Xena helps you answer questions like:

- What is the pattern between my two genes?
- What is the relationship between this frequent mutation and expression?
- Where does my sample fall within the larger genomics space?
- How did drug treatment change expression of a gene in my cell line?

We offer:

- Tutorials
- Forums
- Mailing lists
- Help guides
- FAQ

Data types: SNPs, Indels, SVs, expression, CNV, methylation, sample annotations (age, tumor type, clustering results, etc)

Watch to learn more

http://xena.ucsc.edu