

#Obtaining the list of genes that vary according to NEDD8 and NAE1 expression levels

First, the Liver Hepatocellular Carcinoma (TCGA, PanCancer Atlas) (372 samples) dataset is selected in cBioPortal.

Filters:

- Cancer Type Detailed: Hepatocellular Carcinoma
- Tumor Type: Hepatocellular Carcinoma
- Subtype: LIHC

Add gene specific charts, with the following specification

NEDD8: mRNA Expression, RSEM (Batch normalized from Illumina HiSeq\_RNASeqV2)

NAE1: mRNA Expression, RSEM (Batch normalized from Illumina HiSeq\_RNASeqV2)

These charts can be found at the bottom of the page, where they are grouped by median. Thus, under the mRNA tab, there are separate lists of high and low genes according to NEDD8 and NAE1.

Please select significant only