# Installation

## EMERGE package

EMERGE is distributed as a self-extracting package. It automatically detects or installs the proper version of the MATLAB Compiler Runtime (MCR)
( http://www.mathworks.com/products/compiler/mcr/ ). The EMERGE package can be extracted to a location of your choice. Do not move the program to another directory because the program and the Genome subdirectory are required to remain at the same location. The Genome subdirectory lists text files containing chromosome lengths and gene positions. For mouse genome version 9 (mm9), human genome version 19 (hg19) and fruitfly genome version 3 (dm3), the files have been provided with the program package. When using data from other genomes, such files need to be placed in the Genome subdirectory and the EMERGE program will detect them automatically.

You'll find the *EMERGE* paper here