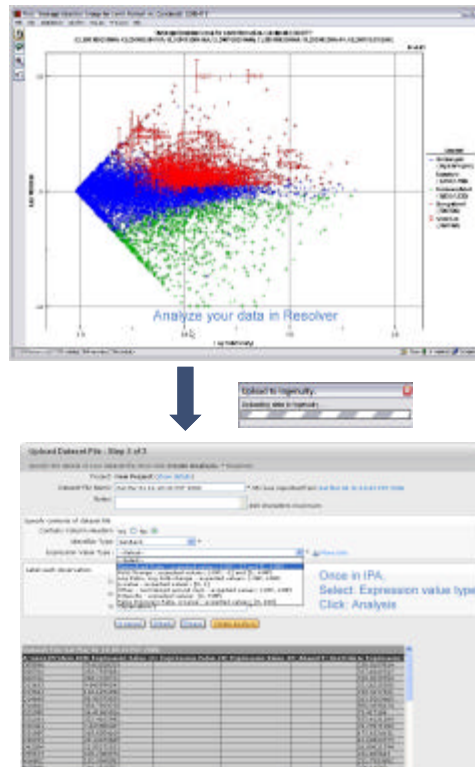


Integration Module for Ingenuity Pathway Analysis and Rosetta Resolver® System

BENEFITS

- ⇒ Combines the analytical power of the Resolver system with the biology of IPA.
- ⇒ Allows data to be analyzed using various tools in the Resolver system, which can then be further analyzed in IPA to infer the biology.
- ⇒ The integration allows useful interoperability between the two systems that allows researchers to iteratively analyze their data with minimal effort.



Now send an experiment from the Rosetta Resolver system to IPA and perform data analysis.

INTRODUCTION

The Rosetta Resolver – Ingenuity IPA integration module enables users of Rosetta Biosoftware’s Resolver system and Ingenuity’s IPA to exchange information between the two systems. Users of the Resolver system will be able to send Gene Lists and Experiment information to IPA. Once in IPA, researchers can perform network analyses, view relevant canonical pathways, create graphical and interactive diagrams, and use the search function to obtain information on genes or proteins of interest, as well as add proprietary information and interactions or pathways. Users will be able to send a list of genes back to the Resolver system, which allows further iterative analysis of those genes.

FEATURES

- All Rosetta Resolver system genomes will connect with IPA Gene View.
- Resolver users can send user-specified gene lists or experiments from the Resolver system to IPA and perform data analysis or map pathways.
- IPA users will be able to send user-specified gene lists from IPA back to the Resolver application.

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