An Introduction to the ISB-CGC Web App SeqPeek

brought to you by

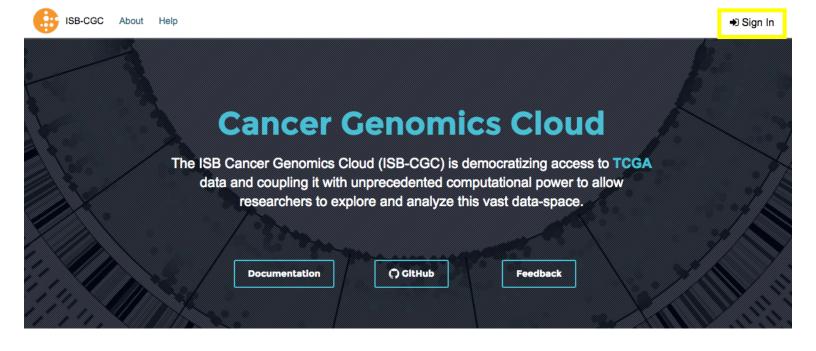
The ISB Cancer Genomics Cloud







https://isb-cgc.appspot.com



Your Dashboard > Saved Cohorts > **TCGA HNSC** New Workbook Comments Duplicate Delete View Files Download IDs Share Shared With (0) **Selected Filters** SAMP:Project: TCGA SAMP:Study: HNSC **Details** Total Number of Samples: 1123 1. Log into the system Total Number of Participants: 528 2. Create a cohort of TCGA HNSC Samples Your Permissions: OWNER Revision History:

There is no revision history.

TCGA CESC New Workbook Comments Duplicate Delete View Files Download IDs Share Shared With (0) **Selected Filters** SAMP:Project: TCGA SAMP:Study: CESC **Details** Total Number of Samples: 621 3. Create a cohort of TCGA CESC Samples Total Number of Participants: 308 Your Permissions: OWNER Revision History: There is no revision history.

Your Dashboard > Saved Cohorts >

- 4. Create a gene list with the following genes:
- ERBB2
- FN1
- PVT1
- SERPINB11
- RAD51L1
- TMPRSS3

Your Dashboard > Saved Gene Favorites >

Gene Favorite Details





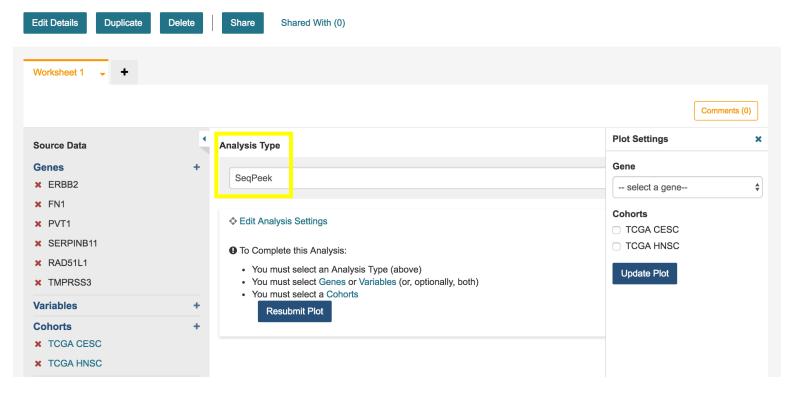
5. Apply gene list and cohorts to new workbook.

Edit Details Duplicate Share Shared With (0) Worksheet 1 Comments (0) **Plot Settings** × Source Data Analysis Type Genes + Gene SeqPeek × ERBB2 * -- select a gene--× FN1 Edit Analysis Settings × PVT1 ☐ TCGA CESC **★** SERPINB11 ☐ TCGA HNSC ① To Complete this Analysis: × RAD51L1 • You must select an Analysis Type (above) Update Plot You must select Genes or Variables (or, optionally, both) × TMPRSS3 • You must select a Cohorts **Variables** Resubmit Plot X TCGA CESC X TCGA HNSC



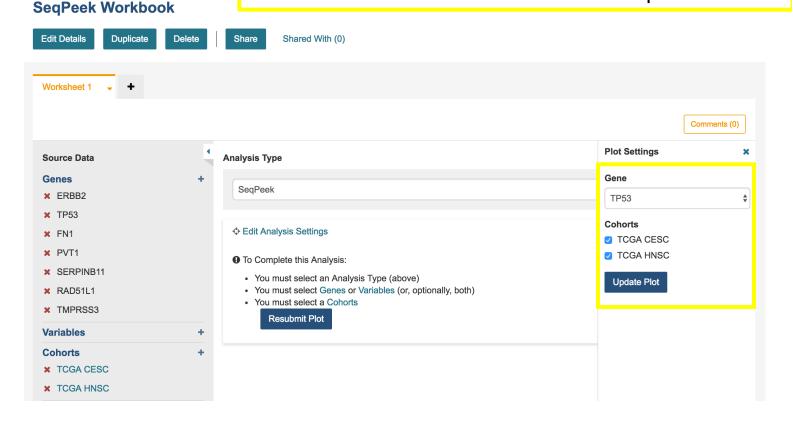
6. Select SeqPeek analysis type.

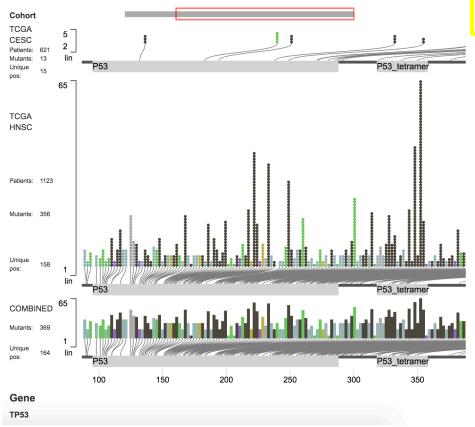
SeqPeek Workbook



Your Dashboard > Saved Workbooks >

7. Select Gene TP53 and cohorts and Update Plot.





8. Explore resulting plot.



- Nonsense_Mutation
- Silent
- Frame_Shift_Del
- Frame_Shift_Ins
- Missense_Mutation
- In_Frame_Ins
- In_Frame_Del