1. Record Nr. UNINA9911022162503321 Autore Tiwari Shalini Titolo Decoding Non-Coding RNA: Enhancing Plant Abiotic Stress Tolerance / / edited by Shalini Tiwari, Jose L Gonzalez-Hernandez, Charu Lata Singapore:,: Springer Nature Singapore:,: Imprint: Springer,, 2025 Pubbl/distr/stampa **ISBN** 9789819698592 9789819698585 Edizione [1st ed. 2025.] Descrizione fisica 1 online resource (408 pages) Collana Concepts and Strategies in Plant Sciences, , 2662-3196 Gonzalez-HernandezJose L Altri autori (Persone) LataCharu Disciplina 580 Soggetti **Botany** Plant genetics Stress (Physiology) **Plants** Plant Science Plant Genetics Plant Stress Responses Lingua di pubblicazione Inglese **Formato** Materiale a stampa Livello bibliografico Monografia Nota di contenuto 1. Introduction to Non-Coding Regions -- 2. Structural and Functional Diversity of Non-Coding Regions -- 3. Promoter regulatory regions of abiotic stress responses -- 4. Long Non-Coding RNAs (IncRNAs) in Abiotic Stress Adaptation -- 5. Epigenetic Regulation of Abiotic Stress Responses -- 6. Mobile Genetic Elements and Abiotic Stress Tolerance -- 7. Genome-Wide Profiling of Abiotic Stress-Responsive Non-Coding RNAs -- 8. Non-Coding RNA-Mediated Signaling Pathways in Abiotic Stress Tolerance -- 9. Evolutionary Dynamics of Stress-Responsive Non-Coding Regions -- 10. Genomic Resources for Studying Stress-Responsive Non-Coding Regions -- 11. Utilization of non-coding regions for enhancing stress resilience in crop species -- 12. Challenges and Future Directions in Non-Coding Region Research for

This book offers an in-depth exploration of non-coding RNA and its

Stress Tolerance.

Sommario/riassunto

critical role in enhancing plant abiotic stress tolerance. It provides a comprehensive overview of non-coding DNA's structural and functional diversity and its regulatory mechanisms in stress responses. The chapters cover topics such as epigenetic regulation, mobile genetic elements, and genomic profiling techniques. It covers latest research on non-coding RNA-mediated regulation, transcriptional regulatory elements controlling stress-responsive gene expression, and the role of long non-coding RNAs (IncRNAs) in fine-tuning gene expression networks under stress conditions. The book also discusses chromatin modifications, transposon-mediated regulation, high-throughput sequencing approaches for identifying stress-responsive ncRNAs, and CRISPR-based functional analysis. In addition to exploring these advanced topics, the book highlights practical implications for crop improvement strategies. Researchers in plant biology, genomics, molecular biology, and agricultural sciences will find this book particularly beneficial. It is also an essential resource for educators looking to provide their students with up-to-date information on cutting-edge research techniques. By offering both theoretical perspectives and practical case studies from renowned experts worldwide, this volume equips readers with actionable knowledge to address real-world challenges in agriculture.