

The Physiological Pathways Portal:

A new interactive online tool providing efficient access to genomic and phenomic information through biological pathway analysis



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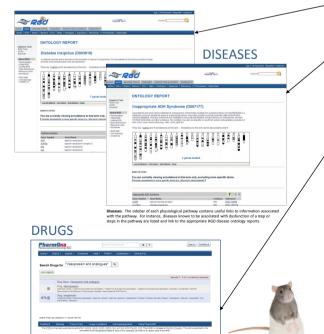
Abstract

Aim: The Rat Genome Database (RGD, http://rgd.mcw.edu) has been developing tools and resources to aid investigators in the integration of genotype and phenotype data derived from rat research for the past 10 years. The new Physiologica Pathway Portal consists of navigational hubs which allow physiologists to easily link phenotype and genotype data through intuitive mechanistic pathway flowcharts.

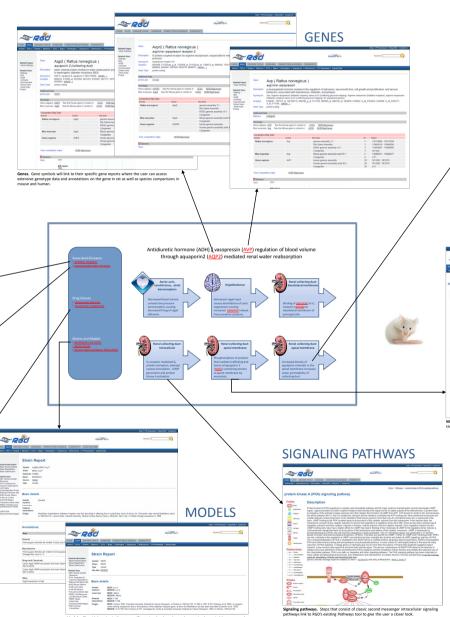
Methods: RGD has always contained valuable information for physiologists, however the focus has been primarily or genomic data, making it somewhat challenging for physiologists to navigate. The Physiological Pathway diagrams provide a familiar man for the user including links to relevant genotype and phenotype data through flowcharts depicting related physiological and regulatory mechanisms and pathways. Additionally, pharmacological actions and drug-gene interactions that influence these pathways are depicted within the diagrams and also link to related phenotype and genotype data. The diagrams are also linked to reports for individual genes involved in any of the pathways. In addition, links from the diagram to phenotype data allow the user to compare physiological measurements across multiple rat strains under a variety of experimental conditions or to access strain-specific variation data such as SNPs and other polymorphisms for direct linkage of the genotype and phenotype data. Examples of the types of pathways included in this portal are varied and complex regulatory and physiological systems such as blood pressure regulation, hematopoiesis and angiogenesis

Conclusion: The new Physiological Pathways Portal at RGD promises to be an effective and user-friendly means to access and retrieve physiological and genomic data for researchers who use the rat as a model to study specific physiological processes and pharmacological modulation of these processes.

Physiological Pathways are graphical depictions of multi-organ biological processes that provide the user with a systems biology approach to RGD. Physiologists will find the Physiological Pathways logically organized making it easy to locate and access pertinent information. Discrete steps within each process are arranged temporally and regionally with organ/tissue location of each step indicated by symbol and text label for clarity. Each pathway consists of clickable elements leading to information regarding genes involved in the process, experimental data at specific points in the process, associated diseases due to dysfunction of a step in the process, genetic strains that have been characterized as disease models, drugs classes that can favorably or adversely affect the process, and associated intracellular pathways.



Drugs. The sidebar lists generic classes of pharmaceuticals that are used in rat research and clinically. The classes link to the PharmGKB (www.pharmgkb.org) website where information on specific drugs in each



their strain report pages in RGD.

PHENOTYPE DATA



MOLECULAR PATHWAYS



Conclusion

The new Physiological Pathways portal serves as a gateway into RGD for researchers in the biological sciences searching for gene function annotation and links to information regarding genotypes, phenotypes and disease associations. It is organized in a systems biology context through sequential temporal and regional steps in physiological processes with links to gene reports, phenotype data, intracellular molecular pathways. The genetics researcher will also find it a valuable resource for determining the functional aspects and potential interactions of a particular gene of interest. New pathways will continually be added on a regular basis.

Physiological Pathways seamlessly ties together many of the best features of RGD into a functionally coherent access point making it easier and more efficient than ever to navigate through the diverse but extensive wealth of genomic and phenomic information that is RGD.



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