Cell Circuits

gcn* gal4 GO:0003677 "DNA binding"

Search   Load Example Query

Species to include:

Drosophila melanogaster
Homo sapiens
Plasmodium falciparum
Saccharomyces cerevisiae

Sort results:
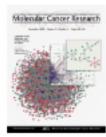
By Number of Query Terms Matching Model

P-value cutoff:

0.0001

Select publication(s) to search: Select All   Unselect All

☑ Begley TJ, Rosenbach AS, Ideker T, Samson LD. (2002)
Damage recovery pathways in Saccharomyces cerevisiae revealed by genomic phenotyping and interactome mapping.
Mol. Cancer Res. Dec;1(2):103-12.

[PubMed]

☑ Bernard A, Hartemink AJ. (2005)
Informative structure priors: joint learning of dynamic regulatory networks from multiple types of data.
Pac. Symp. Biocomput 2005 :459-70.

[PubMed]

☑ de Lichtenberg U, Jensen LJ, Brunak S, Bork P. (2005)
Dynamic complex formation during the yeast cell cycle.
Science Feb 4;307(5710):724-7.

[PubMed]

☑ Gandhi TK, Zhong J, et al. (2006)
Analysis of the human protein interactome and comparison with yeast, worm and fly interaction datasets.
Nat. Genetics Mar;38(3):285-93.

[PubMed]

☑ Hartemink AJ, Gifford DK, Jaakkola TS, Young RA. (2002)
Combining location and expression data for principled discovery of genetic regulatory network models.
Pac. Symp. Biocomput. 2002 :437-49.

[PubMed]

☑ Haugen AC, Kelley R, et al. (2004)
Integrating phenotypic and expression profiles to map arsenic-response networks.
Genome Biol. 5(12):R95. Epub 2004 Nov 29.

[PubMed]