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**BIOMAX INFORMATICS AG
releases a new version of its flagship product,
the BioXM Knowledge Management Environment.**

MARTINSRIED, Germany — (23 May 2007) — Adding new data management capabilities for high-throughput “-omics” research, BioXM™ version 2.4 widens its target market as a premier enterprise platform for the integration of life science knowledge using semantic technologies.

New Web Services support enables organizations to easily build their own applications on top of the BioXM server, further leveraging this central inventory of information based on a user-created, extendible scientific knowledge model. The updated BioXM client application ensures a highly personalized work experience, adding new support for Mac OS X® as a client platform and extended data exchange capabilities now supporting OpenDocument formats.

The updated BioXM framework also includes an optional module for text mining based on the BioLT™ Literature Mining Tool, which combines biological and medical term dictionaries with powerful free-text querying capabilities. By integrating the BioLT engine, the BioXM platform is extended with an efficient mechanism allowing theme-specific knowledge bases to be quickly and easily built.

The BioXM Knowledge Management Environment is a fully customizable knowledge management (KM) solution for knowledge-based drug discovery, drug development and other areas such as translational medicine. This project-centered, distributed software platform facilitates communication and collaboration in a research environment. Users create, manage and visualize scientific models as an extendible network of interrelated concepts describing a particular area of research. Scientists conveniently stay in touch with recent

additions or changes in knowledge that should be available to the entire organization while focusing specifically on generating knowledge in a particular scientific field. Newly generated knowledge is easily published to a broader audience within the BioXM Knowledge Management Environment.

About Biomax

Biomax Informatics AG (Martinsried, Germany), founded in 1997, is a leader in the development of customized bioinformatics solutions. Biomax developed the well-known Pedant-Pro™ Sequence Analysis Suite, the BioRS™ Integration and Retrieval System, the BioXM™ Knowledge Management Environment and other computational solutions for better decision making and knowledge management in the life science industry. Additional information about Biomax can be found at the company's site on the World Wide Web at www.biomax.com.

About the BioXM Knowledge Management Environment

The BioXM Knowledge Management Environment is a project-centered, distributed software platform that provides a central inventory of information and knowledge. Users create, manage and visualize scientific models as an extendible network of interrelated concepts. The BioXM platform facilitates communication and collaboration within research environments, allowing you to focus on science.

About the BioLT Literature Mining Tool

The BioLT Literature Mining Tool is a customizable tool for intuitive and structured text mining. BioLT version 2.0 combines biological and medical term dictionaries with powerful free-text querying capabilities. The tool provides comprehensive and structured answers to complex questions. Search results can be used for iterative refinement and extension of queries.

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