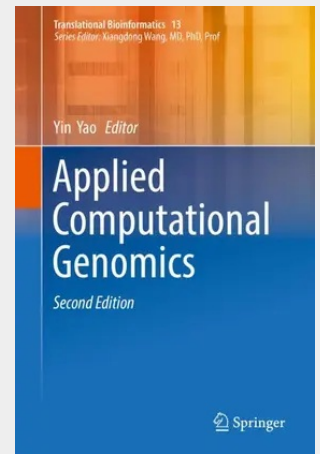


Applied Computational Genomics

The volume provides a review of statistical development and application in the area of human genomics, including candidate gene mapping, linkage analysis, population-based genome-wide association, exon sequencing, and whole genome sequencing analysis. The authors are extremely experienced in the field of statistical genomics and will give a detailed introduction to the evolution of the field, as well as critical comments on the advantages and disadvantages of the proposed statistical models. The future directions of translational biology will also be described.

The volume provides a review of statistical development and application in the area of human genomics, including candidate gene mapping, linkage analysis, population-based genome-wide association, exon sequencing, and whole genome sequencing analysis. The authors are extremely experienced in the field of statistical genomics and will give a detailed introduction to the evolution of the field, as well as critical comments on the advantages and disadvantages of the proposed statistical models. The future directions of translational biology will also be described.



106,99 €

99,99 € (zzgl. MwSt.)

Lieferfrist: bis zu 10 Tage

Artikelnummer: 9789811310706

Medium: Buch

ISBN: 978-981-13-1070-6

Verlag: Springer

Erscheinungstermin: 22.09.2018

Sprache(n): Englisch

Auflage: 2. Auflage 2018

Serie: Translational Bioinformatics

Produktform: Gebunden

Gewicht: 442 g

Seiten: 150

Format (B x H): 160 x 241 mm

