

Program modifications:

Version 1.1 (May 10, 2003):

The Windows version of the program was having difficulty handling very small migration rates. To avoid this problem, a fix was put in where frequencies are not allowed to decrease past a small value. Also, the program was modified to decrease running time, especially when there are only two populations.

Version 1.2 (August 28, 2003):

Some users had data files which exceeded the pre-set size for number of individuals or number of alleles. These values were increased and a note was included in the help file describing the new maximum values.

Three new output options were also added; the change in log likelihood values, the distribution of log likelihood values when the chain was sampled, and the likelihood ratio test of the prior and posterior distributions (which is currently under development).