Data Distillery Apps

Explore Data Distillery data with these apps



Find all genes that are highly expressed in a GTEx tissue and may be perturbed by a specific compound based on data from the LINCS LIOOO dataset, and are known drug targets based on data curated by IDG.



For a specific disease in Metabolomics Workbench, find all IDG and GTEx assertions that may be related to the disease in a specific tissue type.



Select a drug processing enzyme to view drugs that up or down regulate it from LINCS as well as tissues where it is highly expressed based on data from GTEX.

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For a given set of genes that are highly expressed in a specific disease, identify extracellular RNAs (exRNAs) that may be detected in human biofluids and thus serve as non-invasive disease biomarkers.

For a given drug term, return (1) upregulated genes, (2) loci expressed in exRNAX that overlap RBP binding regions, (3) the RBP that overlaps the locus, and (4) the biofluid(s) where locus was expressed in exRNA.