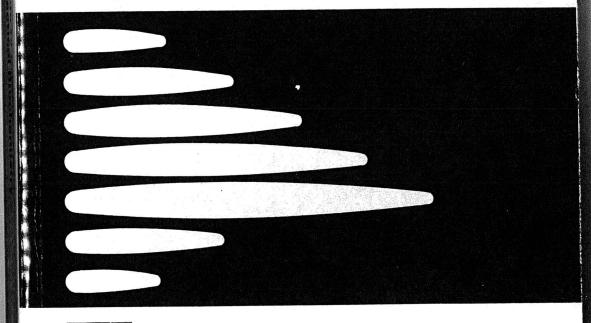
### A Biologist's Guide to

# Principles and Techniques of Practical Biochemistry

Third Edition

Edited by Keith Wilson and Kenneth H. Goulding





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### 170 Molecular biology techniques

5'---TACGCTCG - 32P 3' Single-stranded DNA, labelled only at its 3' end

Modification of 'C' using hydrazine; this removes base, leaving ribosyl urea

---TACGCTCG-32P

---TACGCTCG-32P

Cleavage at modified bases, using piperidine

G-32P

TCG-32P

GCTCG-32P

gCTCG-32P

plus non-radioactive fragments

Separate on sequencing gel alongside products of other

modification / cleavage reactions (as in Fig. 5.13)

**Fig. 5.14** Maxam and Gilbert sequencing of DNA. Only modification and cleavage of deoxycytidine is shown, but three more aliquots of the end-labelled DNA would be modified and cleaved at G, G+A, and T+C, and the products would be separated on the sequencing gel alongside those from the 'C' reactions.

to that produced by the Sanger method, since each sample now contains radioactive molecules of various lengths, all with one end in common (the labelled end), and with the other end cut at the same type of base. Analysis of the reaction products by electrophoresis is as described for the Sanger method.

Because the Sanger method produces oligonucleotides which are radio-actively labelled throughout their lengths, rather than only at one end, the molecules can be made a lot more radioactive, and therefore easier to detect; so less DNA is needed for sequencing. Once M13 cloning has been set up in a laboratory, it provides a very convenient and rapid way to obtain single-stranded DNA. For these reasons, dideoxy sequencing of M13-cloned DNA is probably the most commonly used sequencing method, though the chemical procedure is still used by many laboratories.

### 5.5.3 Protein sequencing

Although protein sequencing may seem out of place in a section dealing with the analysis of DNA, the molecular biologist can often make use of a



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