Tutorial: Interactive Next-Generation Clustered Heat Maps

This slide deck introduces many, but by no means all, features of Next-Generation clustered Heat Maps (NG-CHMs). For concreteness, it's based on:

- A TCGA mRNA expression (rows) vs. samples (columns) NG-CHM for breast cancer (as part of our TCGA NG-CHM Compendium: http://bioinformatics.mdanderson.org/TCGA/NGCHMPortal/
- The specific map:http://bioinformatics.mdanderson.org/chm/chm.html?name=tcga_rnaseq_brca_v1.0_gene_sample&collectionHome=http%3A%2F%2Fbioinformatics.mdanderson.org%2FTCGA%2FNGCHMPortal%3Fview%3D0%26p0DiseaseInput%3Dbrca (too long, sorry)
- If you have a large monitor or two monitors, you might want to open the breast cancer map in a separate window to explore while you're going through the tutorial.
- Each index entry on slide 2 link to the corresponding slide – which links back to slide 2.
- Note: not all features are pertinent to any given map in the compendium (e.g., a gene-name link-out to GeneCards wouldn't be pertinent to for a copy number variation map.
- A more complete, detailed tutorial plus other videos can be found at http://bioinformatics.mdanderson.org/main/Navigating_Clustered_Heatmaps

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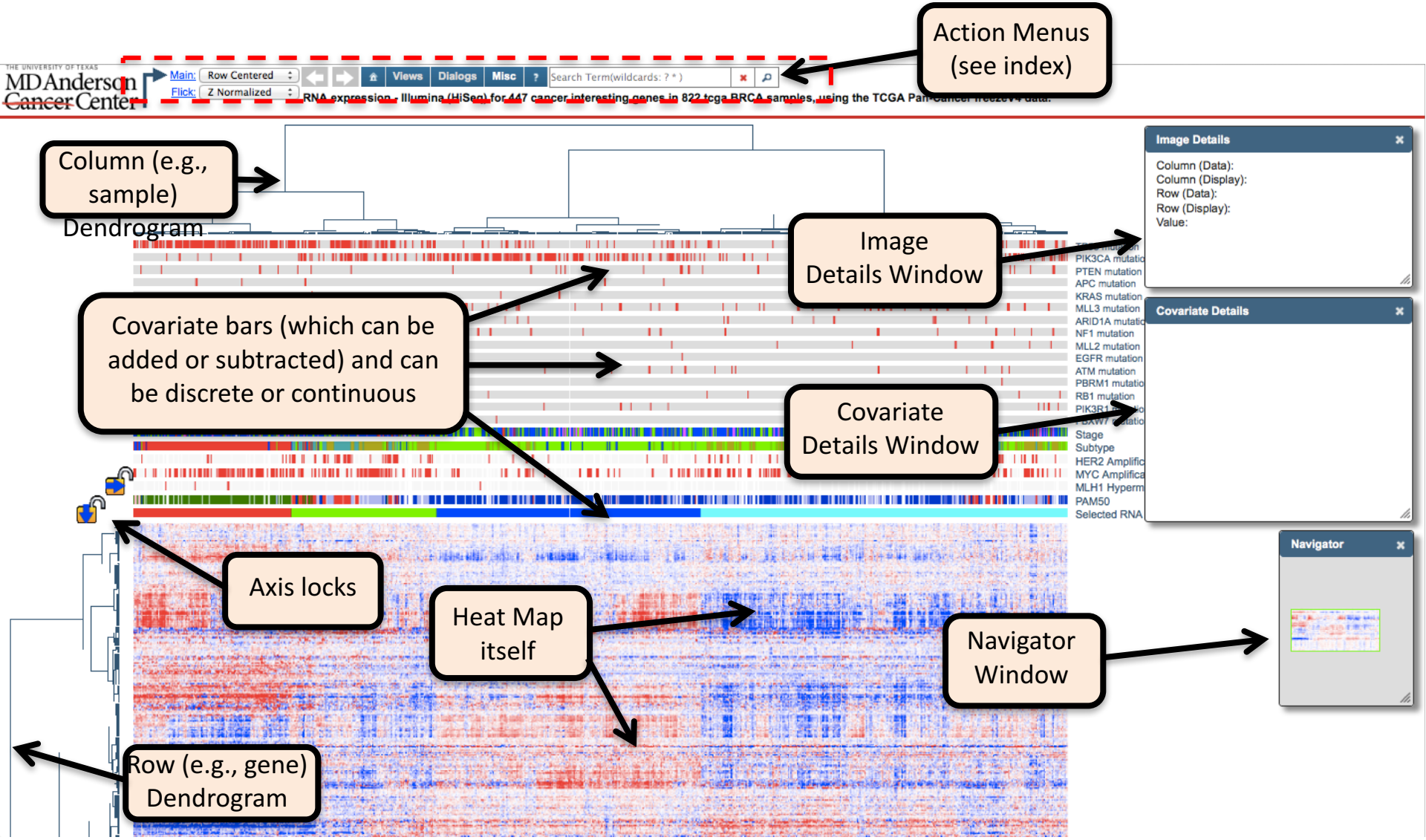
Main Components of the NG-CHM Viewer



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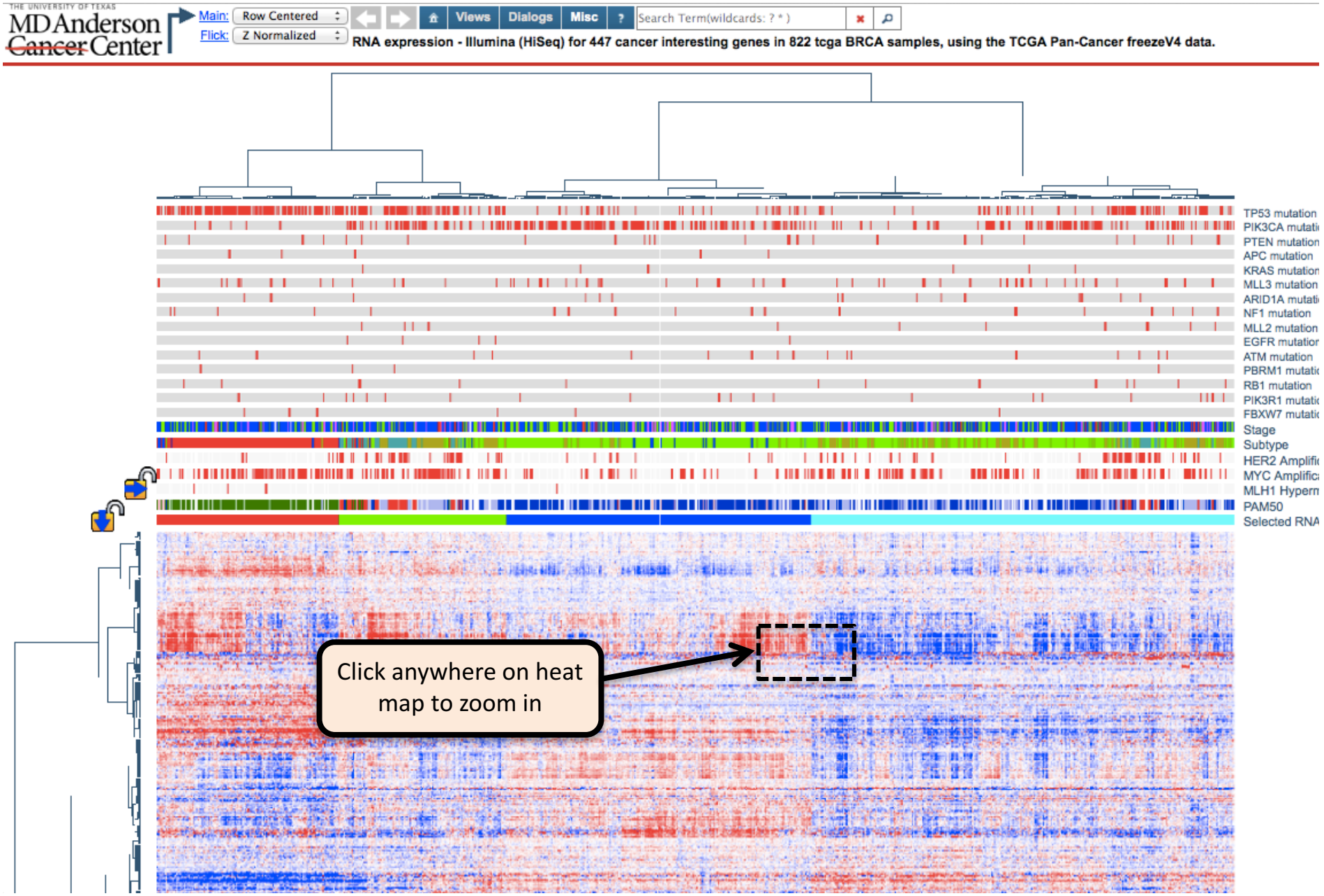

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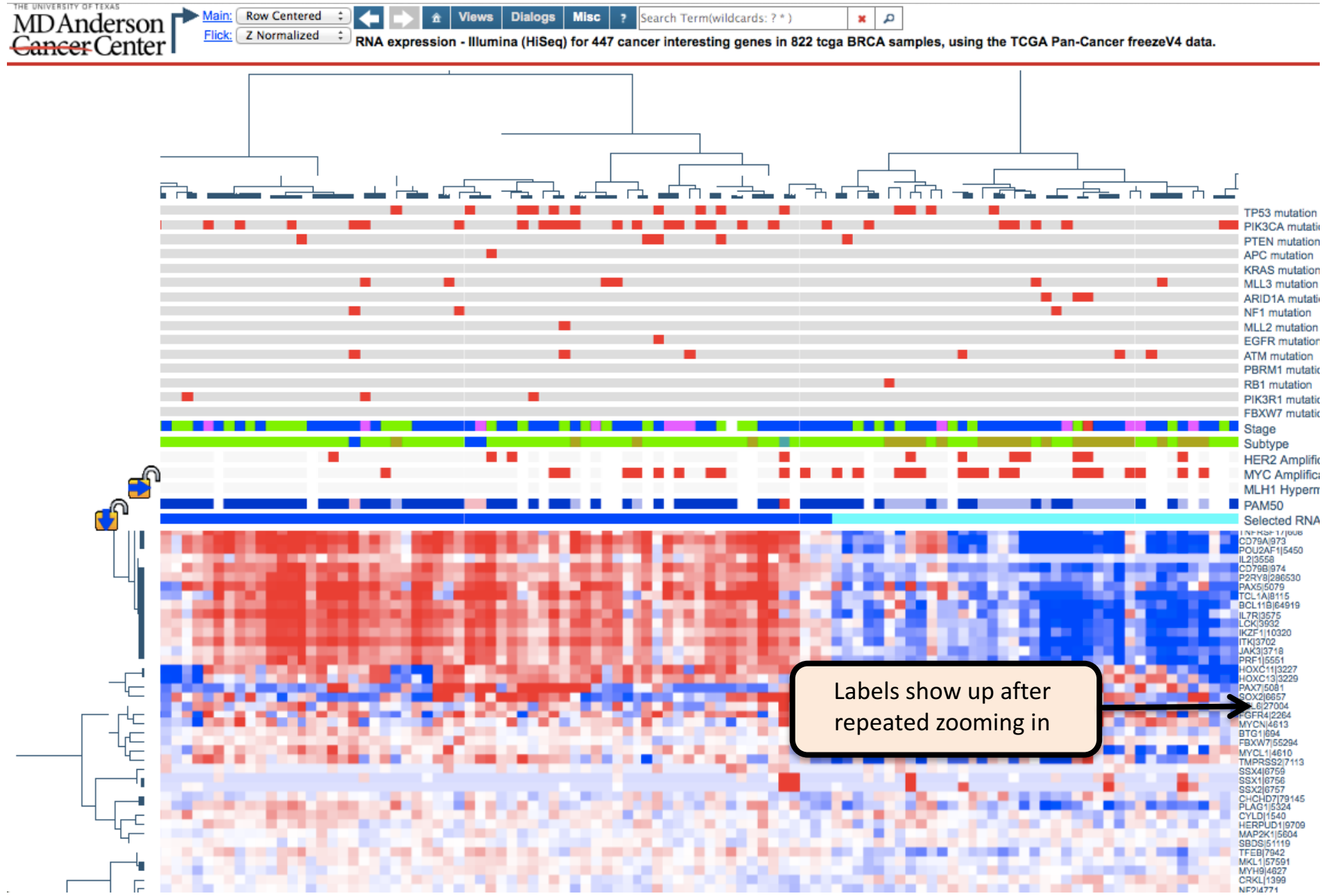

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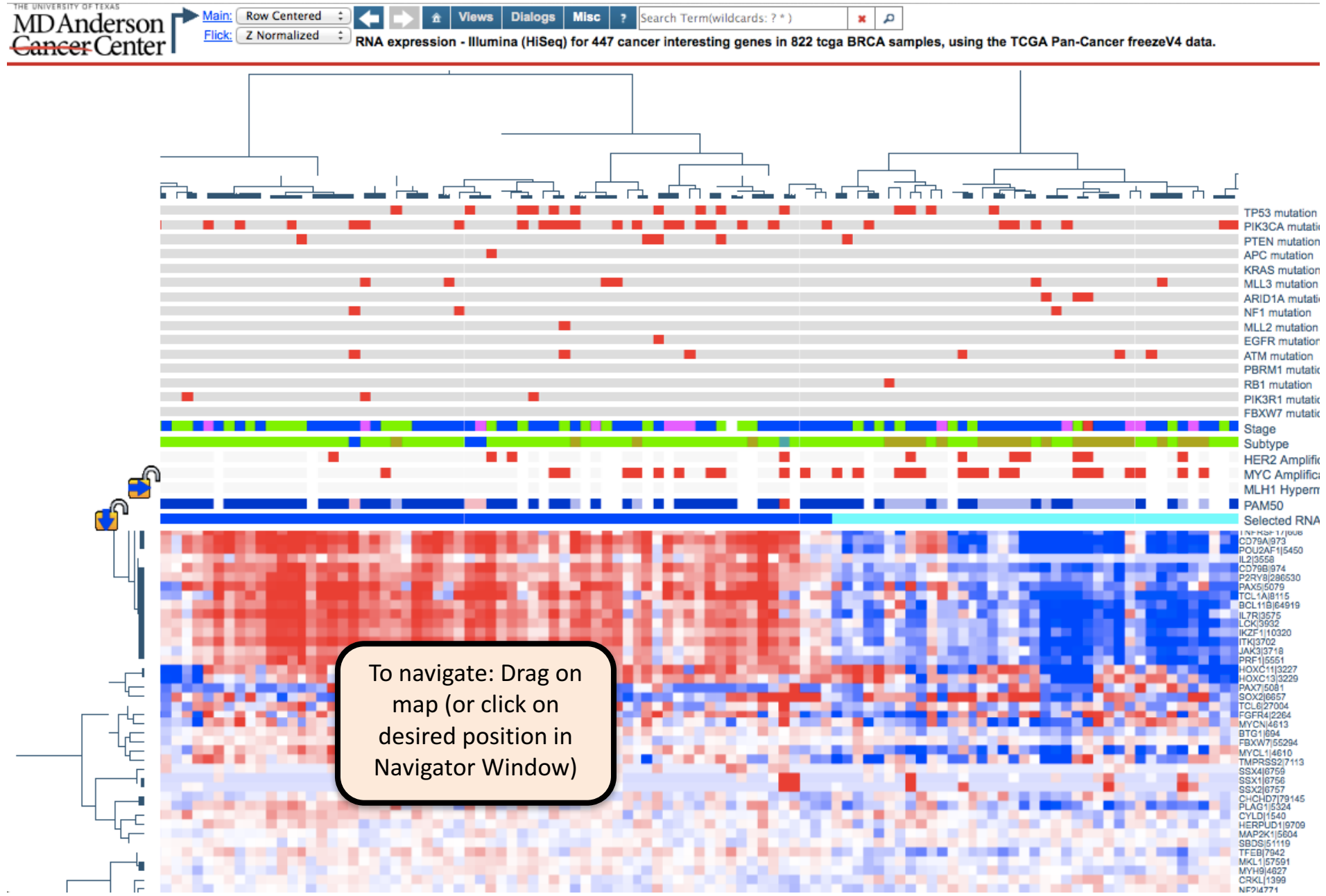

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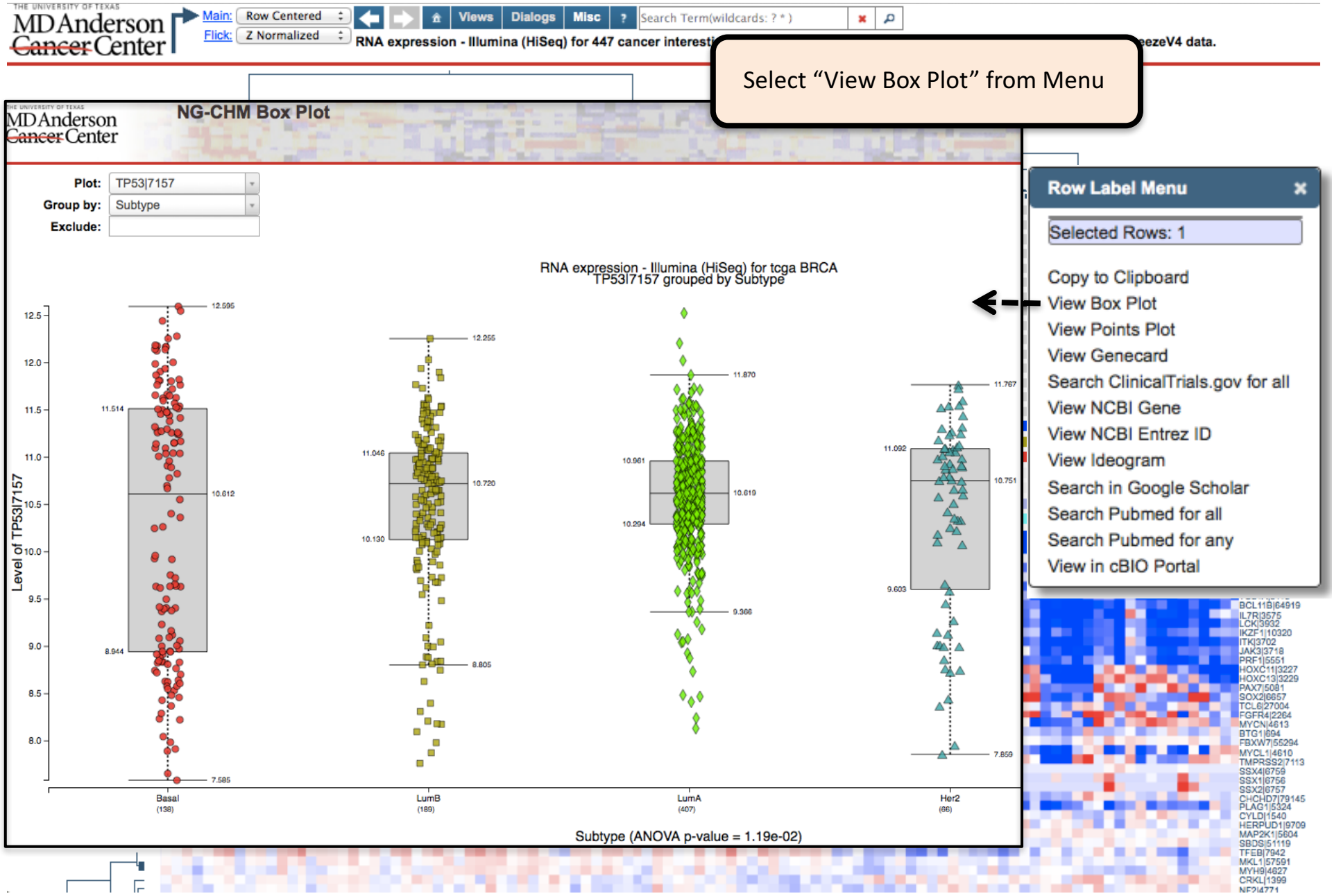

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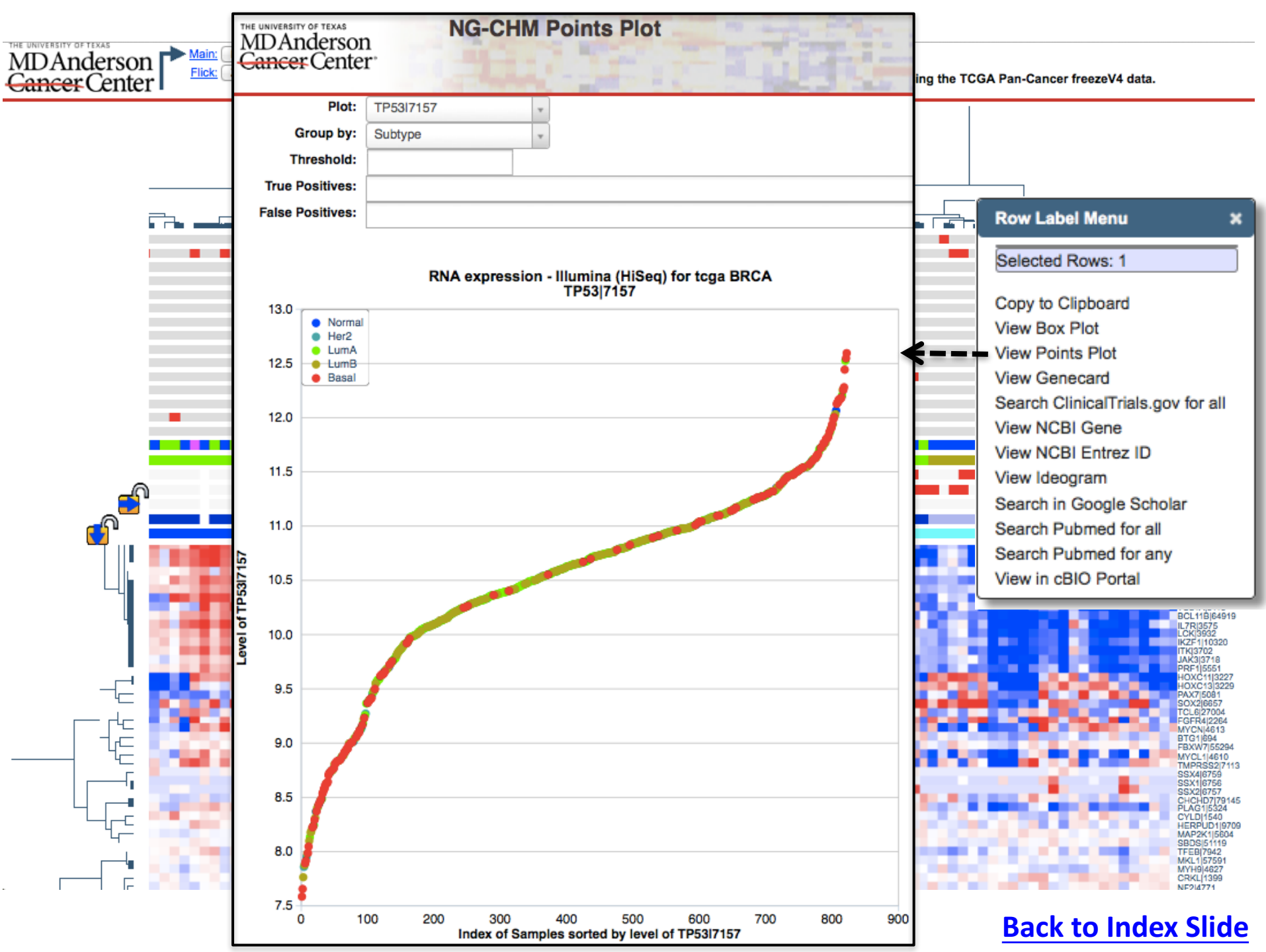

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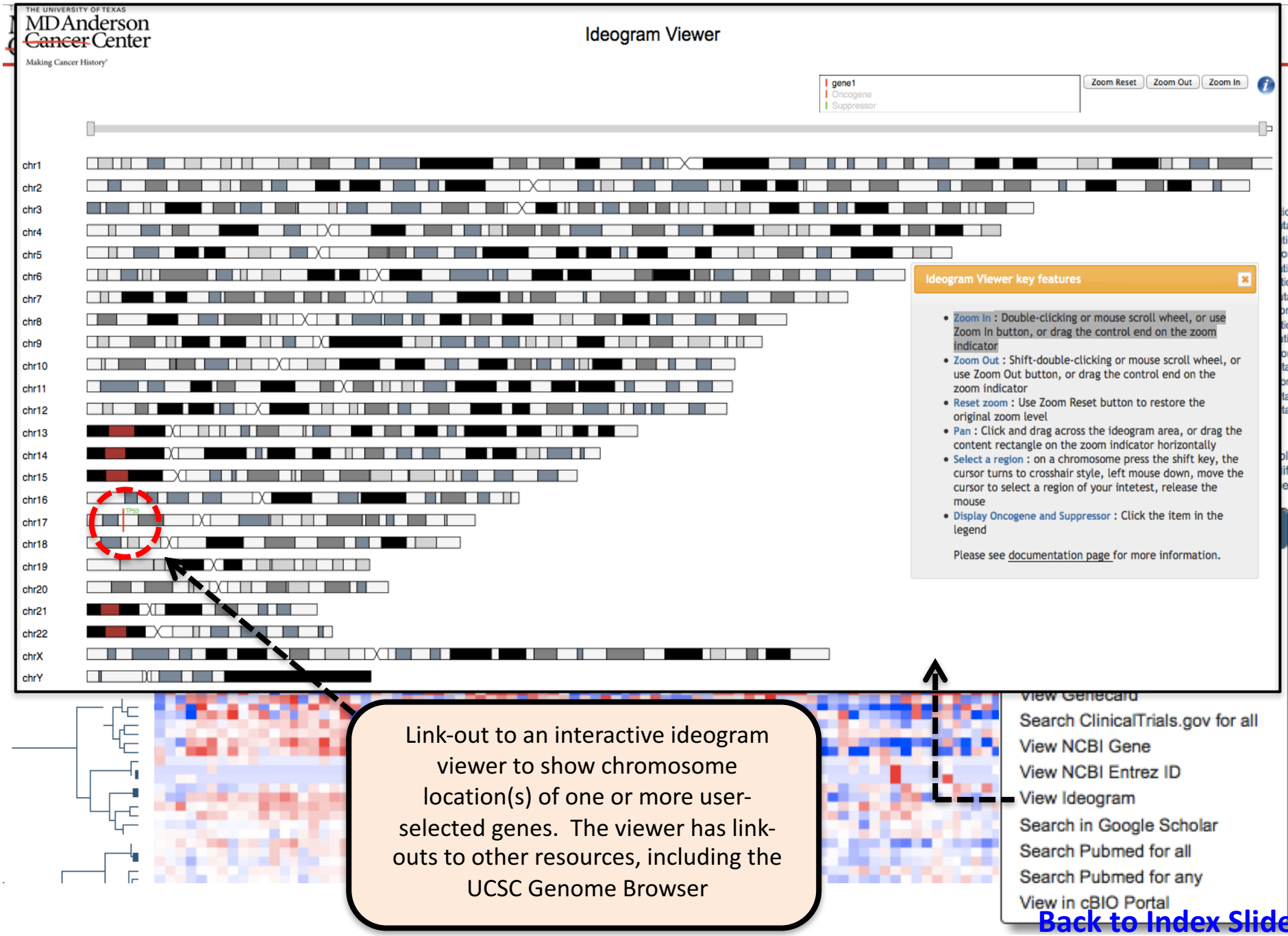

Link-out to an interactive ideogram viewer to show chromosome location(s) of one or more user-selected genes. The viewer has link-outs to other resources, including the UCSC Genome Browser

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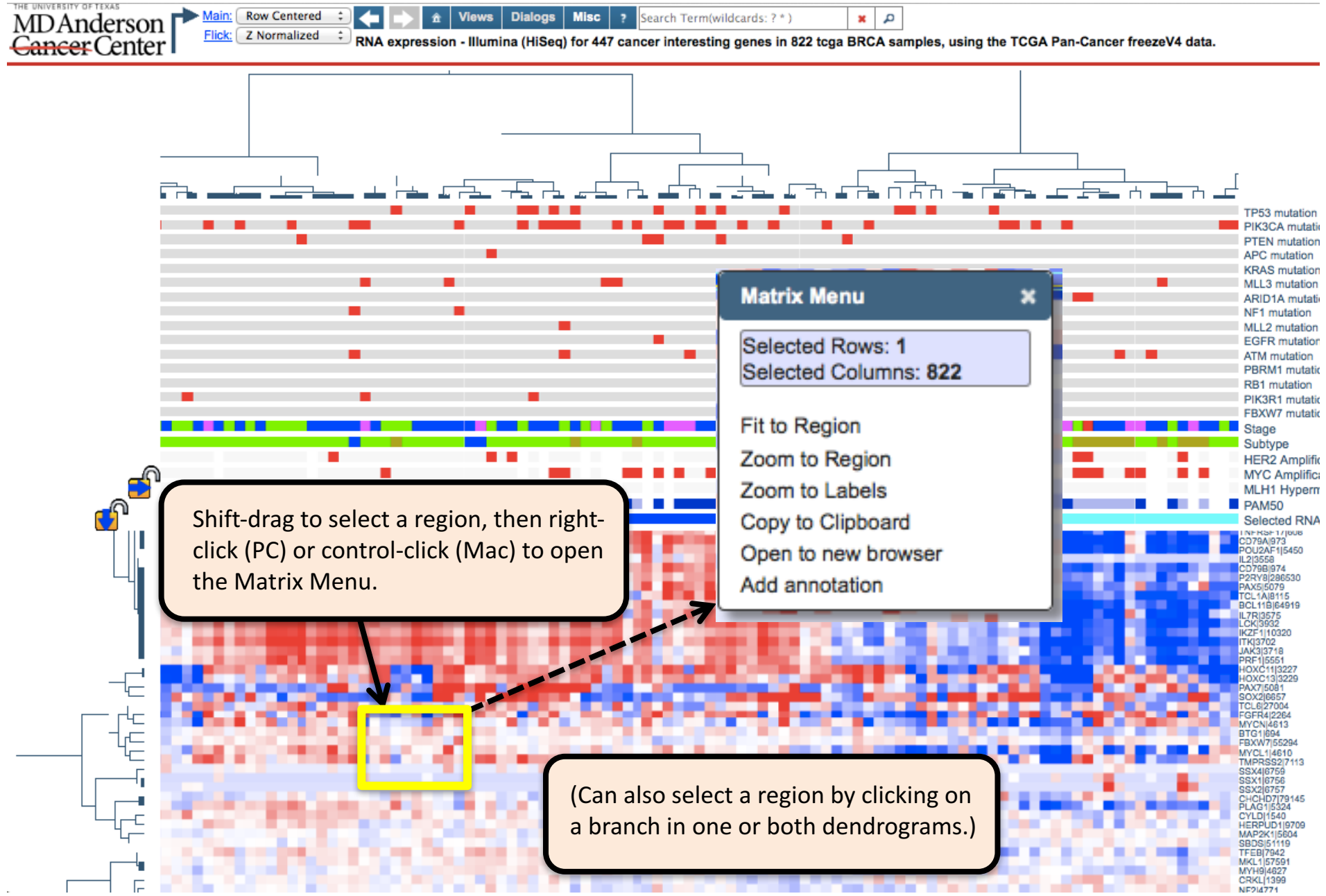

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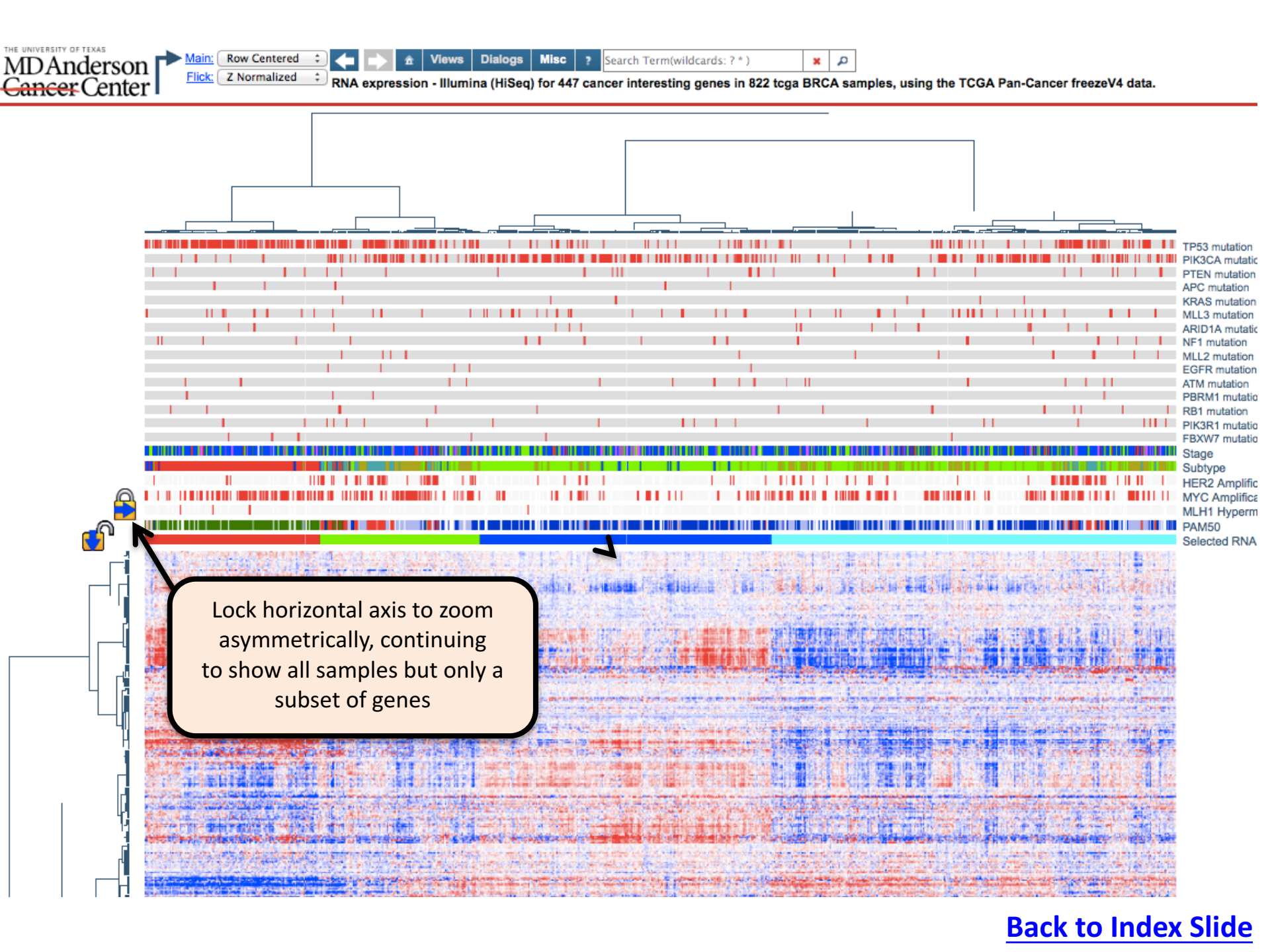

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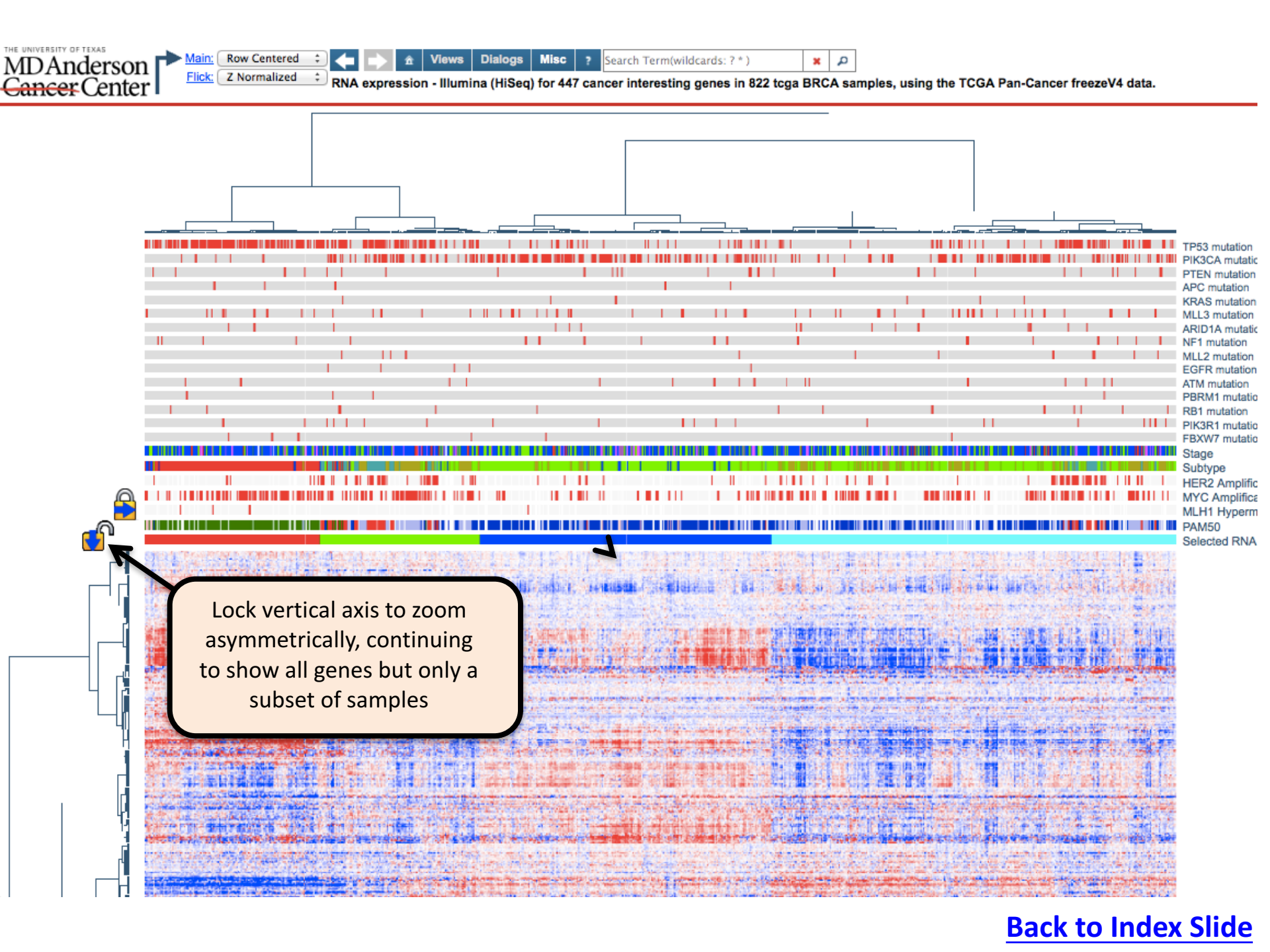

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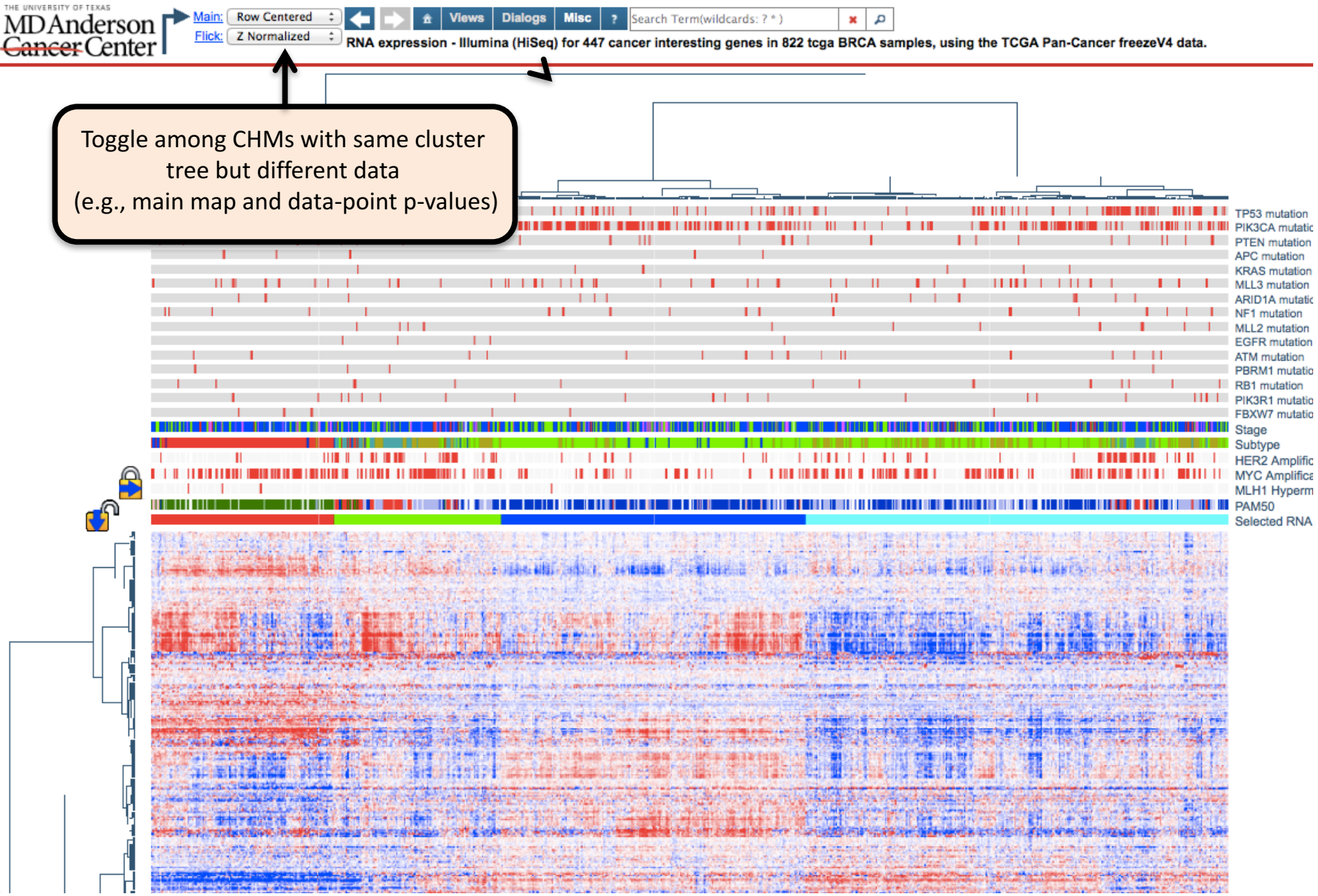

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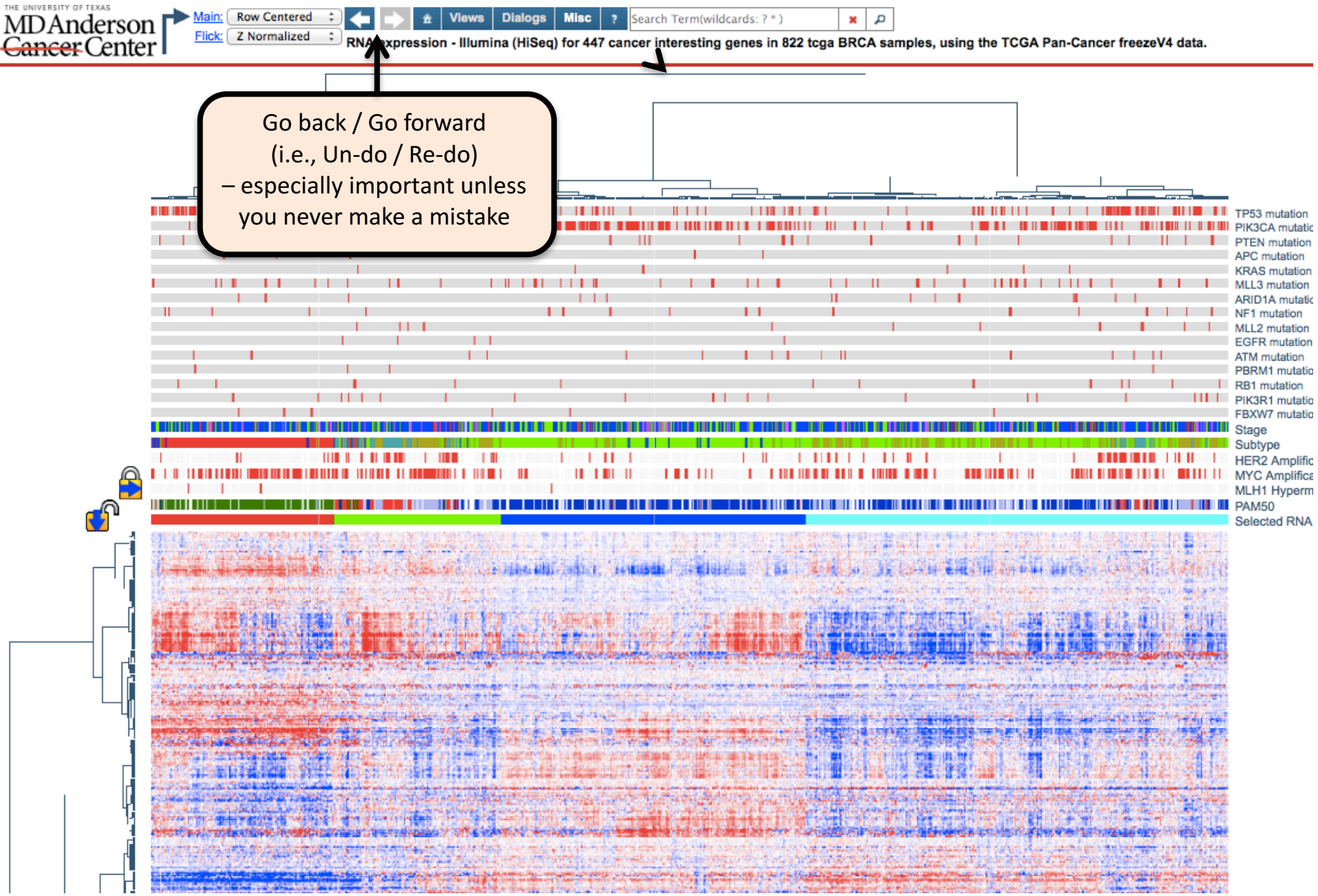

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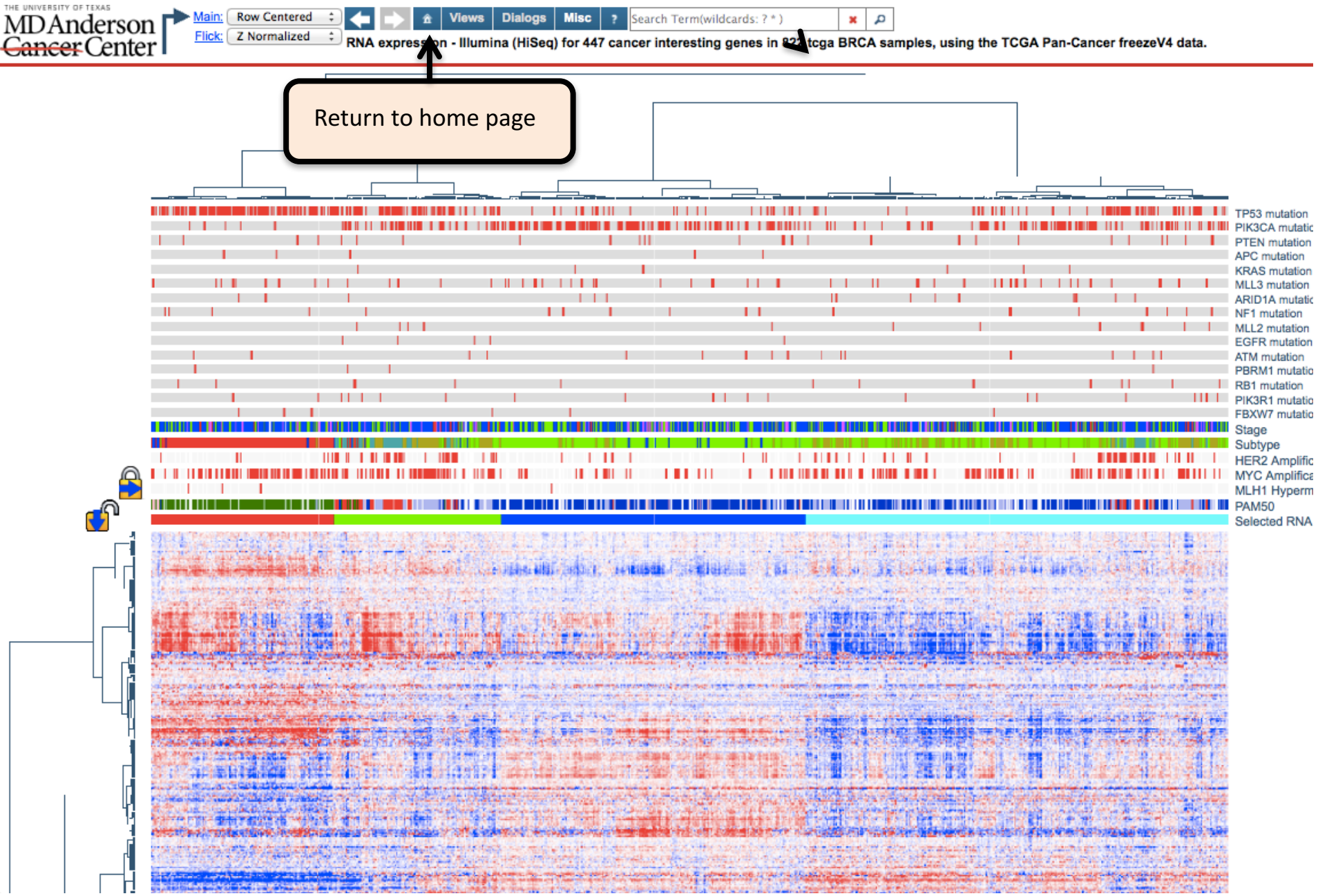

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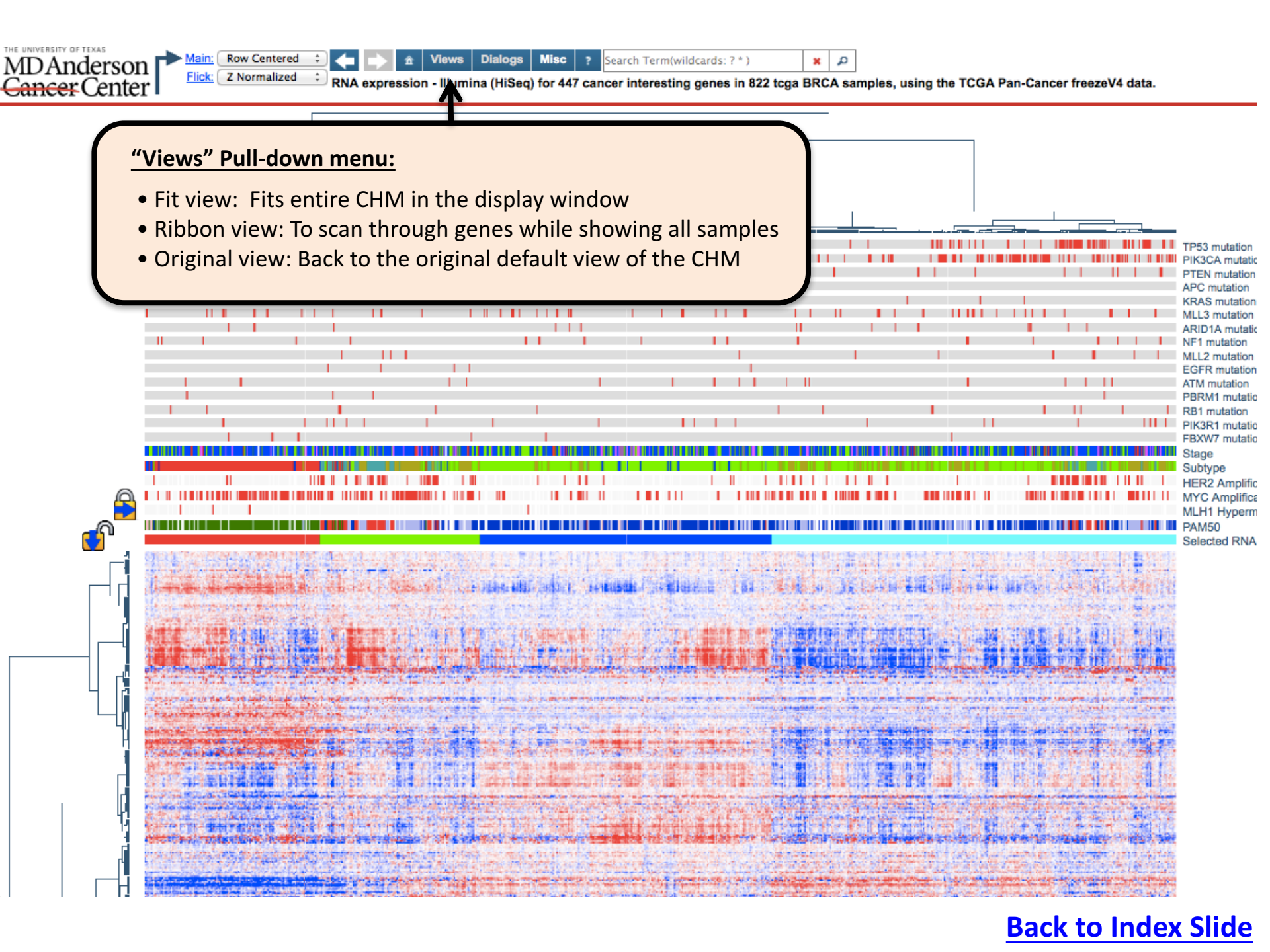

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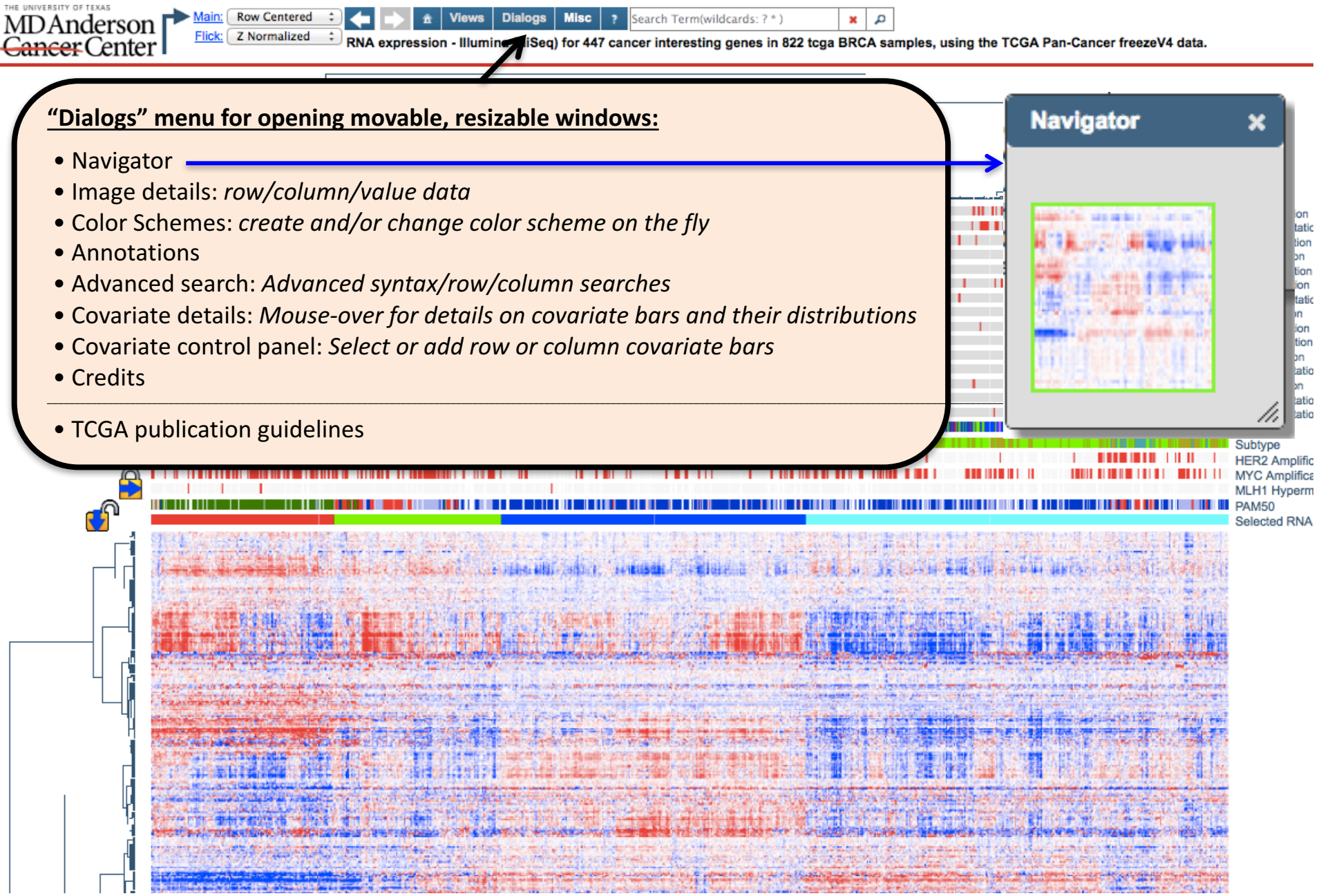

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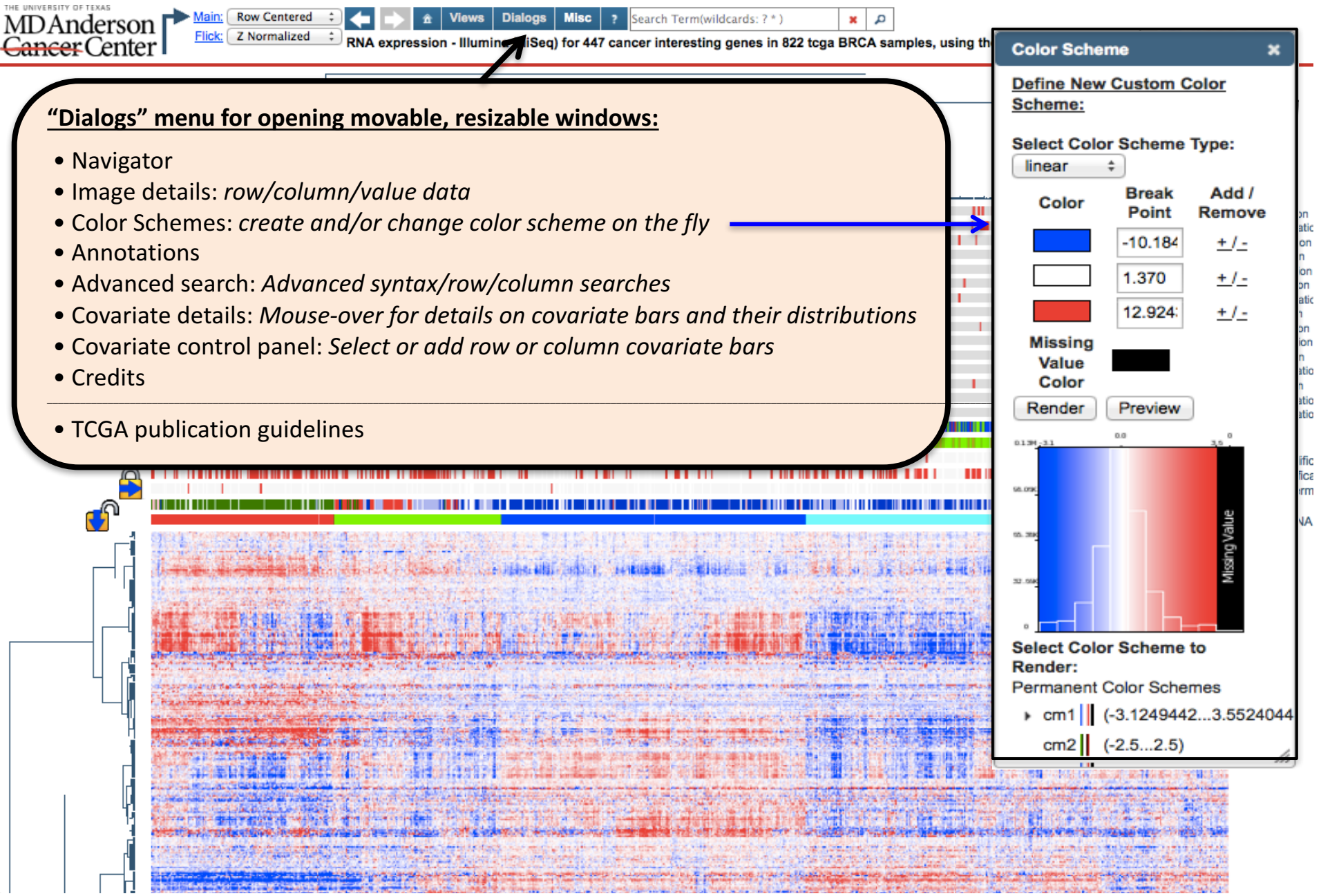

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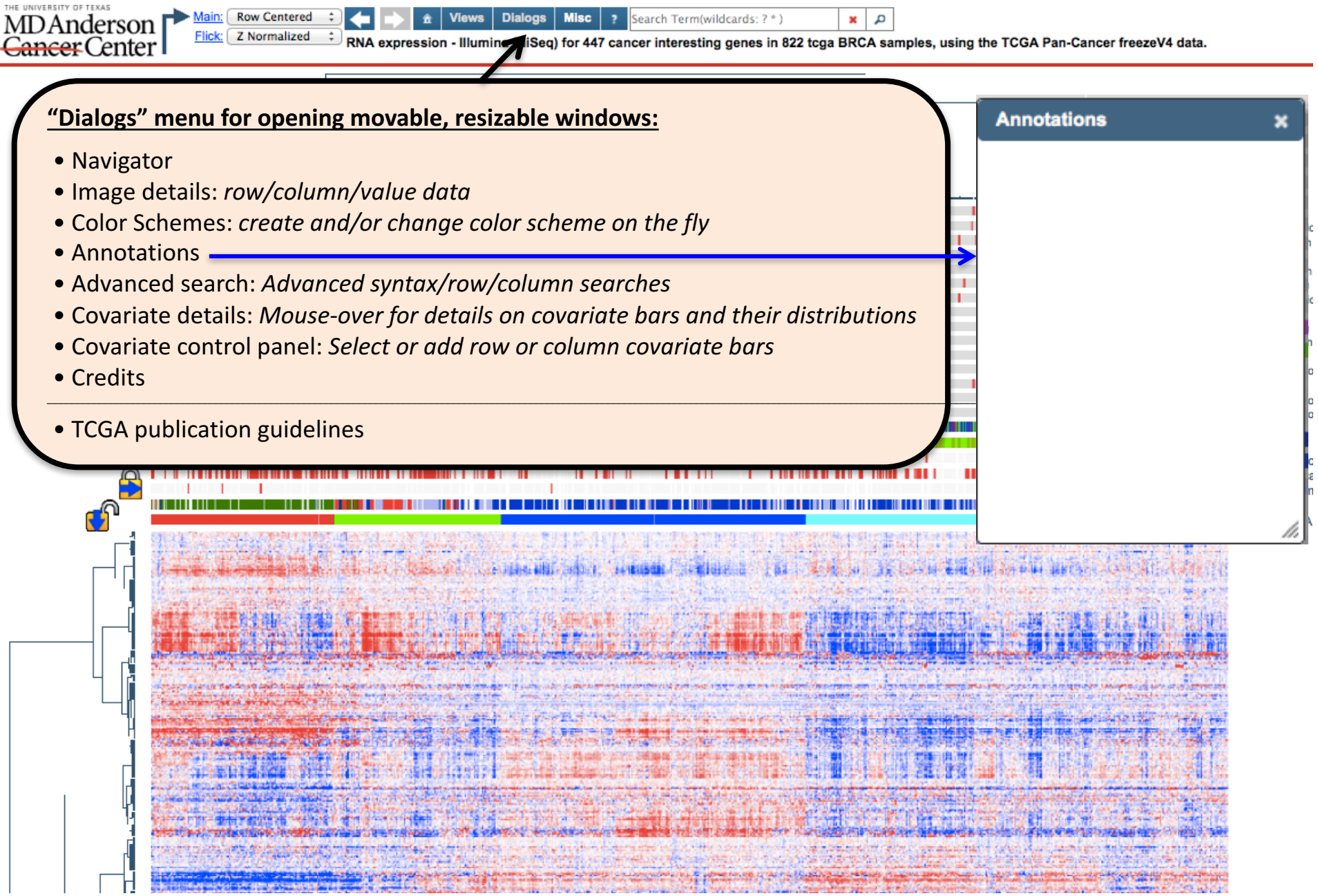

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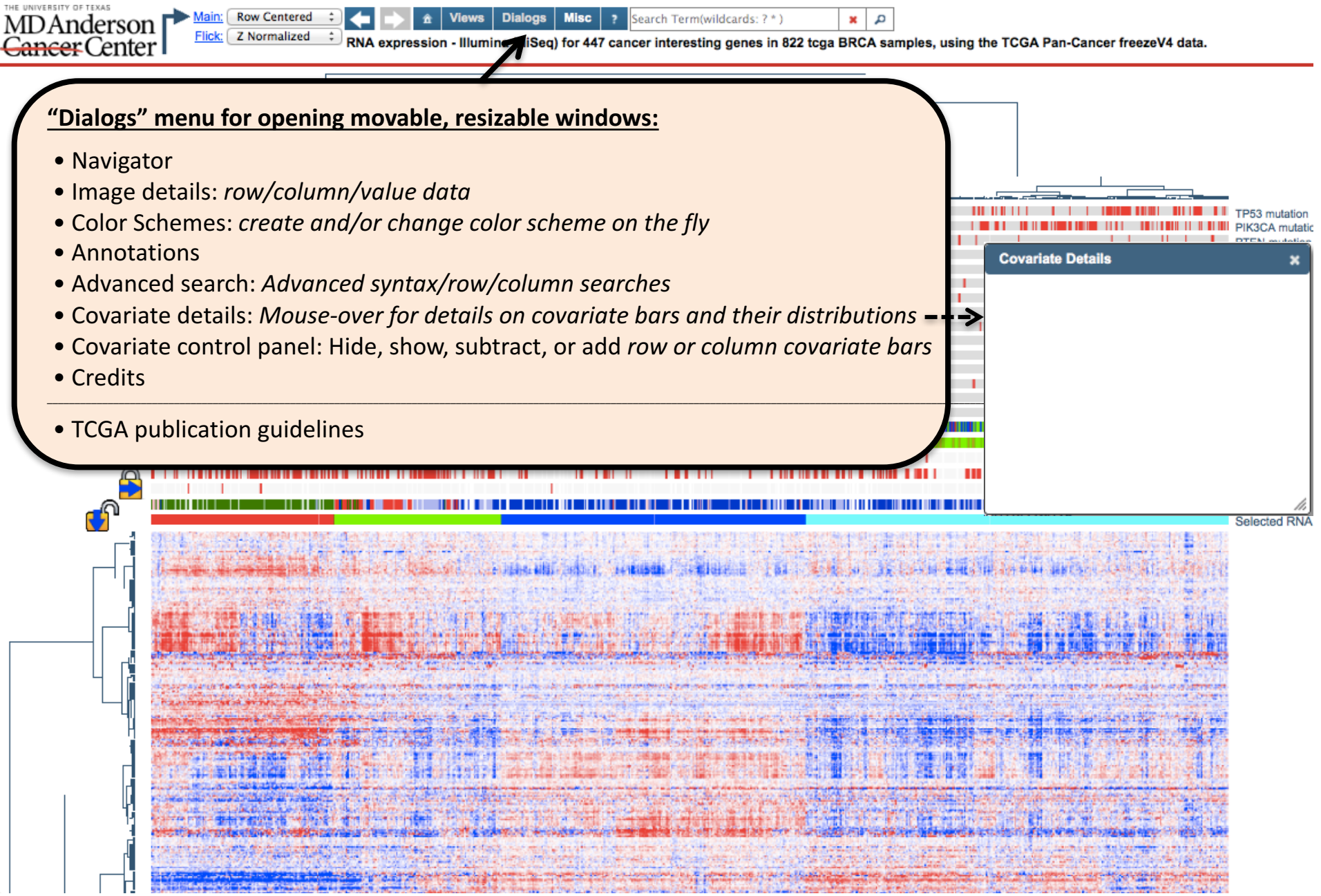

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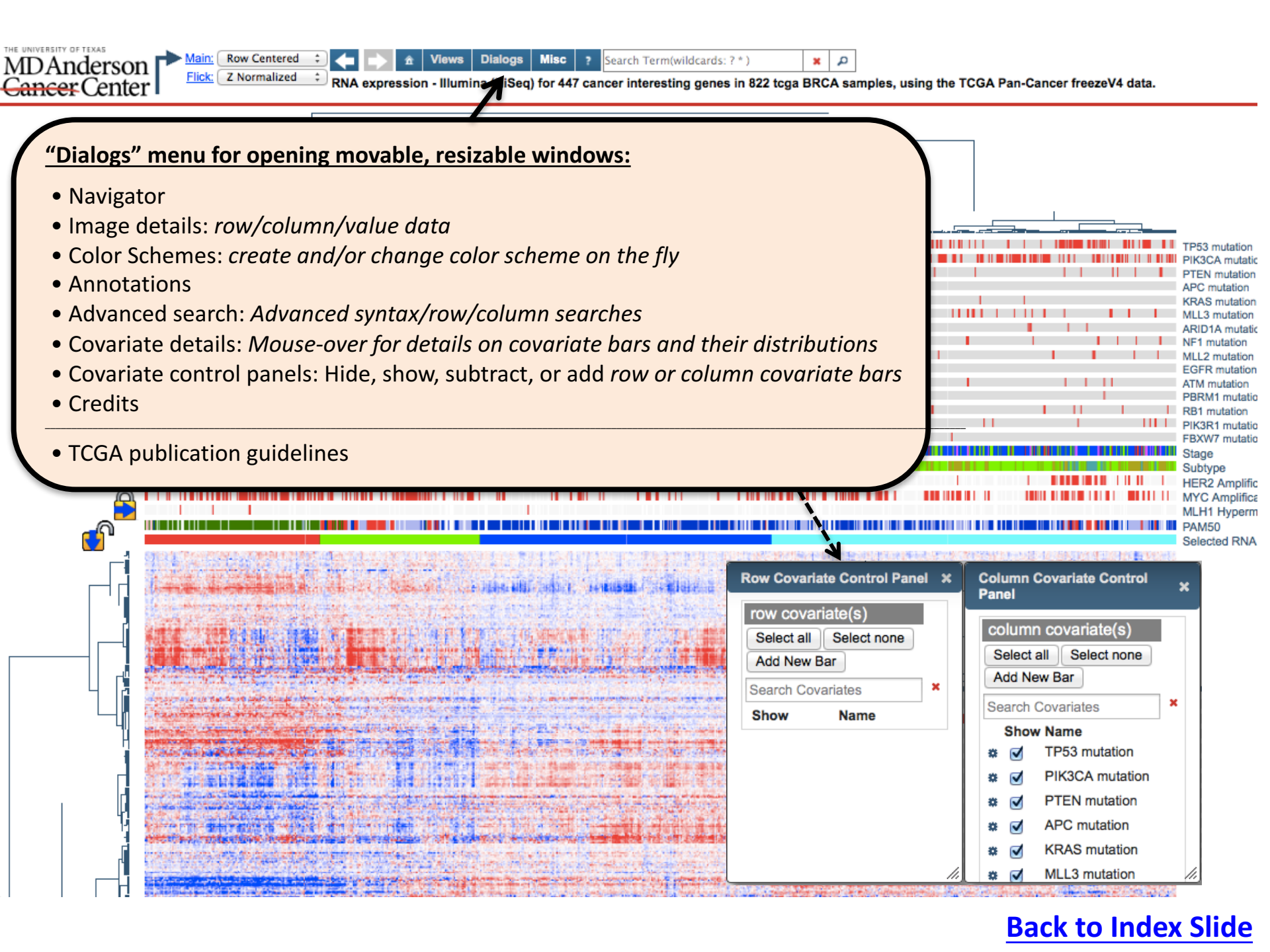

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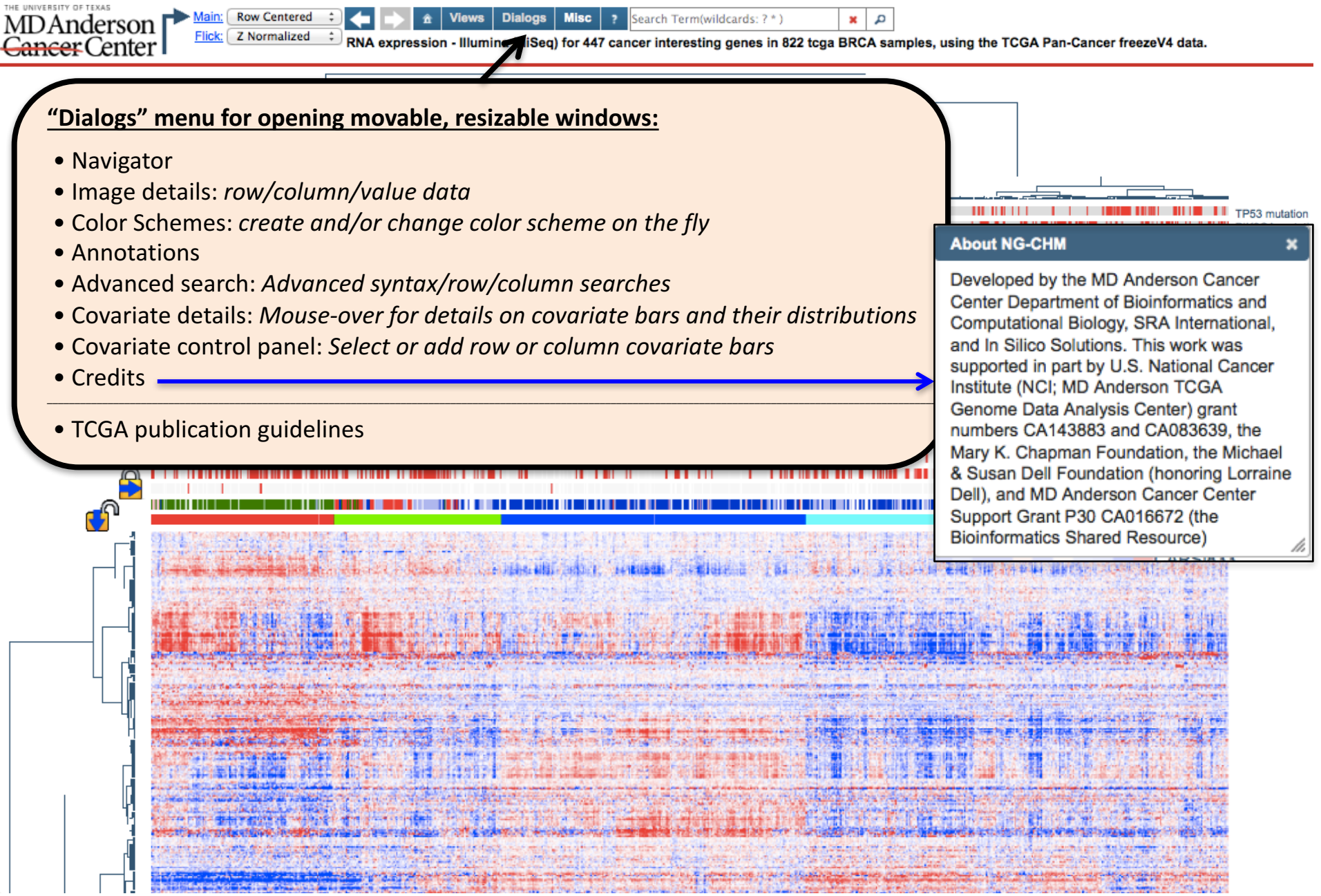

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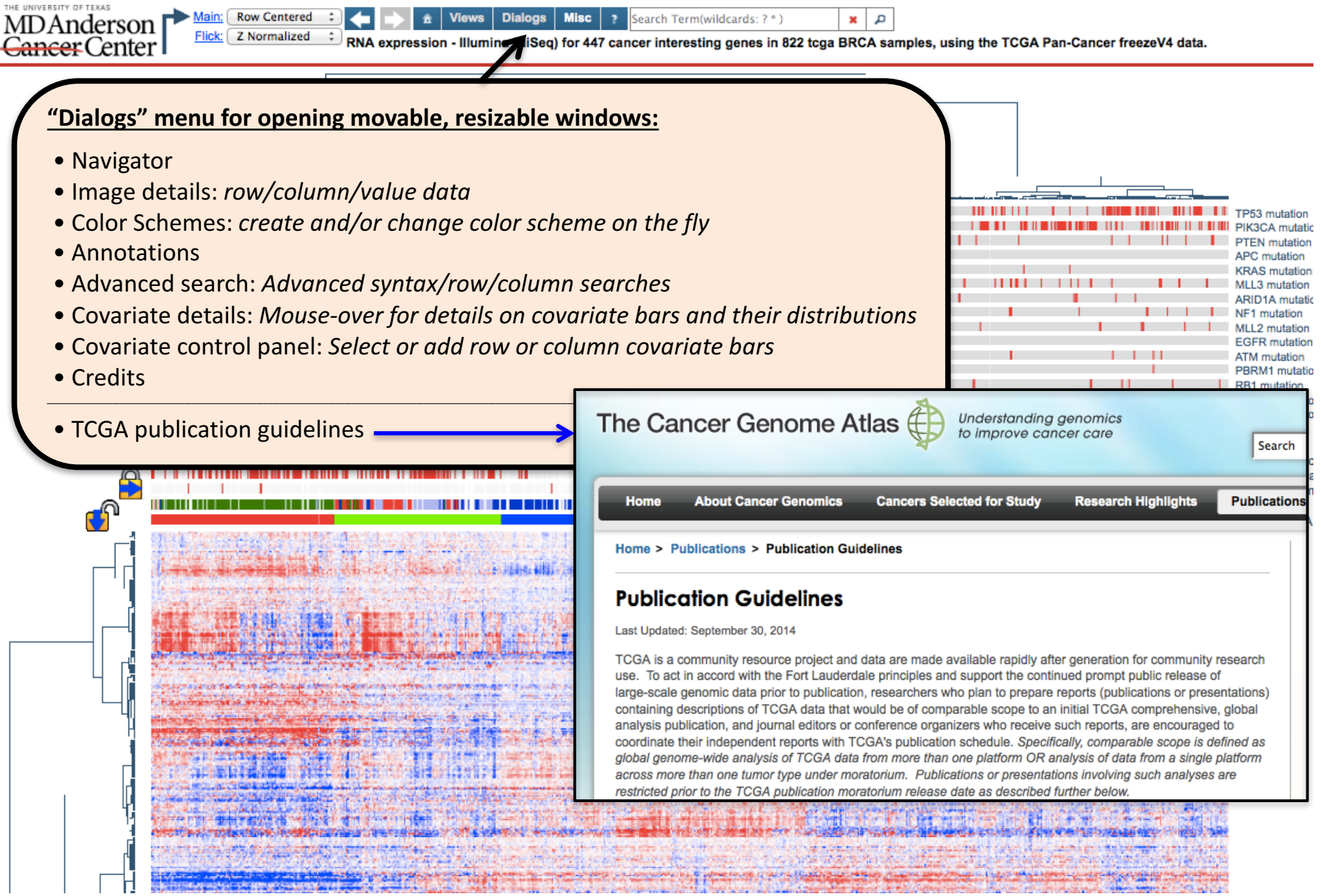

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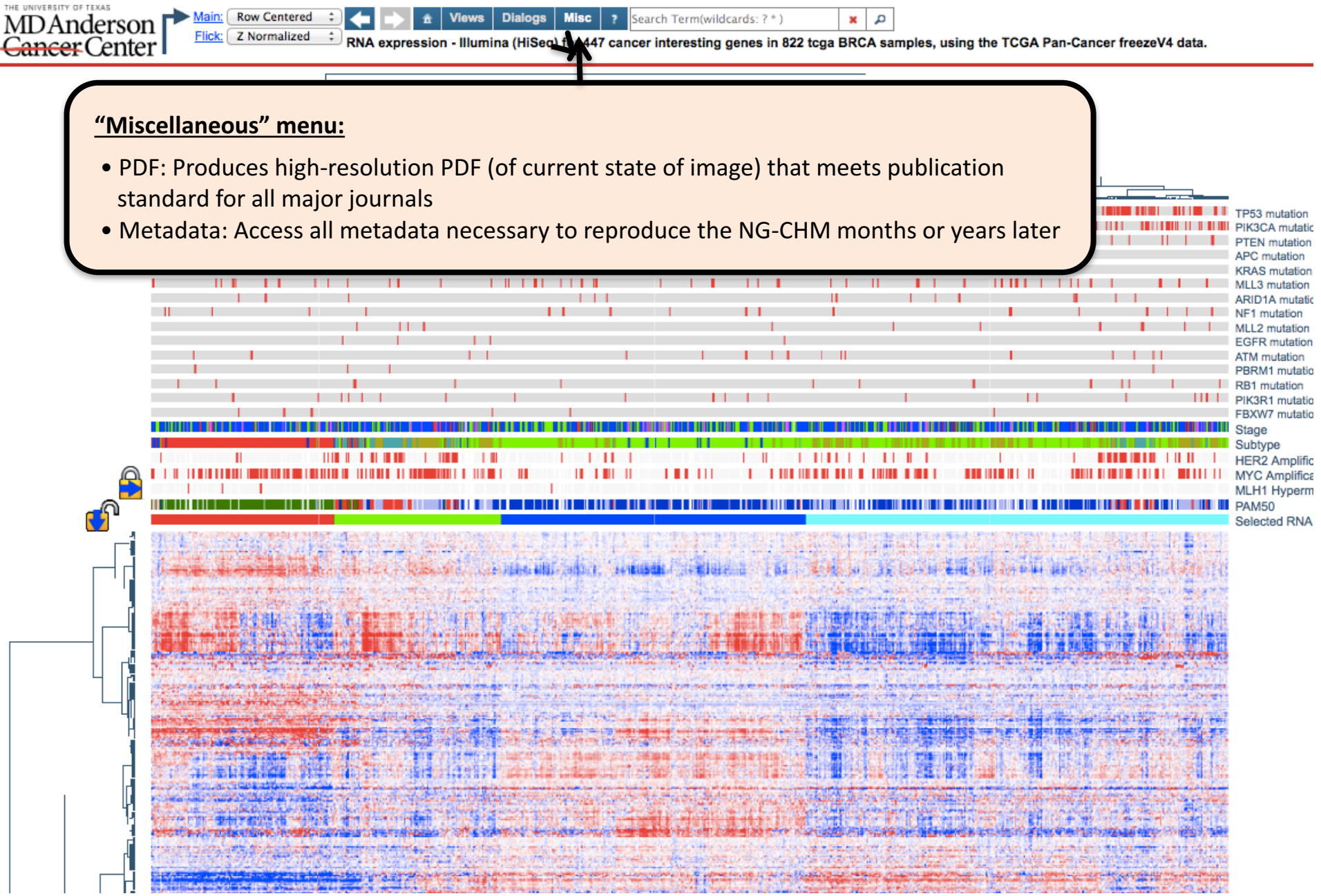

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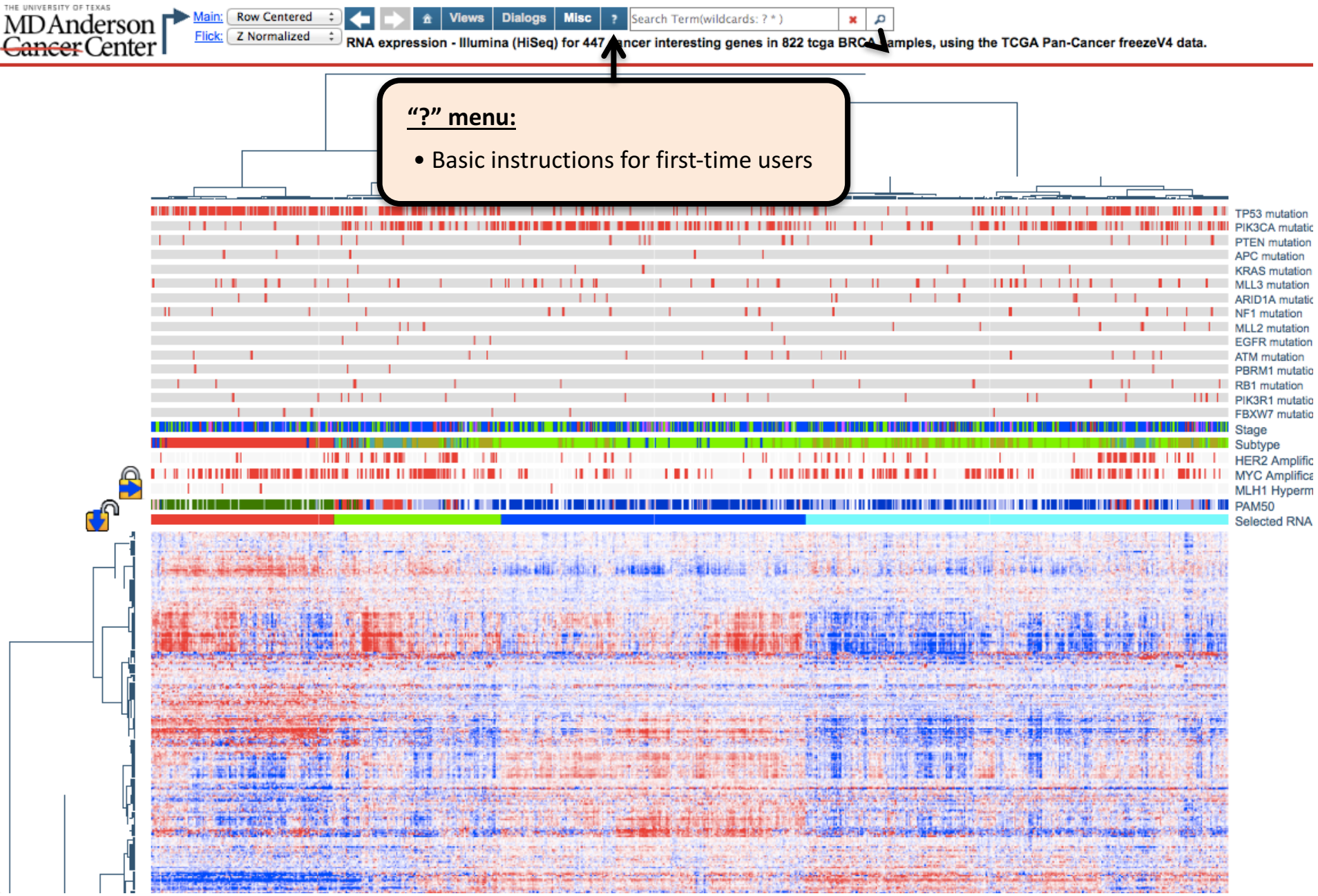

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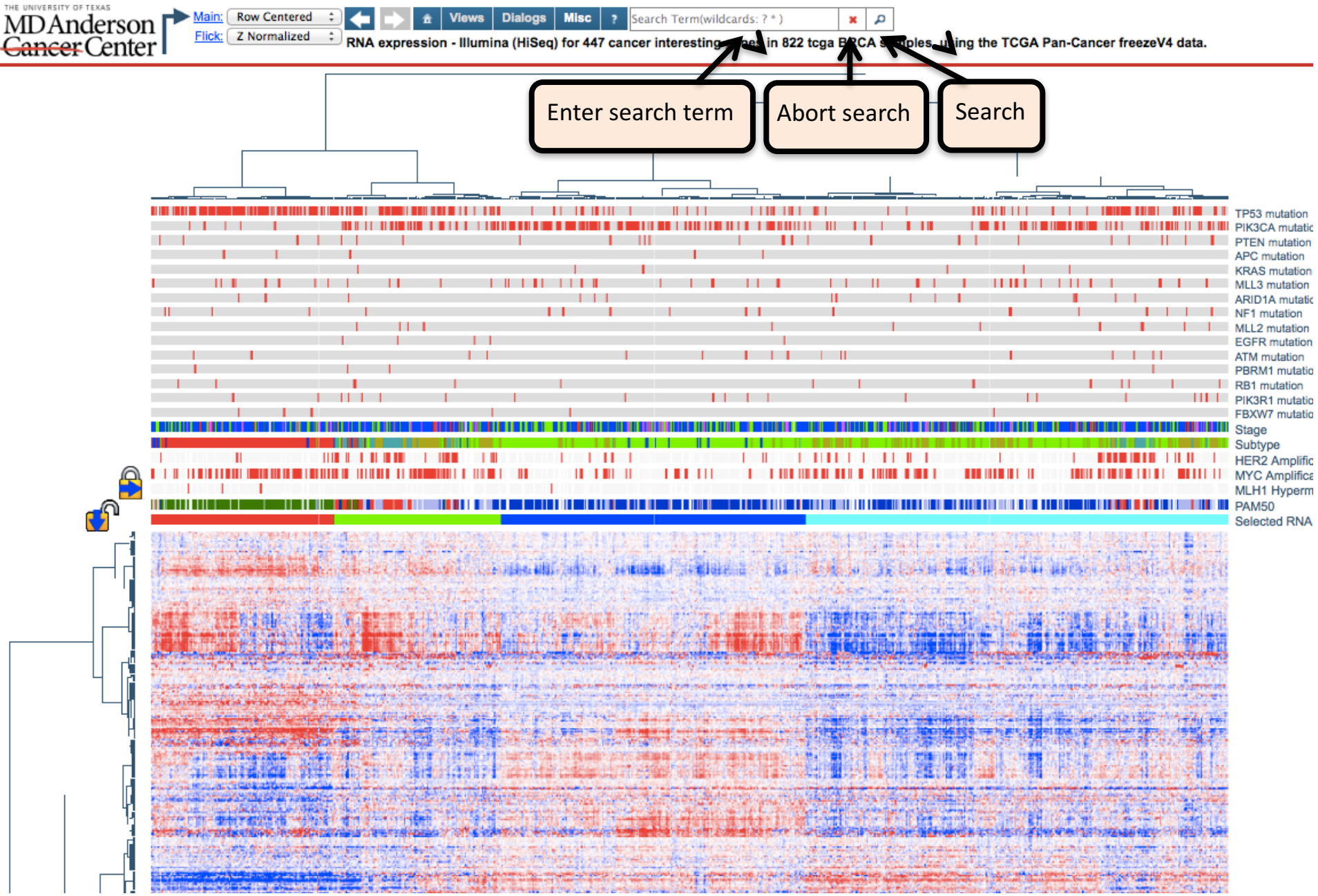

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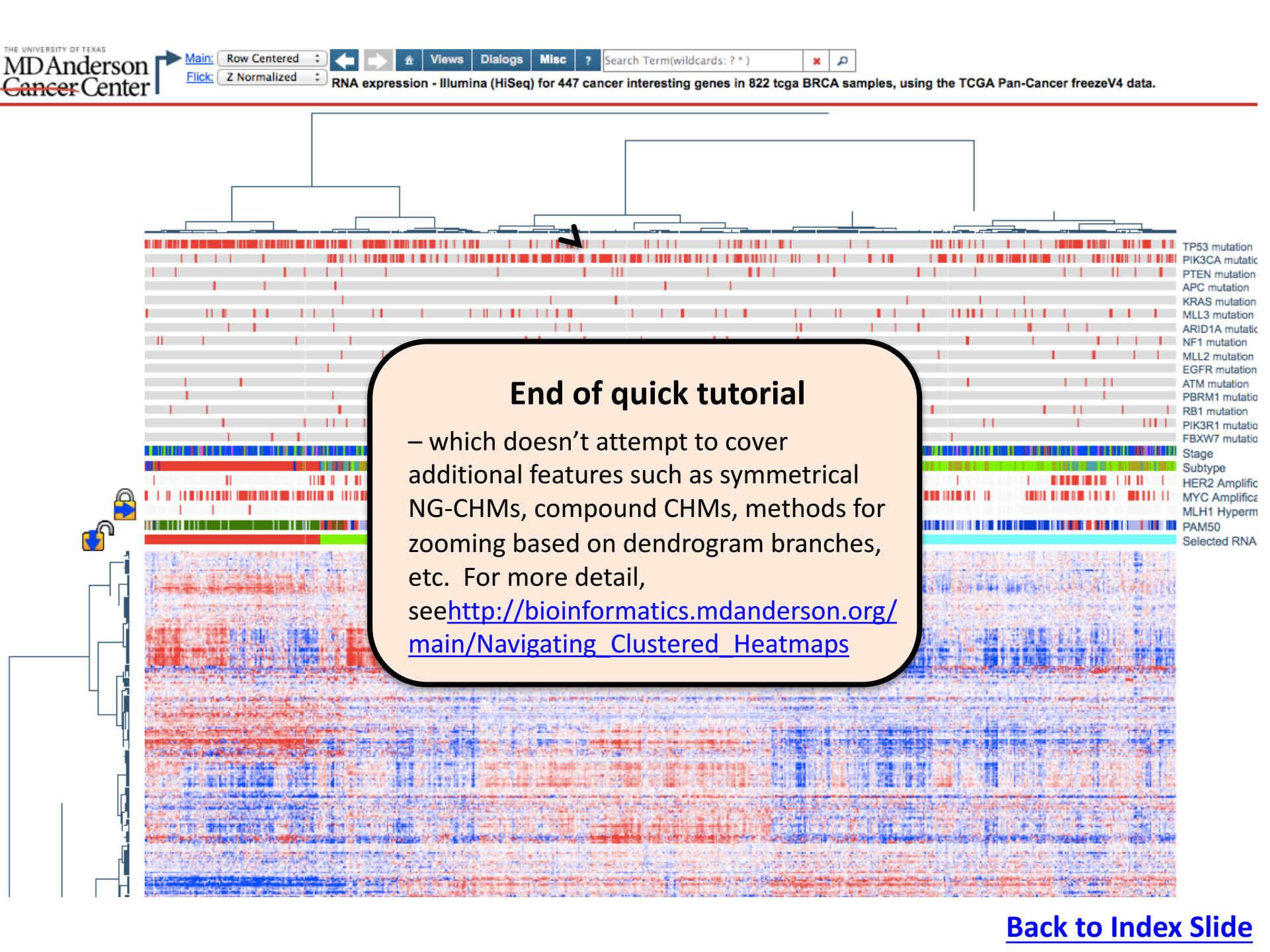

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