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Sophic and Biomax Win Second NCI Cancer Gene Curation Project

Falmouth, MA – October 10, 2005. Sophic Systems Alliance Inc. today announced, together with Biomax Informatics AG, that Sophic signed a second project with National Cancer Institute to text mine Medline abstracts and manually annotate and validate the results to identify all cancer genes, gene-disease relationships and cancer gene-compound relationships.

“The Sophic-Biomax team delivered on our expectations in the Cancer Gene Curation Pilot Project. As the second phase of the project moves forward, this growing, high-quality repository of cancer information will be available to the NCI caBIG community,” said Dr. George Komatsoulis, the NCI Project Manager. “Our goal is to provide a single source for cancer genes with the disease and compound relationships that will accelerate cancer research.”

“Last year, we successfully completed a pilot project with NCI that demonstrated the high quality of results produced through the use of BioLT™, the Biomax Linguistic System, and the accuracy of refinements performed by Biomax scientists in the manual annotation team. For the second phase of this long-term NCI project, Sophic was able to use its recent GSA Award to be the prime integrator for this project,” said Patrick Blake, CEO of Sophic.

Dr. Klaus Heumann, President of Biomax Informatics AG said, “We are honored to support this vital effort and we are building a flexible knowledge management environment, the BioXM Knowledge Management System, to leverage this high quality cancer content.” Heumann went on to say, “The NCI Cancer Gene content is being combined with the manually annotated Biomax Human Genome Database currently integrated within the BioXM Knowledge Management System.”

“The convergence of the NCI Project with BioXM Knowledge Management System development supports our Alliance strategy to focus on helping researchers find cures for complex diseases, starting with cancer,” said Blake.

Additional details on the project are provided in a white paper and a joint NCI-Biomax Poster on the Sophic Systems Alliance Inc. website. Go to www.sophicalliance.com

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About Sophic Systems Alliance.

Sophic Systems Alliance Inc. (SSA) is an integrator of bioinformatics software development companies and provides a full range of professional services. SSA sells, markets, implements and supports the delivery of a broad range of software modules developed by Biomax Informatics AG, Geospiza, Inc. and other members of the alliance. The company is focused on providing professional services to scientists and agencies in Washington, D.C. including scientific consulting and custom bioinformatics software development. Customers include government research organizations, pharmaceuticals, and biotech companies. Sophic is an Optimized IBM Business Partner. More information is available at www.sophicalliance.com.

About Biomax Informatics AG.

Biomax, a recognized pioneer in the development of customized bioinformatics solutions, was founded in 1997 as a spin-off of the GSF-MIPS academic research group, now the German Research Center for Environment and Health-Institute for Bioinformatics (GSF-IBI). Founded by Prof. Dr. H. W. Mewes, Prof. Dr. D. Frishman and Dr. K. Heumann, Biomax developed a broad portfolio of bioinformatics solutions, including the well-known Pedant-Pro™ Sequence Analysis Suite, the BioRS™ Data Integration and Retrieval System, and other bioinformatics tools and services used in metabolic analysis, proteomics, and gene expression analysis. Additional information about Biomax can be found at the company's site on the World Wide Web at www.biomax.com.

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