Web-based dynamic simulation environment for system-level understanding of cellular network systems

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Abstract: We develop WebCellTM which is a java-based simulation environment for system-level understanding of biological systems. A web-accessible repository of cellular network models has been established to investigate the dynamics of such models. In addition, its efficient and user-friendly web interface allows users to import their own models described by SBML (Hucka et al., 2003) for representing computational models in systems biology. Consequently, dynamic simulations of the imported models can be carried out from anywhere an internet connection is available. Acknowledgements: This work was supported by the National Research Laboratory Program (2000-

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