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PHYLIP version 3.6 is my own package. It is available free, from its Web site, in C source code, or as executables for Windows, Mac OS X, and Mac OS 8 or 9. The C source code can easily be compiled on Unix or Linux systems. Top Categories. DNA Gel Electrophoresis. Carolina makes DNA gel electrophoresis easy when studying forensics or genetics. There's a simple set up with consistent results. Jun Adachi and Masami Hasegawa have written a package MOLPHY, version 2.3b3, carrying out maximum likelihood inference of phylogenies for either nucleotide sequences or protein sequences. Their protein sequence maximum likelihood program, ProtML, is a successor to the one they made available to me and which I formerly distributed on a nonsupported basis in PHYLIP. Medicine. Discover high-impact journals, award-winning books and eBooks and online solutions. Browse the catalog of books and journals