

## Acknowledgments

### ACKNOWLEDGMENTS

The HIV Sequence Database and Analysis Project is funded by the Vaccine and Prevention Research Program of the AIDS Division of the National Institute of Allergy and Infectious Diseases (Dr. James Bradac, Project Officer) through an interagency agreement with the U.S. Department of Energy.

We thank the many researchers who have made their sequences available prior to publication, and authors who help by contributing to our review section.

We also thank editors from previous editions, Simon Wain-Hobson, Kuan-Teh Jeang, Lou Henderson, Thomas Leitner, Gerald Myers and George Pavlakis, among others, who were not editors of this year's publication, but who may at some future time return in this role.

### The Cover

The cover of the 1998 database depicts the 3-dimensional crystal structure of gp120 with the asparagine residues (including side chains) modified by N-linked glycosylation in mammalian cells shown in blue. Note the concentration of glycosylation on the gp120 surface believed to face outward on the trimeric envelope glycoprotein spike. This figure is from a review of the *Structure of the Core of HIV-1 gp120 Exterior Envelope Glycoprotein* by Richard Wyatt, Peter D. Kwang, Wayne A. Hendrickson and Joseph G. Sodroski, in this volume, page III-1.