

References

1. Judson, HF. (1987). *Mapping the Human Genome: Historical Background* (Mapping Our Genes contractor reports, Vol. 1, NTIS Order No. PB 88-160-783/AS). Office of Technology Assessment, US Congress. September 1987.
2. Sanger, F, and Coulson, AR. (1975). Rapid Method for Determining Sequences in DNA by Primed Synthesis with DNA-Polymerase. *Journal of Molecular Biology* 94 : 441-448.
3. Sanger, F. (1975). The Croonian Lecture, 1975: Nucleotide Sequences in DNA. *Proceedings of the Royal Society of London B* 191 : 317-333.
4. Maxam, AM, and Gilbert, W. (1977). A New Method for Sequencing DNA. *Proceedings of the National Academy of Sciences (USA)* 74 (February): 560-564.
5. Smith, LM, Sanders, JZ, Kaiser, RJ, Hughes, P, Dodd, C, Connell, CR, Heiner, C, Kent, SBH, and Hood, LE. (1986). Fluorescence Detection in Automated DNA Sequence Analysis. *Nature* 321 (12 June): 674-679.
6. Smith, TF. (1990). The History of the Genetic Sequence Databases. *Genomics* 6 (April): 701-707.
7. Bishop, MJ, and Rawlings, CJ. (1987). *Nucleic Acid and Protein Sequence Analysis: A Practical Approach*. IRL Press, Oxford, UK.
8. Waterman, MS. (1989). *Mathematical Methods for DNA Sequences*. CRC Press, Boca Raton, FL.