



Bacterial Artificial Chromosomes: Volume 2: Functional Studies (Methods in Molecular Biology)

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Several developmental and historical threads are woven and displayed in these two volumes of Bacterial Artificial Chromosomes, the first on Library Construction, Physical Mapping, and Sequencing, and the second on Functional Studies. The use of large-insert clone libraries is the unifying feature, with many diverse contributions. The editors have had quite distinct roles. Shaying Zhao has managed several BAC end-sequencing projects. Marvin Stodolsky during 1970–1980 contributed to the elucidation of the natural bacteriophage/prophage P1 vector system. Later, he became a member of the Genome Task Group of the Department of Energy (DOE), through which support flowed for most clone library resources of the Human Genome Program (HGP). Some important historical contributions are not represented in this volume. This preface in part serves to mention these contributions and also briefly surveys historical developments. Leon Rosner (deceased) contributed substantially in developing a PAC library for drosophila that utilized a P1 virion-based encapsidation and transfection process. This library served prominently in the Drosophila Genome Project collaboration. PACs proved easy to purify so that they substantially replaced the YACs used earlier. Much of the early automation for massive clone picking and processing was developed at the collaborating Lawrence Berkeley National Laboratory. However, the P1 virion encapsidation system itself was too fastidious, and P1 virion-based methods did not gain popularity in other genome projects.

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