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**GENEGO ADDS BRISTOL-MYERS SQUIBB TO CUSTOMER BASE**

*Company licenses MetaCore™ systems biology database*

**St. Joseph, MI** – August 10th, 2004 – GeneGo, Inc., a leading systems biology company, today announced that Bristol-Myers Squibb has licensed its MetaCore™ data mining suite that combines analytical tools, data content and algorithms for understanding complex interconnected pathways that are affected in common human diseases.

MetaCore™'s manually curated content covers all levels of cellular functionality from GPCRs and other receptors to signal transduction cascades, transcriptional factors, core effector networks, endogenous and xenobiotics metabolism and the relevant chemical compounds (metabolites and xenobiotics). The main application of MetaCore™ is in visualization and analysis of complex experimental data including microarray gene expression, SAGE and SNP DNA sequences, proteomics and metabolites profiles.

“BMS is the fifth major life science company which licensed MetaCore for internal development in recent months” said Tatiana Nikolskaya, Ph.D., President and CSO of GeneGo. “We are pleased that MetaCore’s features, in particular its curated content and ability to integrate customer-generated data into the database, meet Bristol-Myers Squibb’s expectations in systems biology.”

**About GeneGo**

GeneGo is a Michigan-based company developing systems biology technology for life science research. The original computational platform allows an integration and expert analysis of different kinds of experimental data (mRNA expression, proteomics, metabolites, phenotypic data etc.) and relevant active chemistry (metabolites, drugs, other xenobiotics) within the framework of biological pathways and networks. GeneGo's first product, MetaCore™, assists pharmaceutical scientists in the areas of target selection and validation, identification of biomarkers for disease states and toxicology. For more information, please visit the company's Web site at [www.genego.com](http://www.genego.com).

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