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This book, inclusive of 21 chapters, explores the impact of genome plasticity of both pathogens and host to human infections and their control. Diseases discussed cover a wide range of infections caused by diverse pathogens, including diseases caused by bacteria (i.e., *Escherichia coli*, enterohaemorrhagic *E. coli*, *Helicobacter pylori*, coagulase-negative staphylococci, *Legionella pneumophila*, L....

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mechanisms of genome plasticity include dna methylation noncoding rnas dna variants chromatin and histone modifications genome plasticity has been associated with multiple human diseases and related phenotypes human genomics has led to large advances in a deeper understanding of the etiology and pathophysiology of human diseases

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Introduces the rapidly evolving field of genome plasticity, presents the latest research findings, and explores the relevance of these findings to infection and infection control. Compiles and analyzes current investigations on the genome fluidity of pathogenic microbes.

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