The applications of computers to research on nucleic acids II Part 2

Edited by D Söll & R J Roberts

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PREFACE

Two years have passed since the first special issue of NUCLEIC ACIDS RESEARCH devoted to the applications of computers to research on nucleic acids (vol 10, 1-456, 1984). In that time many laboratories have acquired on-line access to thei university computer or have invested in a personal nicrocomputer. No longer are these machines viewed as toys, but rather have assumed the importance of other, more familiar items of equipment in a molecular biology laboratory. Two major data bases, GENBANK and the EMBL data base, have come into existence in response to the need for computer access to published sequences. The software needed to analyze nucleic acid and protein sequences continues to be developed, and already one commercial enterprise has emerged that both develops and maintains a large software package for such analysis.

Despite the widespread interest in programs for manipulating nucleic acid sequences rather few of them are well known since most have been developed locally and do not enjoy widespread circulation. It is the aim of this second special issue to heighten the awareness of both scientists and programmers to the broad range of software that is currently available. It is our hope that this issue, like the previous one, will provide a useful starting point for those investigators newly discovering a need for programs in this area and will stimulate those already deeply involved to exchange and improve existing programs.

Richard J. Roberts

Dieter Söll

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