

RGD'S NEW PHYSIOLOGICAL PATHWAY DIAGRAMS: LINKING BIOLOGICAL SYSTEMS TO THE GENOME

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Abstract

The Bat Genome Database (RGD) has developed new Physiological Pathway Diagram that function as navigational hubs, allowing physiologists to easily follow links from known process mechanisms to phenotype and genotype data. These diagrams, which dence physiological processes occurring on the systems and tissue levels, are designed to seamlessly integrate with RGD's existing intracellular molecular pathway diagrams, allowing users to drill down from whole animal function to cellular and molecular mechanisms and back up again.

Physiological Pathway Diagrams provide a familiar and logical map for the physiologist and include links to relevant genotype data with R60, such as gene reports and sequence data, as well as phenotype data, such as experimental data, strains and models, through flowcharts depicting related physiological and regulatory mechanisms and pathways. Additionally, pharmacological actions, drug-gene interactions, as well as genetic manipulations such as gene knockouts that influence these pathways are depicted within the diagrams and also link to related phenotype and genotype data.

The first diagram to be released depicts the role of vasopressin (AVP. also known as antiduretic hormone, ADPI) in the activation of aquapoint -2 water channels in the renal collecting ducts to promote increased water reabscrytion by the kidney when blood and fluid volume has been reduced due to dehydration, hemorrhage, etc. As with all of the diagrams within this tool, this consists of an overview depiciting the stimulus, result, and feedback mechanism. Then, each involved organ or tissue can be zoomed in to reveal the specific mechanisms of the process at the tissue or multicellular level. Links to genotype and phenotype data are available at both the system and tissue levels.

Examples of the types of pathways included in this portal are complex regulatory and physiological systems such as blood pressure regulation (including blood vessel reactivity, cardiac contractility, and blood volume regulation as well as neurohumoral modulation of these processes) and glucose homeostasis (including regulation of insulin and glucago neosetsis and willows incroported on a onegoing basis.

The Physiological Pathway Diagrams tool at RGD promises to be a dynamic and user-friendly tool with which to access and retrieve physiological physiological processes and the modulation of these processes through pharmacologic or genetic interventions or for the elucidation of disease mechanisms.

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.

Let any RGDer know if you have a favorite pathway that you would

like to see developed for RGD Physiological Pathways!



Users can follow the pathways throughout the process from "challenge" (green star) to "resolution" (red target) by hovering their mouse over the shapes on each arrow to reveal a description of each step in the path. In this example, the challenge to the system is low blood volume. The resolution is increased renal fluid reabsorption to return blood volume to normal. Clicking on the kidney allows users to see a zoomed in view of how the process works on the tissue level (below).

Tissue level view



This level illustrates the intercellular mechanisms that are required in order to achieve the desired "resolution", shown in the systems level view. Descriptions of each step in the process appear upon mouse-over. The user can further zoom in to see associated canonical intracellular pathways available in RGD (e.g. PKA pathway). Links to RGD gene reports and phenotype data as well as data and information from outside RGD are available by clicking within the diagram. Alterations in the process that occur as a result of disease or drug intervention are illustrated on associated tabs.



Conclusion

The new *Physiological Pathways* tool 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, and 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. Pathways currently under construction are: vascular smooth muscle contraction and insulin-mediated glucose transport.

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.