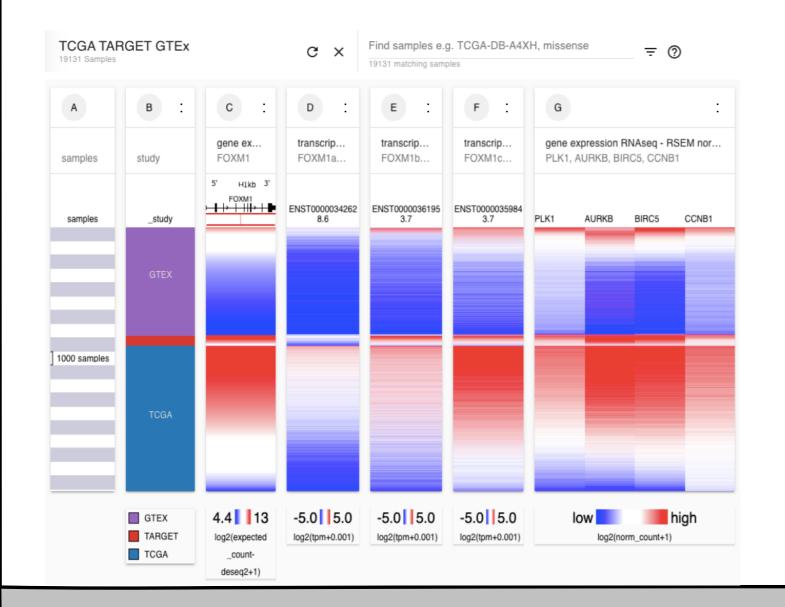
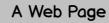




Ⅲ ♣ ७ ७









19131 Samples

Find samples e.g. TCGA-DB-A4XH, missense

19131 matching samples

- ②





Choose a column

Choose a column |-

Back







19131 Samples

Find samples e.g. TCGA-DB-A4XH, missense

19131 matching samples

= ②







Choose a column



Choose a column -

Study

FOXM1 gene expression RNA ...

FOXM1a/ENST0000034262 ...

FOXM1b/ENST00000361953 ...

FOXM1c/ENST0000035984 ...

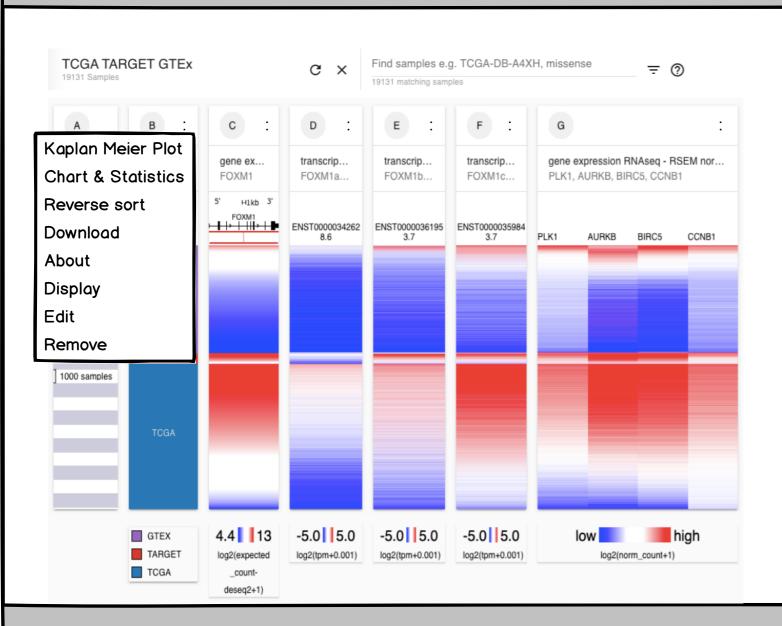
PLK1, AURKB, BIRC5, CCNB1 ...

Back





Ⅲ ♣ ₽ 8











Find samples e.g. TCGA-DB-A4XH, missense

19131 matching samples

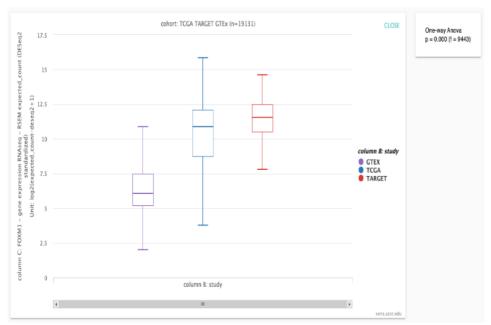
= ②

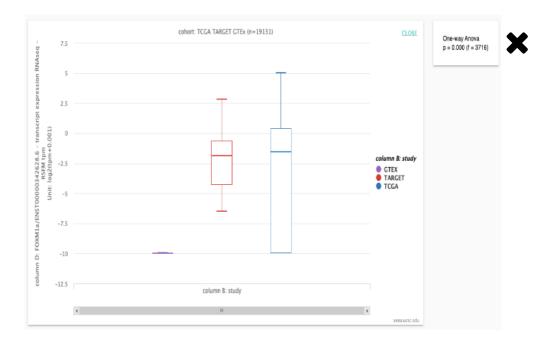
























Find samples e.g. TCGA-DB-A4XH, missense

19131 matching samples



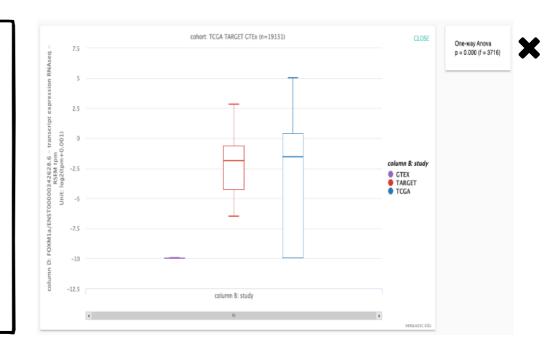


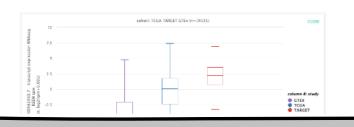




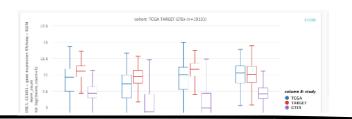
Suggested graphs

FOXM1 gene expression RNAseq - RSEM expected_count (DESeq2 standardized) by Study













TCGA TARGET GTEX 19131 Samples

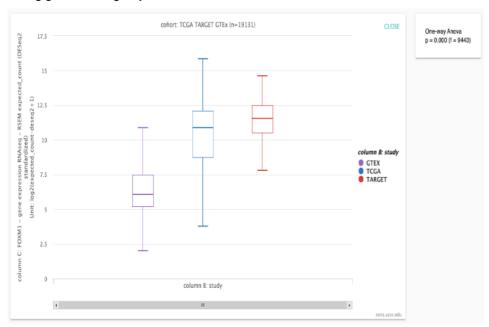
G >

Find samples e.g. TCGA-DB-A4XH, missense

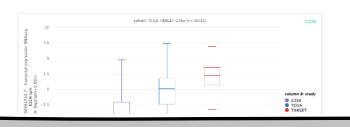
- ②



Suggested graphs



ENST00000342628.6
Transcript expression RNAseq
- RSEM expected_count
(DESeq2 standardized)
by
Study













19131 Samples

G X

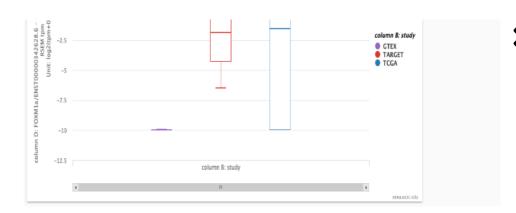
Find samples e.g. TCGA-DB-A4XH, missense

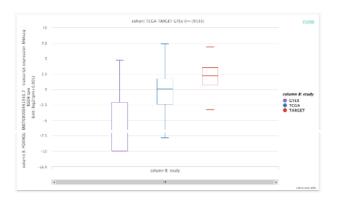
19131 matching samples

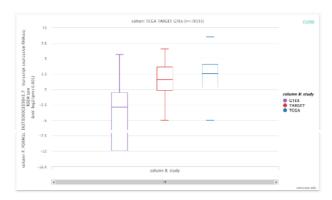
- ?

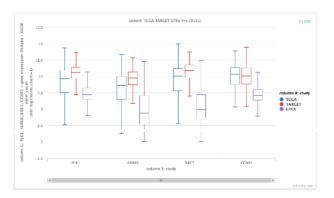


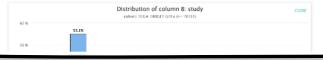
















19131 Samples



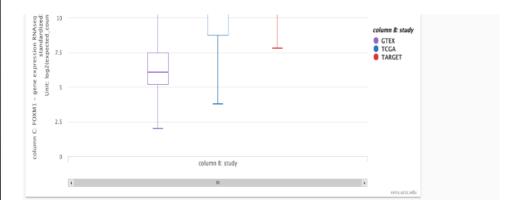
Find samples e.g. TCGA-DB-A4XH, missense 19131 matching samples

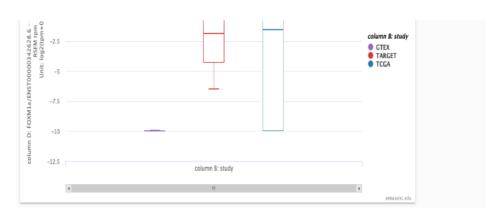
= ②



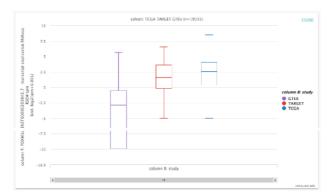


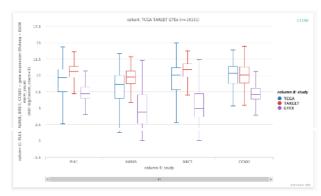






ENST00000361953.7 Transcript expression RNAseq - RSEM expected_count (DESeq2 standardized) by Study











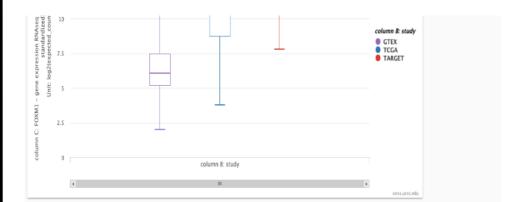
19131 Samples

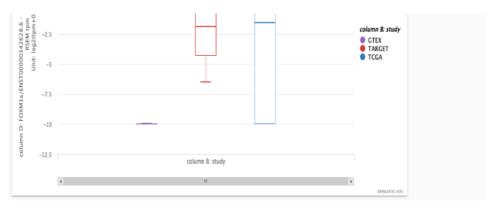


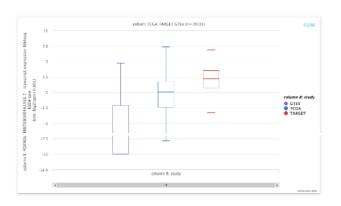
Find samples e.g. TCGA-DB-A4XH, missense



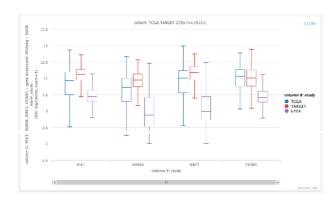








ENST00000359843.7
Transcript expression RNAseq
- RSEM expected_count
(DESeq2 standardized)
by
Study









19131 Samples

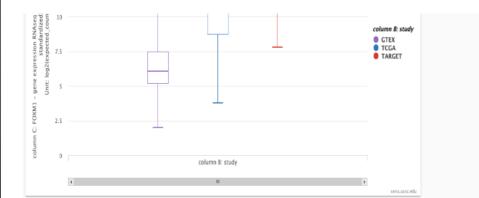


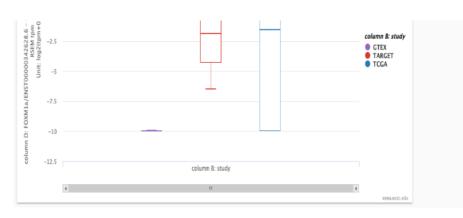
Find samples e.g. TCGA-DB-A4XH, missense 19131 matching samples

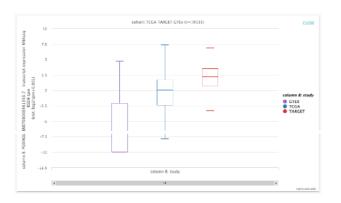
- ②

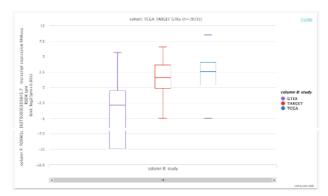




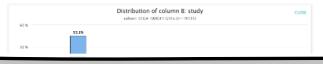








PLK1, AURKB, BIRC5, CCNB1 gene expression RNAseq -RSEM expected_count (DESeq2 standardized) by Study







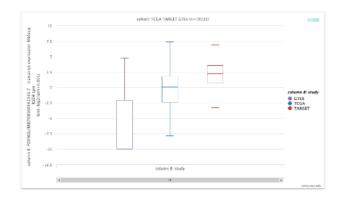
19131 Samples

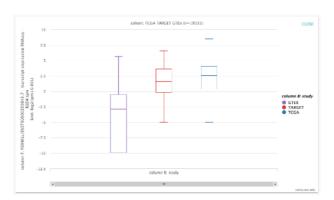


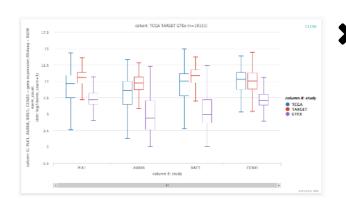
Find samples e.g. TCGA-DB-A4XH, missense
19131 matching samples

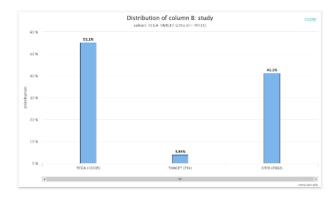
















19131 Samples

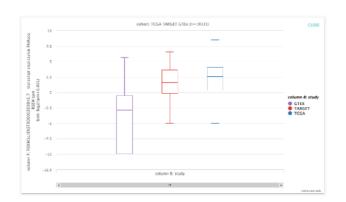


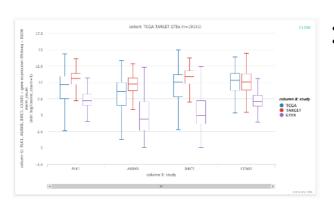
Find samples e.g. TCGA-DB-A4XH, missense 19131 matching samples

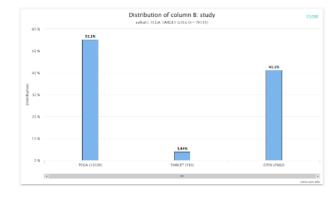




ENST00000361953.7 Transcript expression RNAseq - RSEM expected_count (DESeq2 standardized) by Study











19131 matching samples



TCGA TARGET GTEX

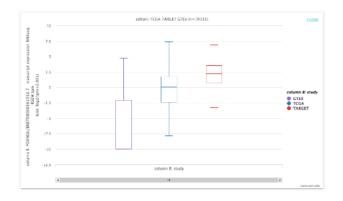
19131 Samples

G X

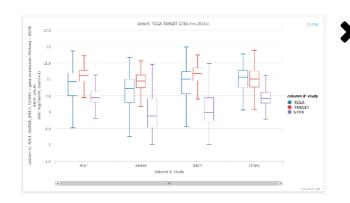
Find samples e.g. TCGA-DB-A4XH, missense $\,$

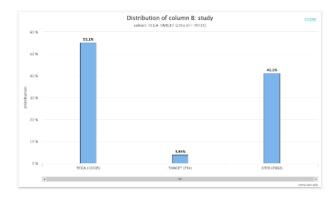
= ②





ENST00000359843.7
Transcript expression RNAseq
- RSEM expected_count
(DESeq2 standardized)
by
Study









19131 Samples

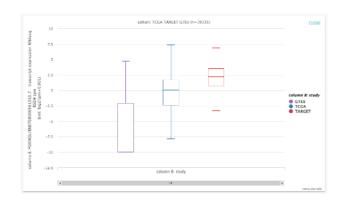


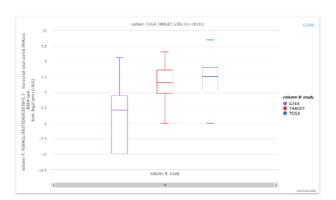
Find samples e.g. TCGA-DB-A4XH, missense

19131 matching samples

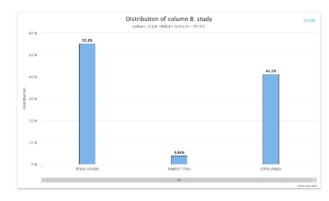
= ②







PLK1, AURKB, BIRC5, CCNB1 gene expression RNAseq -RSEM expected_count (DESeq2 standardized) by Study



- ②





TCGA TARGET GTEX

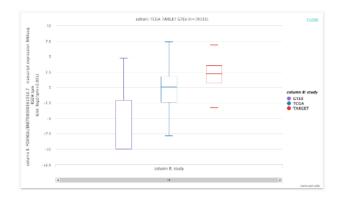
19131 Samples

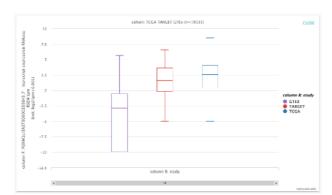
G X

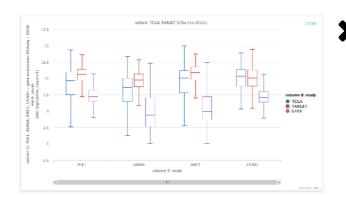
Find samples e.g. TCGA-DB-A4XH, missense

19131 matching samples







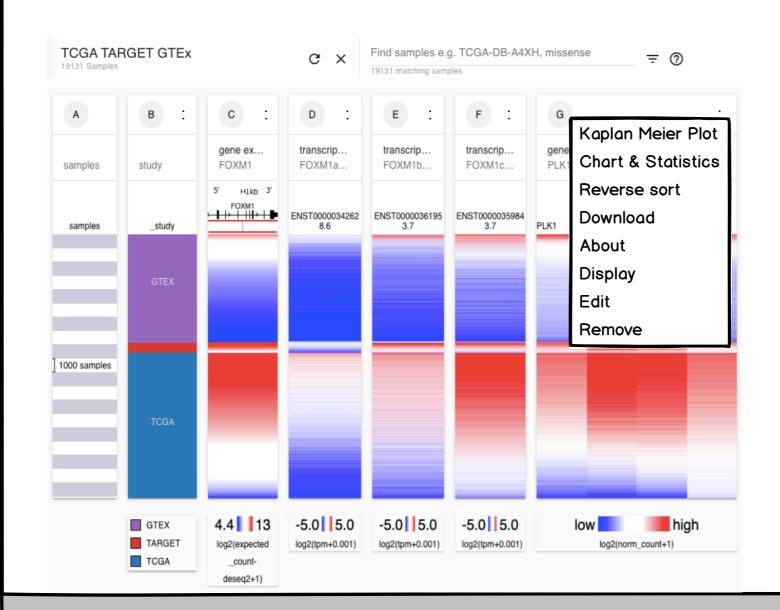


Study Histogram





Ⅲ ♣ ₽ 8









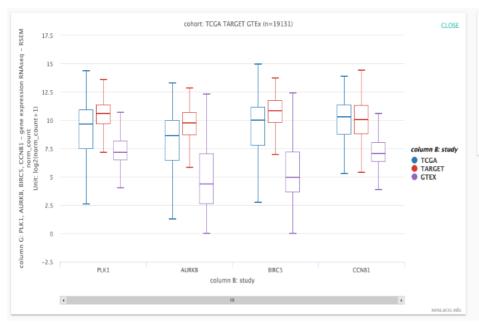


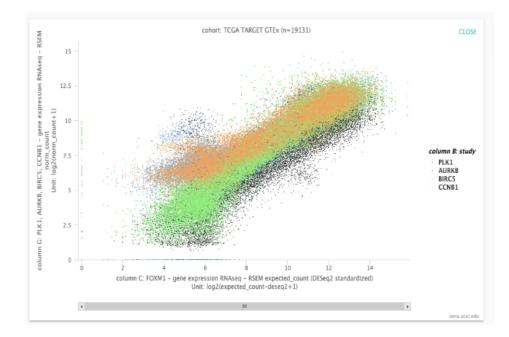
Find samples e.g. TCGA-DB-A4XH, missense 19131 matching samples

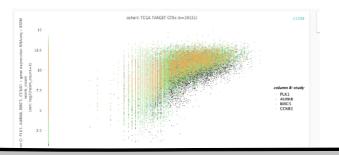




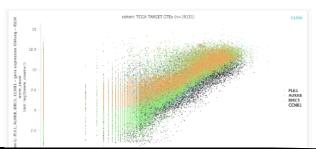
Suggested graphs

















TCGA TARGET GTEX 19131 Samples

Find samples e.g. TCGA-DB-A4XH, missense

19131 matching samples

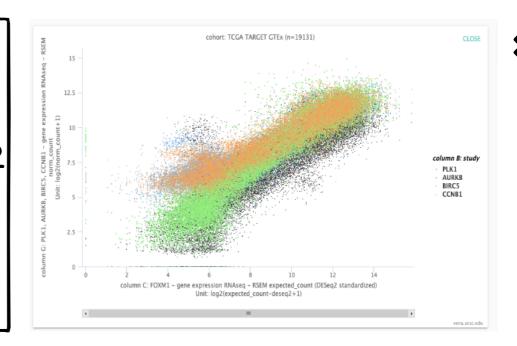


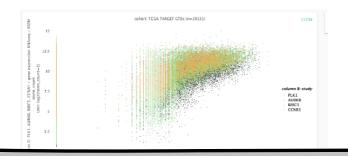


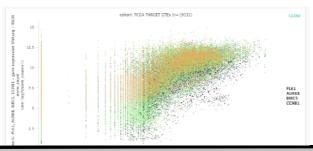


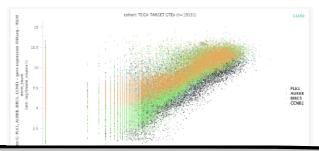
Suggested graphs

PLK1, AURKB, BIRC5, CCNB1 gene expression RNAseq -RSEM expected_count (DESeq2 standardized) Study













TCGA TARGET GTEX 19131 Samples

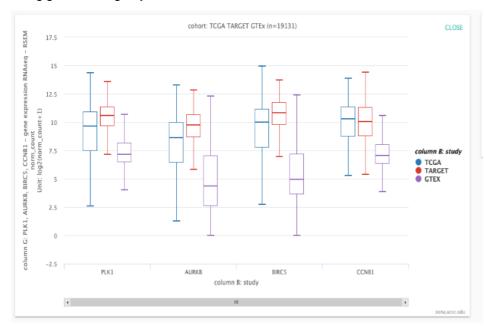


Find samples e.g. TCGA-DB-A4XH, missense

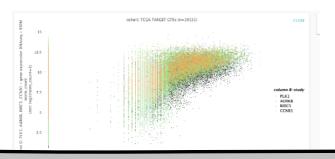


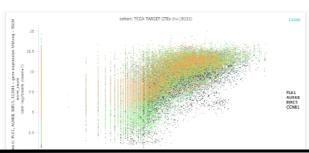


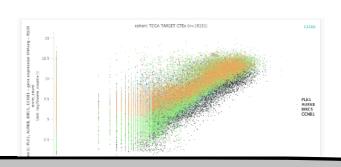
Suggested graphs



PLK1, AURKB, BIRC5, CCNB1
gene expression RNAseq RSEM expected_count (DESeq2
standardized)
by
ENST00000342628.6 Transcript
expression RNAseq - RSEM
expected_count (DESeq2 ...













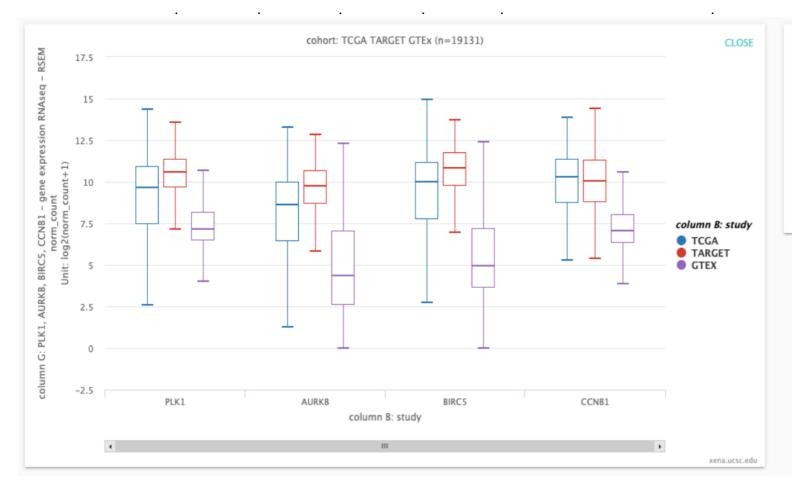
19131 Samples

G X Find samples e.g. TCGA-DB-A4XH, missense

19131 matching samples

= ②





One-way Anova PLK1 p = 0.000 (f = 2257) AURKB p = 0.000 (f = 4424)BIRC5 p = 0.000 (f = 6587) CCNB1 p = 0.000 (f = 6332)

Make another graph

Flip X & Y axes

Advanced Options