st to 3rd May 2021 Online	Taleghani Street, Tehran, Iran https://iranalgae2021.inacc.ir/			
02nd International Conference on Marine Science and Technology	24th to 24th July 2021	Aachen, Germany http://sustainableconference.science/aache n/			
2nd International Conference on Biomedical Engineering, Bio- Informatics and Design & Software Applications	24th to 25th July 2021	langkawi, Malaysia https://aniceas.com/conferences/bbds-july- 2021/			
2021 International Agriculture Innovation Conference	3rd to 4th September 2021 Online	TOKYO, JAPAN https://iaic2021.iaas.org.sg/			

QUERY AND FEEDBACK FORM

Name:

Designation:

Email:

Area of specialization:

Views on our Newsletter:

Suggestion for Improvement:

I would like to collect information on Environmental Biotechnology on the following:

Subject:

Key words:

	BOOK POST
	то
FROM:	
ENVIS RP	
DEPARTMENT OF	
ENVIRONMENTAL SCIENCE,	
UNIVERSITY OF KALYANI	
KALYANI-741235, NADIA	
WEST BENGAL	