



Software

- BioSpice (<http://biospice.lbl.gov/>), Berkeley BioSpice Network Representation and Modeling System (NRMS)
- Copasi (<http://www.copasi.org>), Mendes group at VBI and Kummer group at EML Research
- Dizzy (<http://labs.systemsbioology.net/bolouri/software/Dizzy/>), Bolouri Group, Institute for Systems Biology, Seattle
- E-Cell (<http://ecell.sourceforge.net/>), Institute for Advanced Biosciences, Keio University
- MCell (<http://www.mcell.cnl.salk.edu/>), Salk Institute and Pittsburgh Supercomputing Center
- SBML (<http://sbml.org/index.psp>), XML-based representation language for biochemical network models
- STOCKS (<http://www.sysbio.pl/stocks/>), by Kierzek group at SysBio.pl.
- The Silicon Cell (<http://www.bio.vu.nl/hwconf/Silicon/>), BioCentrum Amsterdam, CWI, University of Amsterdam
- Virtual Cell (<http://www.nrcam.uchc.edu/>), National Resource for Cell Analysis and Modeling, NRCAM