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Collana	Compendium of Plant Genomes, , 2199-479X
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Nota di contenuto	Introduction Genomics resources for functional genomics in poplars The poplar Pangenome Epigenomics in poplar Chromosomal rearrangements and 3D organization of the poplar genome Sexbased identity in poplars Phylogenomics, comparative genomics, interspecific hybridization in poplar Genomics of Chinese poplars Genomic landscape of adaptation in European poplars Genomic landscape of adaptation in North American poplars Poplar and the associated microbiome Wood formation and valorization in poplar Genomics of secondary metabolism in poplar Applied Genomics in poplar: isoprene emissions Applied Genomics in poplar: genetic modifications Applied Genomics in poplar: adaptation to climate change Applied Genomics in poplar: phytoremediation Applied Genomics in poplar: innovative breeding Applied Genomics in poplar: bioenergy Prospects: The Poplar Genome, a Model for Understanding Angiosperm Evolution and Supporting Tree Improvement Efforts
Sommario/riassunto	This book is the first comprehensive compilation of research on state of the art genomics on the most advanced model tree species

including genome assemblies, insights into genomic structural features and methylation patterns, wholegenome resources used for population genomics and adaptation to climate, enabled breeding vs. classical genetics and traditional breeding, comparative genomics, and elucidations on functional genomics. The latest developments in the genomics of wood formation are particularly highlighted. Altogether, the book contains over 300 pages in over 15 chapters authored by globally reputed experts in the relevant fields of this tree crop's genomics research. This book is useful for students, teachers, and scientists in academia and governmental or private tree improvement agencies or companies interested in genetics, pathology, entomology, physiology, molecular genetics and breeding, in vitro culture and genetic engineering, land restoration, and agroforestry solutions.