

***Development of Intracellular Flow Karyotype Analysis**

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Mapping and Sequencing with BACs and Fosmids

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Towards a Globally Integrated, Sequence-Ready BAC Map of the Human Genome

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Generation of Normalized and Subtracted cDNA Libraries to Facilitate Gene Discovery

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Mapping in Man-Mouse Homology Regions

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Positional Cloning of Murine Genes

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Human Artificial Episomal Chromosomes (HA ECS) for Building Large Genomic Libraries

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***Cosmid and cDNA Map of a Human Chromosome 13q14 Region Frequently Lost at B Cell Chronic Lymphocytic Leukemia**

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Informatics

BCM Server Core

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A Freely Sharable Database-Management System Designed for Use in Component-Based, Modular Genome Informatics Systems

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A Software Environment for Large-Scale Sequencing

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Generalized Hidden Markov Models for Genomic Sequence Analysis

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Identification, Organization, and Analysis of Mammalian Repetitive DNA Information

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***TRRD, GERD and COMPEL: Databases on Gene-Expression Regulation as a Tool for Analysis of Functional Genomic Sequences**

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Data-Management Tools for Genomic Databases

Victor M. Markowitz and I-Min A. Chen

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The Genome Topographer: System Design

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A Flexible Sequence Reconstructor for Large-Scale DNA Sequencing: A Customizable Software System for Fragment Assembly

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The Role of Integrated Software and Databases in Genome Sequence Interpretation and Metabolic Reconstruction

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