**README**

**Data from: Data and code from: Automated Identification of Insect Vectors of Chagas Disease in Brazil and Mexico: The Virtual Vector Lab**

This file includes all programming code and images connected with the paper listed above, which is published in *PeerJ*. The overall mass of code and data is quite large (about 13 GB), so the file had to be split into 15 pieces, each just less than 1 GB. This splitting was achieved using the freeware program hjsplit, which is available from <http://www.hjsplit.org/>.

Specific descriptions of files and code contained within the file are provided in an additional “README” file contained therein.