1. Record Nr. UNINA9910877873803321 **Titolo** Batch effects and noise in microarray experiments, sources, and solutions / / edited by Andreas Scherer Pubbl/distr/stampa Chichester, West Sussex: Hoboken,: J. Wiley, 2009 **ISBN** 1-282-37950-X 9786612379505 0-470-68598-0 0-470-68599-9 Descrizione fisica 1 online resource (282 p.) Collana Wiley Series in Probability and Statistics;; v.868 Altri autori (Persone) SchererAndreas <1966-> Disciplina 572.8/636 Soggetti DNA microarrays DNA microarrays - Experiments Lingua di pubblicazione Inglese **Formato** Materiale a stampa Livello bibliografico Monografia Note generali Description based upon print version of record. Nota di bibliografia Includes bibliographical references and index. Nota di contenuto Batch Effects and Noise in Microarray Experiments; Contents; List of Contributors; Foreword; Preface; 1 Variation, Variability, Batches and Bias in Microarray Experiments: An Introduction; 2 Microarray Platforms and Aspects of Experimental Variation; 2.1 Introduction; 2.2 Microarray Platforms; 2.2.1 Affymetrix; 2.2.2 Agilent; 2.2.3 Illumina; 2.2.4 Nimblegen; 2.2.5 Spotted Microarrays; 2.3 Experimental Considerations; 2.3.1 Experimental Design; 2.3.2 Sample and RNA Extraction; 2.3.3 Amplification; 2.3.4 Labeling; 2.3.5 Hybridization; 2.3.6 Washing; 2.3.7 Scanning 2.3.8 Image Analysis and Data Extraction 2.3.9 Clinical Diagnosis; 2.3.10 Interpretation of the Data; 2.4 Conclusions; 3 Experimental Design; 3.1 Introduction; 3.2 Principles of Experimental Design; 3.2.1 Definitions: 3.2.2 Technical Variation: 3.2.3 Biological Variation: 3.2.4 Systematic Variation; 3.2.5 Population, Random Sample, Experimental and Observational Units; 3.2.6 Experimental Factors; 3.2.7 Statistical Errors: 3.3 Measures to Increase Precision and Accuracy: 3.3.1 Randomization; 3.3.2 Blocking; 3.3.3 Replication; 3.3.4 Further Measures to Optimize Study Design

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Sommario/riassunto

Batch Effects and Noise in Microarray Experiments: Sources and Solutions looks at the issue of technical noise and batch effects in microarray studies and illustrates how to alleviate such factors whilst interpreting the relevant biological information. Each chapter focuses on sources of noise and batch effects before starting an experiment, with examples of statistical methods for detecting, measuring, and managing batch effects within and across datasets provided online. Throughout the book the importance of standardization and the value of standard operating procedures in the devel

7.1.1 Microarray Gene Expression Data