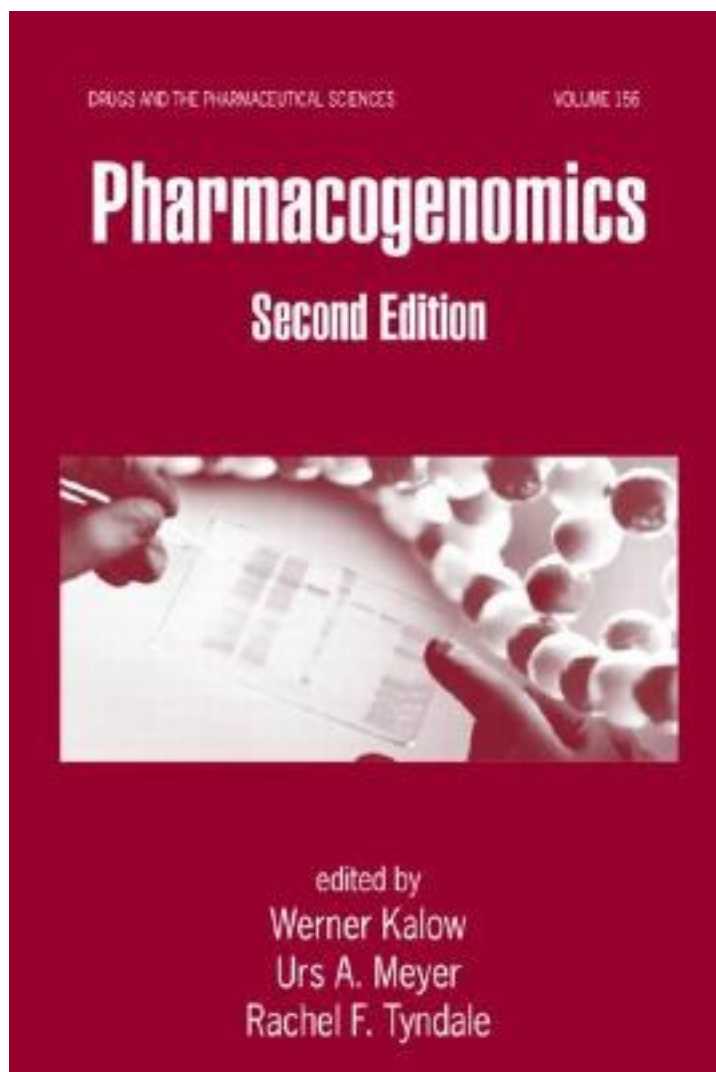


Pharmacogenomics



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Expert researchers and physician/clinicians describe in detail the newest and most commonly used technologies today in this rapidly advancing field. The authors provide readily reproducible methods for assessing the functional consequences of a certain polymorphism, evaluate the variety of genotyping platforms currently available, and discuss the management of pharmacogenomic information. Highlights include techniques for making a snapshot of the allele-specific variation in human gene expression, genome wide analysis of allele-specific expression using oligo microarrays, in vivo assays with HaploChIP, SNP genotyping in DNA pools, and PharmGKB, the pharmacogenetics and pharmacogenomics knowledge base. The methodologies for genotyping include denaturing high-performance liquid chromatography, pyrosequencing, kinetic-fluorescence detection, mass spectrometry, and TaqMan assay for insertion/deletions.

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