

## Making a heatmap of expression with MeV

Download the MultiExperiment Viewer onto your computer.

From a copy of your full info transcripts.txt file from Rockhopper do the following:

Filter for significant Q values

Insert a column next to each of your expression value columns and fill with  
=(expression value next door+1)

Then copy and paste into a new worksheet

Your gene name column

Then each of your expression value +1 columns (use paste special and paste  
in values)

Save as tab delimited file on desktop

Open MeV

Go to 'File' -> 'Load Data'

Say Forbid and just close windows for any questions that come up. It is trying to  
access accessory info online and you don't need it.

Browse to your file

Click on the upperleft most expression value

Hit 'Load'

Voila

Go to 'File' -> 'Save Image' and choose the jpg format which you can then pdf which  
should be scalable and hold resolution.