Exercise 1B: Explore and Compare Genetic Alterations ("Copy Number") in two Genes (ERBB2 and KRAS) in TCGA Ovarian Cancer Data set in the USCS Cancer Browser

our new tool, UCSC Xena.

Add Datasets

LGG copy nun

The UCSC Cancer Browser is no longer under development. For the latest data try

3,080,419,480 bp

1. Go to the UCSC Cancer Browser at <a href="https://genome-cancer.ucsc.edu/proj/site/hgHeatmap/">https://genome-cancer.ucsc.edu/proj/site/hgHeatmap/</a>. Ignore the message in red at the top.

2. On the left panel, click on "Add Datasets."

3. Find the dataset named "TCGA ovarian serious cystadenocarcinoma."

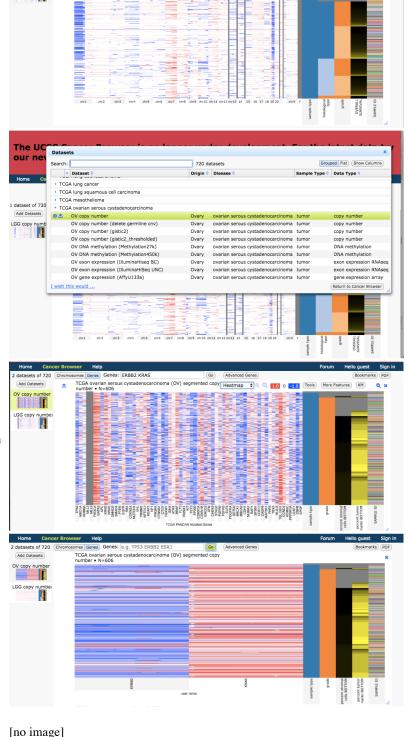
4. Under this dataset, find "OV copy number" data that is available for 606 patients. Click on this dataset once; this will select the dataset and display it in the browser.

5. In the top center of the page (above the image of the dataset), you will see two tabs: "Chromosomes" and "Genes."

6. Click on "Genes" and in the text box, enter: "ERBB2 KRAS." (Space is the delimiter between the two genes.)

[no image]

5. Click "Go." You should see one gene



whose copy number is mostly amplified (red color) and the other gene that has many deletions (blue color).

[no image]