OMICS-Based Approaches in Plant Biotechnology



Rintu Banerjee Garlapati Vijay Kumar P. Jeevan Kumar





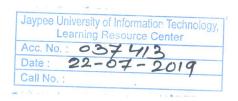
WILEY

OMICS-Based Approaches in Plant Biotechnology

Edited by
Rintu Banerjee,
Garlapati Vijay Kumar
and S.P. Jeevan Kumar







This edition first published 2019 by John Wiley & Sons, Inc., 111 River Street, Hoboken, NJ 07030, USA and Scrivener Publishing LLC, 100 Cummings Center, Suite 541J, Beverly, MA 01915, USA © 2019 Scrivener Publishing LLC

For more information about Scrivener publications please visit www.scrivenerpublishing.com.

All rights reserved. No part of this publication may be reproduced, stored in a retrieval system, or transmitted, in any form or by any means, electronic, mechanical, photocopying, recording, or otherwise, except as permitted by law. Advice on how to obtain permission to reuse material from this title is available at http://www.wiley.com/go/permissions.

Wiley Global Headquarters

111 River Street, Hoboken, NJ 07030, USA

For details of our global editorial offices, customer services, and more information about Wiley products visit us at www.wiley.com.

Limit of Liability/Disclaimer of Warranty

While the publisher and authors have used their best efforts in preparing this work, they make no representations or warranties with respect to the accuracy or completeness of the contents of this work and specifically disclaim all warranties, including without limitation any implied warranties of merchantability or fitness for a particular purpose. No warranty may be created or extended by sales representatives, written sales materials, or promotional statements for this work. The fact that an organization, website, or product is referred to in this work as a citation and/or potential source of further information does not mean that the publisher and authors endorse the information or services the organization, website, or product may provide or recommendations it may make. This work is sold with the understanding that the publisher is not engaged in rendering professional services. The advice and strategies contained herein may not be suitable for your situation. You should consult with a specialist where appropriate. Neither the publisher nor authors shall be liable for any loss of profit or any other commercial damages, including but not limited to special, incidental, consequential, or other damages. Further, readers should be aware that websites listed in this work may have changed or disappeared between when this work was written and when it is read.

Library of Congress Cataloging-in-Publication Data

ISBN 978-1-119-50993-6

Cover image: The Editors

Cover design by Russell Richardson

Set in size of 11pt and Minion Pro by Manila Typesetting Company, Makati, Philippines

Printed in the USA

Contents

In	Introduction		
Pa	rt 1	: Genomics	1
1	Exp	loring Genomics Research in the Context of Some	
	Und	lerutilized Legumes—A Review	3
	Patr	ush Lepcha, Pittala Ranjith Kumar and N. Sathyanarayana	
	1.1	Introduction	3
	1.2	Velvet Bean [Mucuna pruriens (L.) DC. var. utilis	
		(Wall. ex Wight)] Baker ex Burck	4
	1.3	Psophocarpus tetragonolobus (L.) DC.	7
	1.4	Vigna umbellata (Thunb.) Ohwiet. Ohashi	8
	1.5	Lablab purpureus (L.) Sweet	9
	1.6	Avenues for Future Research	10
	1.7	Conclusions	12
		Acknowledgments	12
		References	12
2	Ove	rview of Insecticidal Genes Used in Crop	
	Imp	rovement Program	19
	Nee	raj Kumar Dubey, Prashant Kumar Singh,	
		endra Kumar Yadav and Kunwar Deelip Singh	
		Introduction	19
	2.2	Insect-Resistant Transgenic Model Plant	21
	2.3	Insect-Resistant Transgenic Dicot Plants	27
	2.4	Insect-Resistant Transgenic Monocot Plants	34
	2.5	Working Principle of Insecticidal Genes Used in Transgenic	
		Plant Preparation	39
	2.6	Discussion	41
		References	42

vi Contents

3		ances in Crop Improvement: Use of miRNA Technologies	
		Crop Improvement	55
	Clarissa Challam, N. Nandhakumar		
	and	Hemant Balasaheb Kardile	
	3.1		56
	3.2	Discovery of miRNAs	56
	3.3	Evolution and Organization of Plant miRNAs	57
	3.4	Identification of Plant miRNAs	58
	3.5	miRNA vs. siRNA	59
	3.6	Biogenesis of miRNAs and Their Regulatory Action in Plants	60
	3.7	Application of miRNA for Crop Improvement	61
	3.8	Concluding Remarks	62
		References	70
4	Gen	e Discovery by Forward Genetic Approach in the Era	
		igh-Throughput Sequencing	75
		k Thakur and Samart Wanchana	
	4.1	Introduction	75
	4.2	Mutagens Differ for Type and Density of Induced Mutations	76
	4.3	High-Throughput Sequencing is Getting Better and Cheaper	77
	4.4	Mapping-by-Sequencing	77
	4.5	Different Mapping Populations for Specific Need	81
	4.6	Effect of Mutagen Type on Mapping	83
	4.7	Effect of Bulk Size and Sequencing Coverage on Mapping	83
	4.8	Challenges in Variant Calling	85
	4.9	Cases Where Genome Sequence is either Unavailable	
		or Highly Diverged	85
	4.10	Bioinformatics Tools for Mapping-by-Sequencing Analysis	86
		Acknowledgments	87
		References	87
5	Fun	ctional Genomics of Thermotolerant Plants	91
	Nag	endra Nath Das	
		Introduction	91
	5.2	Functional Genomics in Plants	93
	5.3	Thermotolerant Plants	94
	5.4	Studies on Functional Genomics of Thermotolerant Plants	98
	5.5	Concluding Remarks	99
		Abbreviations	100
		References	100

Pa	art 2	2: Met	tabolor	nics	105
6	Pre-Processing and Statistical Analysis to Biological Insight			107	
		1	a B. Misr	ra	
			duction	_	108
	6.2		ods and I		109
			Source		109
		6.2.2		ing of Raw Mass Spectrometry Data	109
		6.2.3		cal Analyses	109
		6.2.4		y Enrichment and Clustering Analysis	110
	6.3	Resul			110
				of the Study and Data Analysis	110
				ard Cell Metabolomics Dataset	110
		6.3.3		riate Analysis for Insights into Data	
			Pre-Pro		113
		6.3.4		f Data Normalization Methods	119
		Discu			122
	6.5	Conc			124
			icts of In		124
			owledgm	ent	125
		Refer	ences		125
7	Met	abolite	Profilin	g and Metabolomics of Plant Systems	
			MR and		129
				esh Lingwan and Shyam K. Masakapalli	
	7.1		duction	<i>y</i>	129
	7.2	Mater	rials and l	Methods	131
		7.2.1	¹ H NMF	R-Based Metabolite Profiling of Plant Samples	132
				Metabolite Extraction	132
				¹ H NMR Spectroscopy	132
			7.2.1.3	Qualitative and Quantitative Analysis	102
				of NMR Signals	134
		7.2.2	Gas Chr	comatography–Mass Spectroscopy (GC-MS)	101
				Metabolite Profiling	134
			7.2.2.1	Sample Preparation	134
			7.2.2.2	GC-MS Data Acquisition	135
			7.2.2.3	GC-MS Data Pretreatment and Metabolite	100
				Profiling	136
			7.2.2.4	Validation of Identified Metabolites	136
		7.2.3		riate Data Analysis	137
					101

viii Contents

	7.3	Selected Applications of Metabolomics and Metabolite Profiling	139
		Acknowledgments	140
		Competing Interests	140
		References	140
8	OM	IICS-Based Approaches for Elucidation of Picrosides	
	Bio	synthesis in <i>Picrorhiza kurroa</i>	14
	Var	un Kumar	
	8.1	Introduction	140
	8.2	Cross-Talk of Picrosides Biosynthesis Among Different	
		Tissues of <i>P. kurroa</i>	148
	8.3	Strategies Used for the Elucidation of Picrosides	
		Biosynthetic Route in <i>P. kurroa</i>	148
		8.3.1 Retro-Biosynthetic Approach	149
		8.3.2 <i>In Vitro</i> Feeding of Different Precursors and Inhibitors	149
		8.3.3 Metabolomics of Natural Variant Chemotypes	
		of P. kurroa	150
	8.4	Strategies Used for Shortlisting Key/Candidate Genes	
		Involved in Picrosides Biosynthesis	15
		8.4.1 Comparative Genomics	151
		8.4.2 Differential Next-Generation Sequencing (NGS)	
		Transcriptomes and Expression Levels of Pathway	
		Genes Vis-à-Vis Picrosides Content	152
	8.5	Complete Architecture of Picrosides Biosynthetic Pathway	153
	8.6	Challenges and Future Perspectives	161
		Abbreviations	162
		References	163
9	Rele	evance of Poly-Omics in System Biology Studies	
	of I	ndustrial Crops	167
	Nag	rendra Nath Das	
	9.1	Introduction	167
	9.2	System Biology of Crops	169
	9.3	Industrial Crops	171
	9.4	Poly-Omics Application in System Biology Studies	
		of Industrial Crops	176
	9.5	Concluding Remarks	177
		Abbreviations	177
		References	178

Pa	art 3	: Bioinformatics	183	
10	Emerging Advances in Computational Omics Tools for Systems Analysis of Gramineae Family Grass Species and Their Abiotic			
	Stress Responsive Functions			
	Pan	diyan Muthuramalingam, Rajendran Jeyasri,		
	Dho	amodharan Kalaiyarasi, Subramani Pandian,		
	Sub	ramanian Radhesh Krishnan, Lakkakula Satish,		
	Shu	nmugiah Karutha Pandian and Manikandan Ramesh		
		Introduction	186	
	10.2	Gramineae Family Grass Species	187	
		10.2.1 Oryza sativa	187	
		10.2.2 Setaria italica	187	
		10.2.3 Sorghum bicolor	188	
	100	10.2.4 Zea mays	188	
		Abiotic Stress	188	
	10.4	Emerging Sequencing Technologies	198	
		10.4.1 NGS-Based Genomic and RNA Sequencing	199	
		10.4.2 Tanscriptome Analysis Based on NGS	200	
	10 5	10.4.3 High-Throughput Omics Layers	201	
		Omics Resource in Poaceae Species	202	
	10.6	Role of Functional Omics in Dissecting the Stress Physiology		
	107	of Gramineae Members	203	
	10.7	Systems Analysis in Gramineae Plant Species	204	
		Nutritional Omics of Gramineae Species	205	
		Future Prospects	205	
	10.10	Conclusion	206	
		Acknowledgments	207	
		References	207	
11	OM	IC Technologies in Bioethanol Production: An Indian		
	Con	text	217	
	Pulk	it A. Srivastava and Ragothaman M. Yennamalli		
	11.1	Introduction	217	
	11.2	Indian Scenario	219	
	11.3	Cellulolytic Enzymes Producing Bacterial Strains Isolated		
		from India	220	
		11.3.1 Bacillus Genus of Lignocellulolytic Degrading Enzymes	222	
		11.3.2 Bhargavaea cecembensis	222	
		11.3.3 Streptomyces Genus for Hydrolytic Enzymes	230	

x Contents

	11.4	Biomass Sources Native to India	230
		11.4.1 Albizia lucida (Moj)	230
		11.4.2 Areca catechu (Betel Nut)	231
		11.4.3 Arundo donax (Giant Reed)	231
		11.4.4 Pennisetum purpureum (Napier Grass)	231
		11.4.5 Brassica Family of Biomass Crops	231
		11.4.6 Cajanus cajan (Pigeon Pea)/Cenchrus americanus	
		(Pearl Millet)/Corchorus capsularis (Jute)/	
		Lens culinaris (Lentil)/Saccharum officinarum	
		(Sugarcane)/Triticum sp. (Wheat)/Zea mays (Maize)	232
		11.4.7 Medicago sativa (Alfalfa)	232
		11.4.8 Manihot esculenta (Cassava)/Salix viminalis	
		(Basket Willow)/Setaria italica (Foxtail Millet)/	
		Setaria viridis (Green Foxtail)	232
		11.4.9 Vetiveria zizanioides (Vetiver or Khas)	232
		11.4.10 Millets and Sorghum bicolor (Sorghum)	233
		Omics Data and Its Application to Bioethanol Production	233
	11.6	Conclusion	239
		References	239
Pa	rt 4	: Advances in Crop Improvement: Emerging	
			245
16	CIIII	ologies	245
12	Gen	ome Editing: New Breeding Technologies in Plants	247
	Kaly	ani M. Barbadikar, Supriya B. Aglawe,	
	Sate	ndra K. Mangrauthia, M. Sheshu Madhav	
	and	S.P. Jeevan Kumar	
	12.1	Introduction: Genome Editing	248
	12.2	GE: The Basics	249
		12.2.1 Nonhomologous End-Joining (NHEJ)	250
		12.2.2 Homology Directed Repair (HR)	251
	12.3	Engineered Nucleases: The Key Players in GE	251
		12.3.1 Meganucleases	251
		12.3.2 Zinc-Finger Nucleases	256
		12.3.3 Transcription Activator-Like Effector Nucleases	257
		12.3.4 CRISPR/Cas System: The Forerunner	258
	12.4	Targeted Mutations and Practical Considerations	259
		12.4.1 Targeted Mutations	259
		12.4.2 Steps Involved	260
		12.4.2 Steps mvolved	_ 0
		12.4.2.1 Selection of Target Sequence	261

		12.4.2.3 Transformation	263
		12.4.2.4 Screening for Mutation	264
	12.5	New Era: CRISPR/Cas9	264
		12.5.1 Vector Construction	264
		12.5.2 Delivery Methods	266
		12.5.3 CRISPR/Cas Variants	266
		12.5.3.1 SpCas9 Nickases (nSpCas9)	266
		12.5.3.2 Cas9 Variant without Endonuclease Activity	266
		12.5.3.3 FokI Fused Catalytically Inactive Cas9	267
		12.5.3.4 Naturally Available and Engineered Cas9	
		Variants with Altered PAM	268
		12.5.3.5 Cas9 Variants for Increased On-Target Effect	268
		12.5.3.6 CRISPR/Cpf1	268
	12.6	GE for Improving Economic Traits	269
		12.6.1 Development of Next-Generation Smart Climate	
		Resilient Crops	271
		12.6.2 Breaking Yield Incompatibility Barriers	0.51
		and Hybrid Breeding	271
		12.6.3 Creating New Variation through Engineered QTLs	271
		12.6.4 Transcriptional Regulation	272
		12.6.5 GE for Noncoding RNA, microRNA12.6.6 Epigenetic Modifications	272273
		12.6.7 Gene Dosage Effect	273
	12.7	Biosafety of GE Plants	273
		What's Next: Prospects	276
	1210	References	276
	-		270
13		ulation of Gene Expression by Global Methylation Pattern	
		ants Development	287
		esh Kumar Yadav, Krishan Mohan Rai, Nishant Kumar	
		Vikash Kumar Yadav	200
		Introduction	288
		Nucleic Acid Methylation Targets in the Genome	289
		Nucleic Acid Methyl Transferase (DNMtase)	290
		Genomic DNA Methylation and Expression Pattern Pattern of DNA Methylation in Early Plant Life	291
		DNA Methylation Pattern in Mushroom	292293
		Methylation Pattern in Tumor	293
		DNA Methylation Analysis Approaches	294
	10.0	13.8.1 Locus-Specific DNA Methylation	295
		13.8.2 Genome-Wide and Global DNA Methylation	295

xii Contents

		13.8.3 Whole Genome Sequence Analysis by Bioinformatics Analysis	296
		References	297
14	High	h-Throughput Phenotyping: Potential Tool for Genomics	303
		vani M. Barbadikar, Divya Balakrishnan, C. Gireesh,	
	Hem	ant Kardile, Tejas C. Bosamia and Ankita Mishra	
	14.1	Introduction	304
	14.2	Relation of Phenotype, Genotype, and Environment	304
	14.3	Features of HTP	306
	14.4	HTP Pipeline and Platforms	310
	14.5	Controlled Environment-Based Phenotyping	311
	14.6	Field-Based High-Throughput Plant Phenotyping (Fb-HTPP)	311
	14.7	Applications of HTP	313
		14.7.1 Marker-Assisted Selection and QTL Detection	314
		14.7.2 Forward and Reverse Genetics	315
		14.7.3 New Breeding Techniques	315
		14.7.3.1 Envirotyping	315
	14.8	Conclusion and Future Thrust	316
		References	316
Inc	lex		323

The book provides a detailed description of how OMICS can help crop science and horticulture to enhance crop yields, resistance and nutritional values.

Burgeoning world population, decreased water supply and land resources, coupled with climate change, result in severe stress conditions which is a great threat to the global food supply. To meet these challenges, exploring OMICS technologies could lead to improved yields of cereals, tubers and grasses that may ensure food security. Improvement of yields through crop improvement and biotechnological means are the need-of-the-hour, and the current book "OMICS-Based Approaches in Plant Biotechnology", reviews the advanced concepts on breeding strategies, OMICS technologies (genomics, transcriptomics and metabolomics) and bioinformatics that help to glean the potential candidate genes/molecules to address unsolved problems related to plant and agricultural crops. The first six chapters of the book are focused on genomics and cover sequencing, functional genomics with examples on insecticide resistant genes, mutation breeding and miRNA technologies. Recent advances in metabolomics studies are elucidated in the next 3 chapters followed by 5 chapters on bioinformatics and advanced techniques in plant biotechnology and crop breeding.

Audience

The information contained in the volume will help plant breeders, plant biotechnologists, plant biochemists, agriculture scientists and researchers as well as policy makers in using this applied research to focus on better crop breeding and stress adaptation strategies.

Rintu Banerjee, Ex-MNRE- Chair-Professor, Indian Institute of Technology, Kharagpur has created a niche of her own in the area of Biomass Deconstruction/Biofuel Production/Enzyme Technology. In the process of her innovative development, she was granted 8 Indian, 3 international (US, Japanese and Chinese) patents. She has published more than 180 papers in peer-reviewed national/international journals.

Garlapati Vijay Kumar is an Assistant Professor at the Department of Biotechnology and Bioinformatics, Jaypee University of Information Technology, Waknaghat, India. He has 3 patents, 33 research articles in peer-reviewed journals and 10 book chapters to his credit. His research interest areas are: Bioprocess engineering / industrial biotechnology, deployment of OMICS technologies for crop improvement, fermentation technology, biofuels, and biocatalysis.

S.P. Jeevan Kumar is a scientist in ICAR-Indian Institute of Seed Science, Mau, U.P, India. His interests include OMICS technologies for plant biotechnology, crop improvement, seed deterioration mechanisms, genetic purity and bioenergy. He has published 25 papers in peer-reviewed journals and multiple book chapters.

Cover Design: Russell Richardson Cover images courtesy of Pixabay.com

WILEY





