**lature** 





# Reactome - a knowledgebase of human biological pathways

Bijay Jassal<sup>1</sup>, Esther E Schmidt<sup>1</sup>, Guanming Wu<sup>2,4</sup>, Imre Vastrik<sup>1</sup>, David Croft<sup>1</sup>, Bernard de Bono<sup>1</sup>, Gopal Gopinath<sup>2</sup>, Marc Gillespie<sup>2,4</sup>, Steven Jupe<sup>1</sup>,

European Bioinformatics Institute is an Outstation of the European Molecular Biology Laboratory.

Gavin O'Kelly¹, Lisa Matthews³, Phani V Ğarapati¹, Michael Caudy².4, Alexander Kanapin².4, Bruce May².4, Shahana Mahajan³, Ewan Birney¹ Peter D'Eustachio3, Lincoln Stein2,

<sup>1</sup>European Bioinformatics Institute, Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SD, United Kingdom <sup>2</sup>Cold Spring Harbor Laboratory, 1 Bungtown Road, Cold Spring Harbor NY 11724, USA <sup>3</sup>NYU School of Medicine, 550 First Avenue, New York NY 10016, USA

<sup>4</sup>OICR 101 College St., Suite 800, Toronto, ON, Canada M5G0A3







# www.reactome.org

# **Abstract** Reactome is a manually curated knowledgebase, created and peer-reviewed by

experts in their field. Its robust data model describes life processes ranging from metabolism to signal transduction and the cell cycle. Data is cross-referenced to publicly available web-based informatics resources. Reactome features the SkyPainter tool for analysis of high-throughput datasets, and the Mart query and data retrieval tool. All data are freely available and can be exported in a number of

#### **Stats** April 09

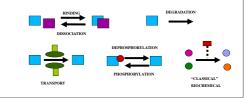
3700 proteins

3200 reactions

960 pathways

## Reactions captured

- · Basic "unit" of Reactome
- Encapsulates many events and states found in biology



### Where Reactome's Data Comes From

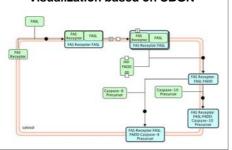


- Expert and curator create outline of new pathway
- Expert provides the biological information
- Curator enters information into database
- Reviewer (another expert) checks biological correctness

**Topics** 

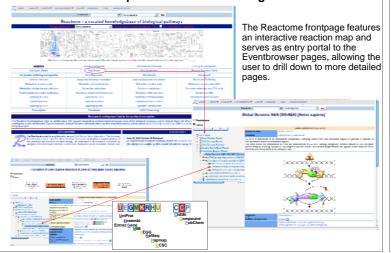
- Apoptosis
- · Cell cycle
- DNA repair
- · Transcription, mRNA processing, translation, post-translational modification
- Signaling pathways (insulin, NOTCH, opioid, NGF, EGFR, FGFR, Rho GTPases II)
- Hemostasis
- Metabolism (energy, amino acid, lipid, nucleotide, xenobiotic)
- HIV and Influenza lifecycles
- · Neurotoxicity of botulinum toxins
- and more..

## Visualization based on SBGN



### **Reactome Data Model** Pathway Pathway Reaction Reaction Input 1 Reaction Output Input 2 CatalystActivity

# Reactome Frontpage With Interactive Reaction Map and Eventbrowser Pages



#### Reactome Resources

- · mysql database dump
- SBML
- BioPAX level 2
- PDF/RTF "Textbook"
- "Interaction" and ID mapping files
- · Local installation
- Author- and curator-tools
- SOAP based webservices
- SVG
- · List of participants

## **SkyPainter**



#### **Reactome Mart**

- · Query submission and data retrieval
- · Canned queries
- · Various output formats (HTML, TSV, XLS)
- Combined queries across UniProt, Ensembl, Wormbase and PRIDE BioMarts

#### **Human versus other species**

- Manual curation focus on human
- Electronic inference to 22 other species based on orthology data from OrthoMCL
- · Cross-species comparison webpage

The development of Reactome is supported by a grant from the US National Institutes of Health (P41 HG003751) and a grant from the EBI Industry Programme. We also acknowledge the OrthoMCL group at the University of Pennsylvania, USA for providing orthology data