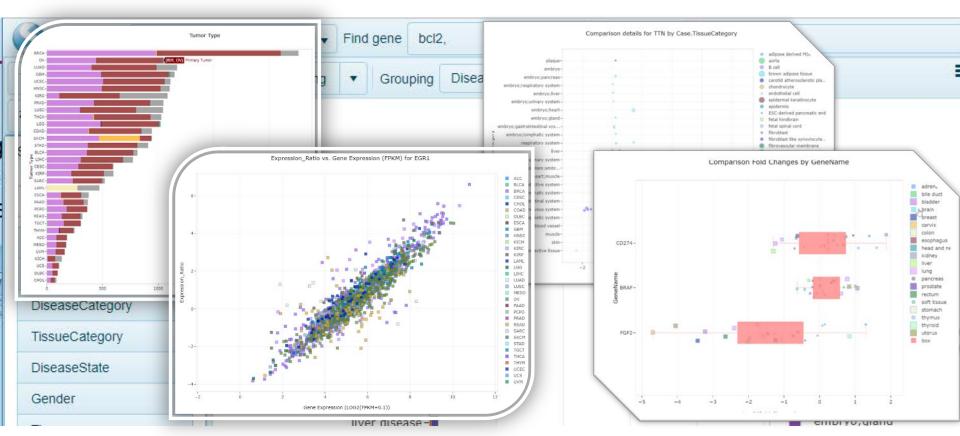


OmicSoft Land Explorer

Making disease-relevant data accessible to biologists



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Bioinformatic Scientist, Discovery Scientific Consulting



Identification of a prognostic RNA-Seq expression signature in endometrial carcinoma

Diagnostic Oncology

Research

Identification of a prognostic RNA-Seq expression signature in endometrial carcinoma

Tim Bonnert*1 and Michelle Engle2

Address: ¹QIAGEN Bioinformatics, Skelton House, Lloyd Street North, Manchester, M₁₅ 6SH, UK and ²QIAGEN OmicSoft, 5001 Weston Parkway, Suite 201 Cary, NC 27513, USA

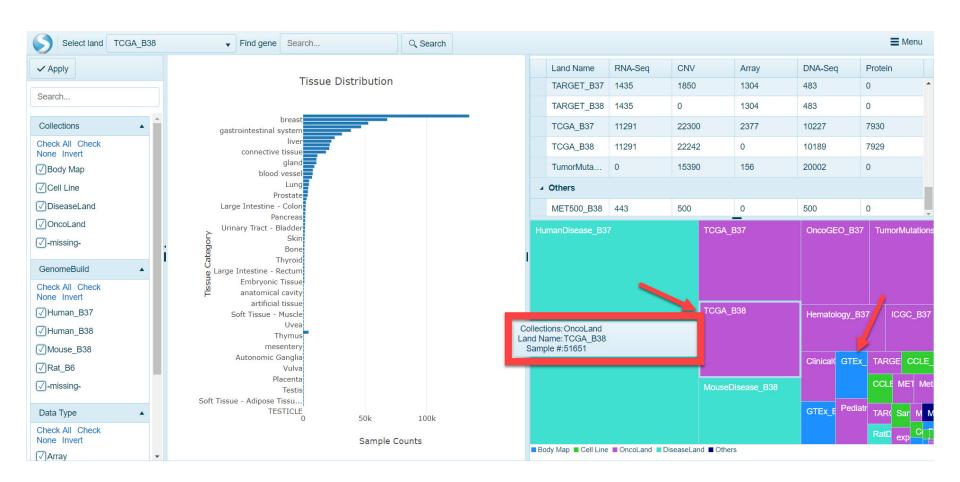
Email: Tim Bonnert* - tim.bonnert@qiagen.com

* Corresponding author

Abstract

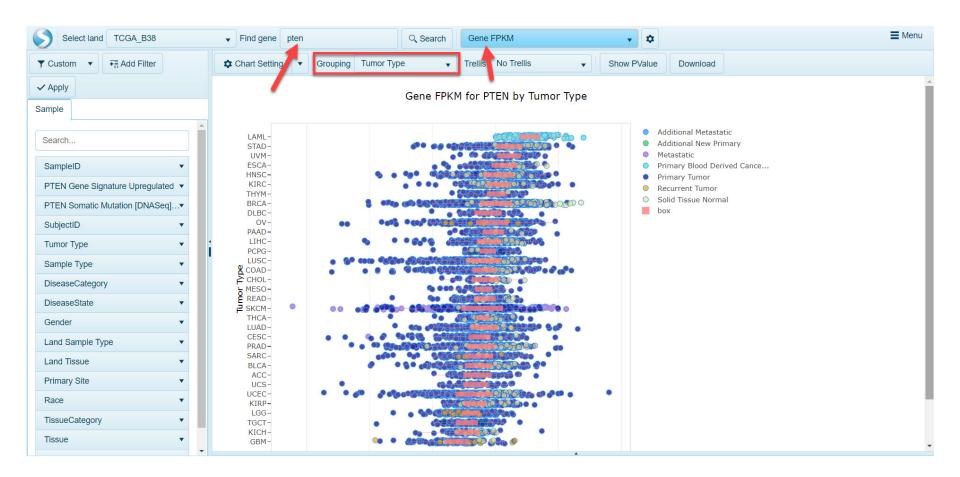


Sample Explorer



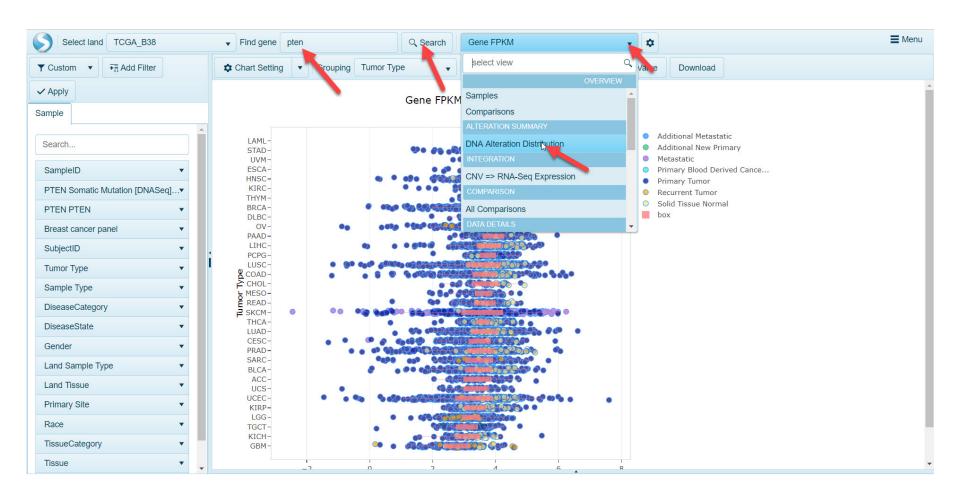


Gene FPKM for PTEN across all tumor types



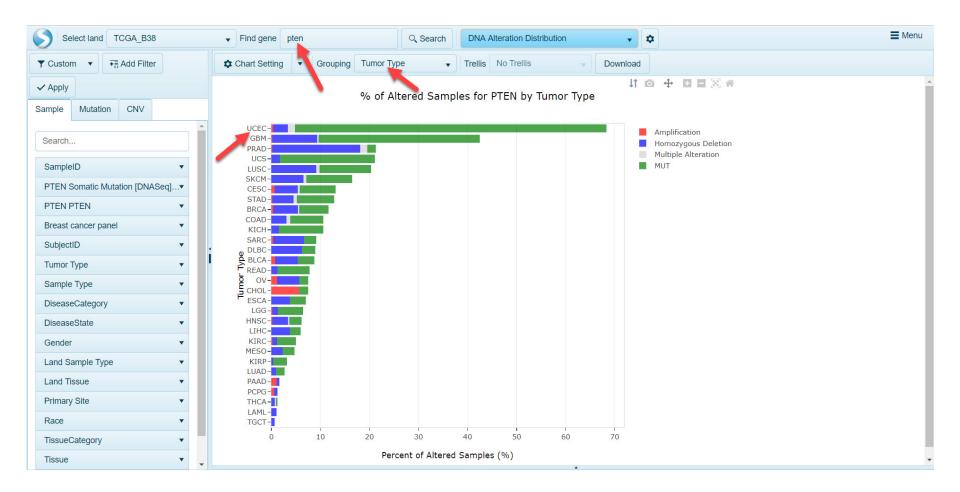






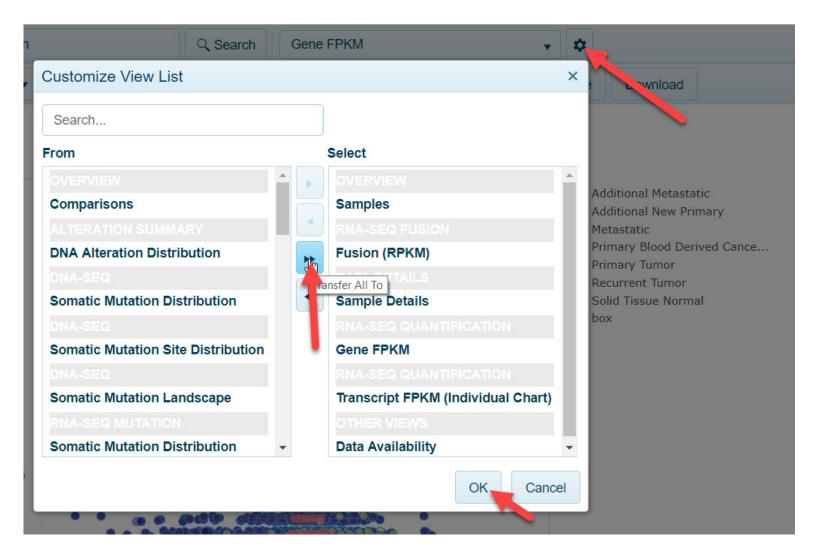


PTEN has the highest mutation rate in UCEC tumors



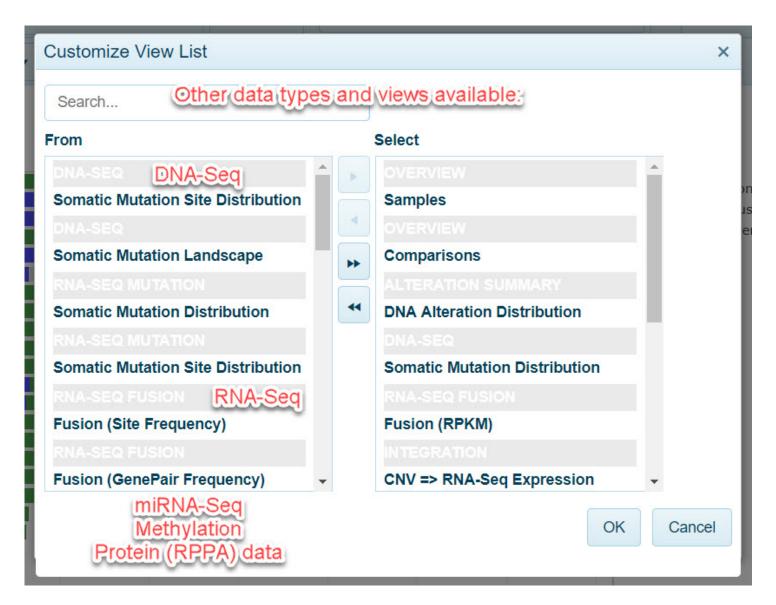






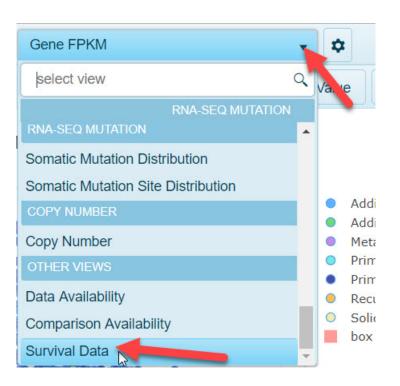






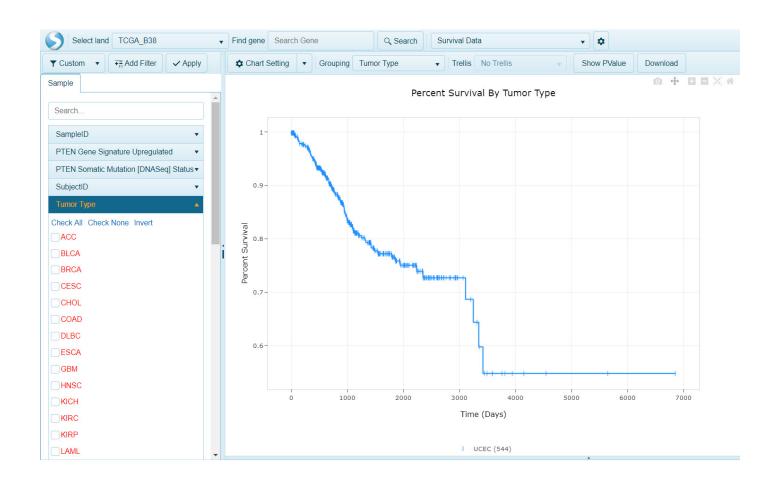






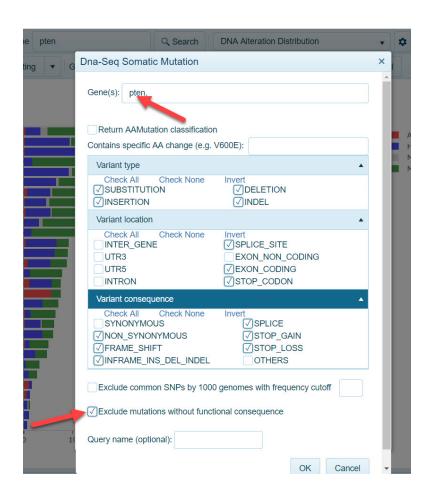


Survival plot of all UCEC patients





Create a custom query based on PTEN mutation status

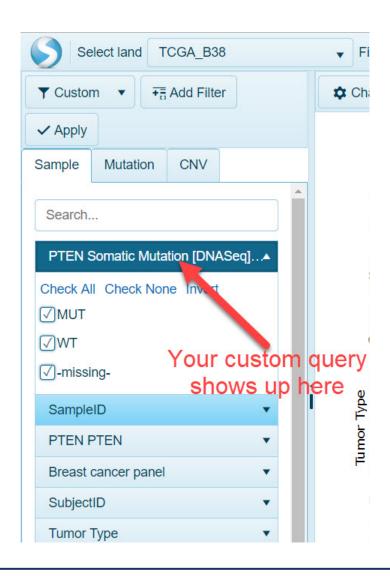


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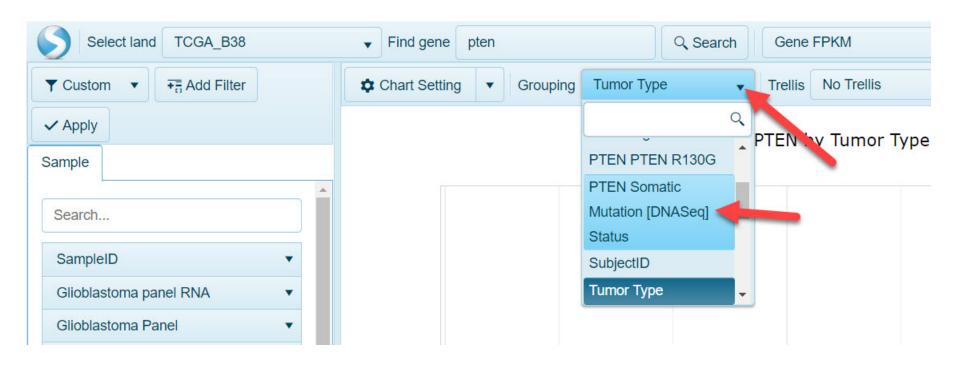






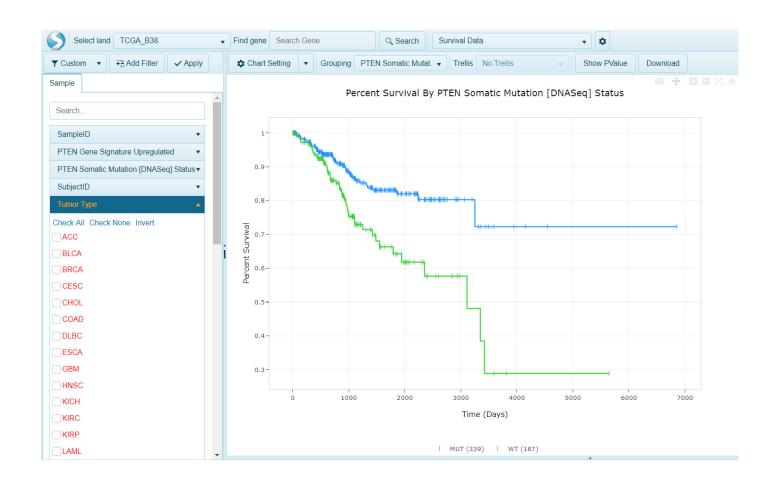






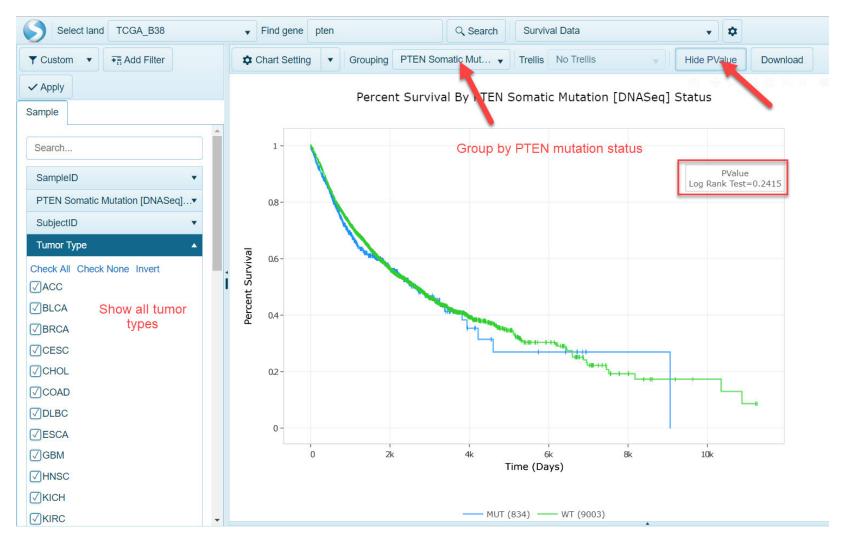


Survival plot for UCEC patients, grouped by PTEN WT or MUT





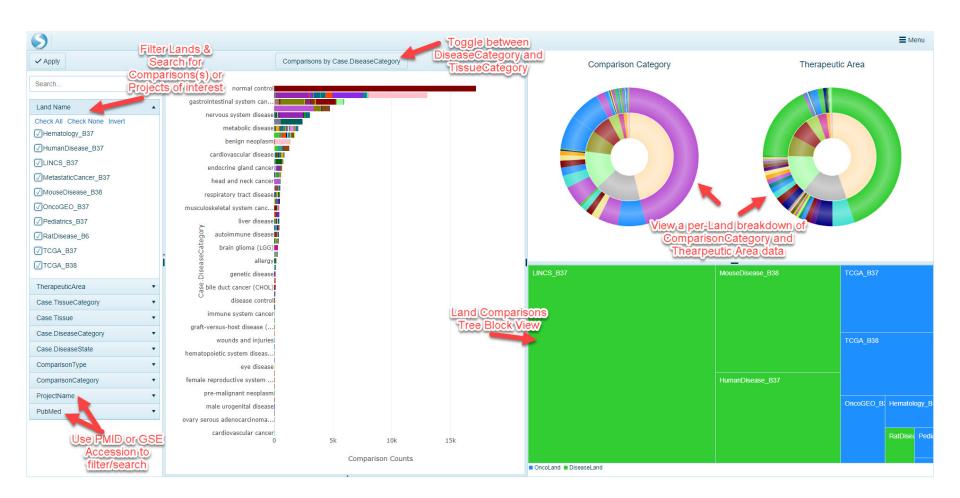




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Comparison Explorer



Sample to Insight

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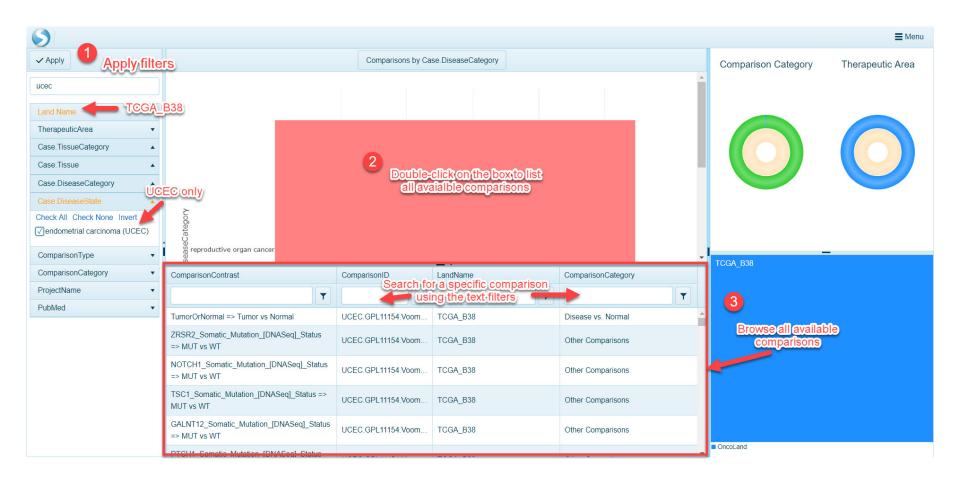
Filter down to TCGA_B38 comparisons



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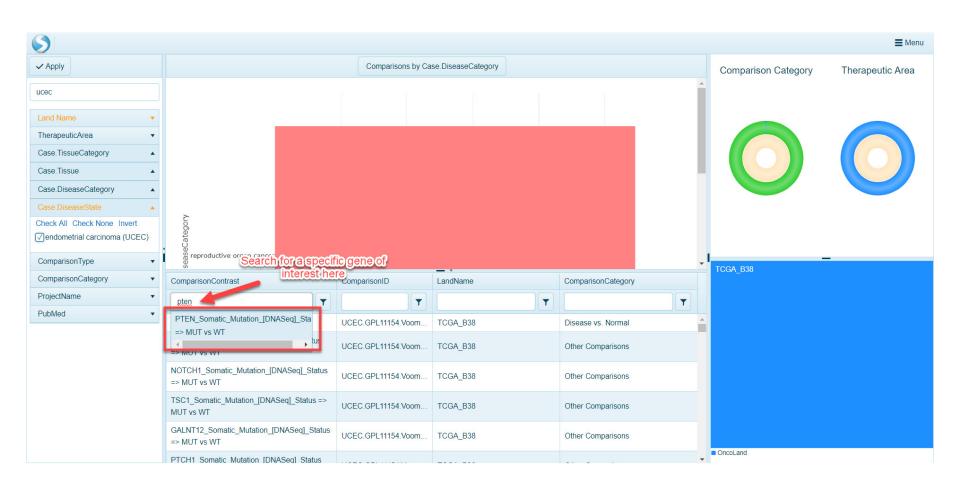
Filter down to UCEC comparisons in TCGA_B38 Land



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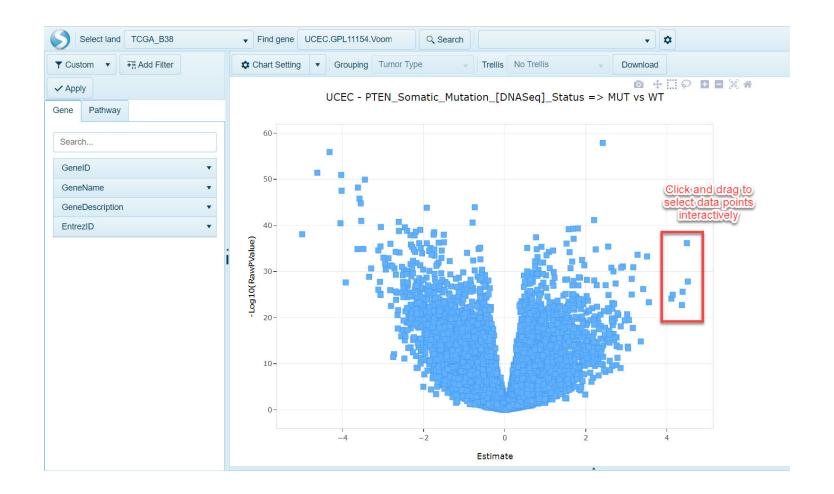






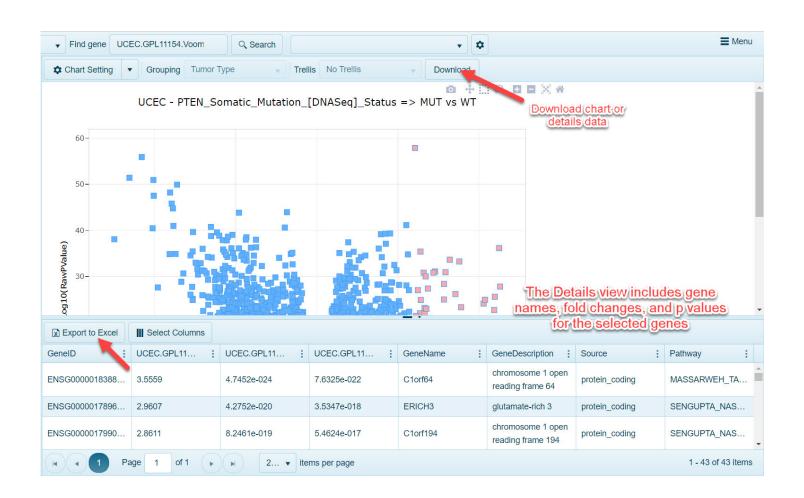


Browse the precalculated data comparing PTEN WT vs MUT expression for all genes



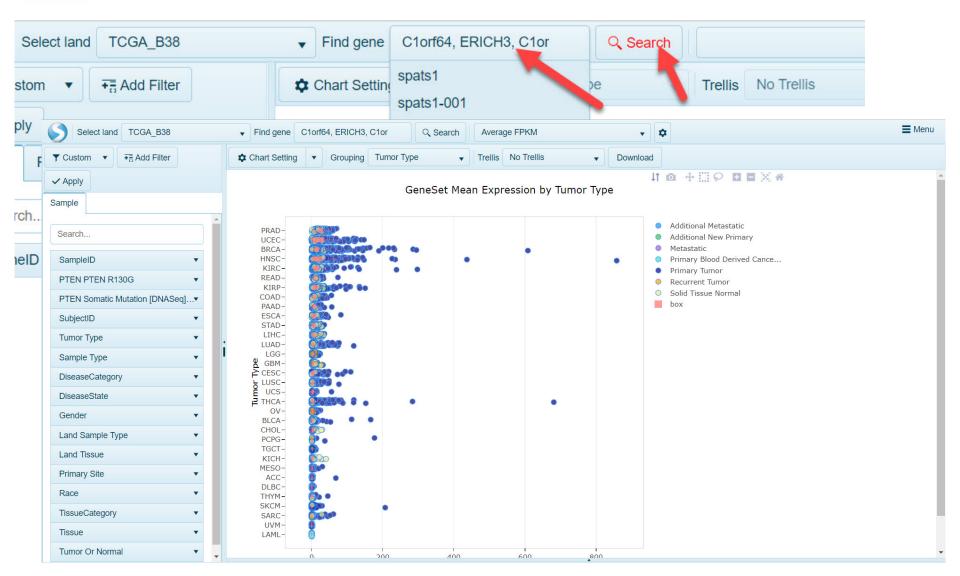


Select genes to see details and create a gene set





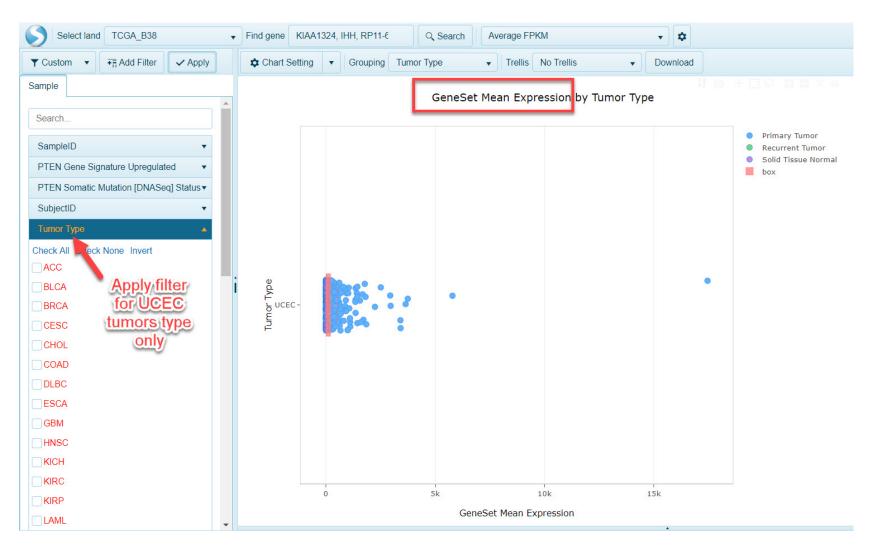
Search for gene set expression using the chosen genes



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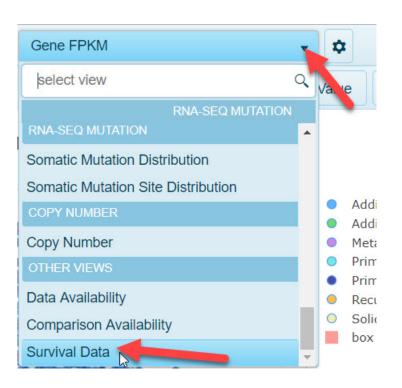
View gene set mean expression in UCEC tumors only



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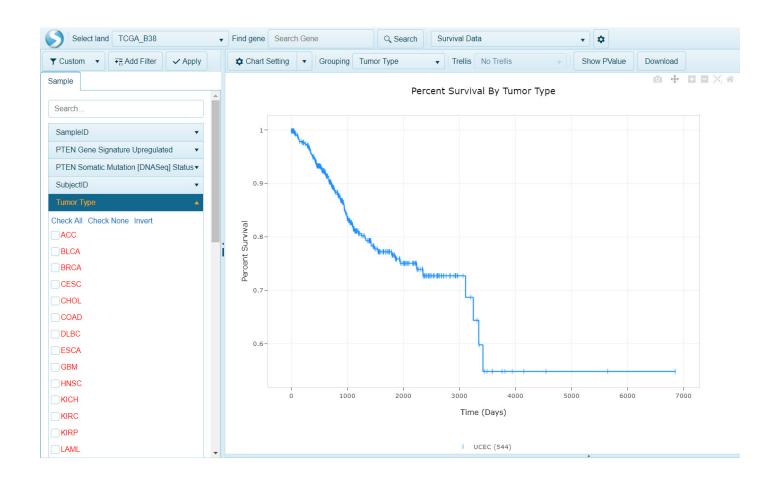






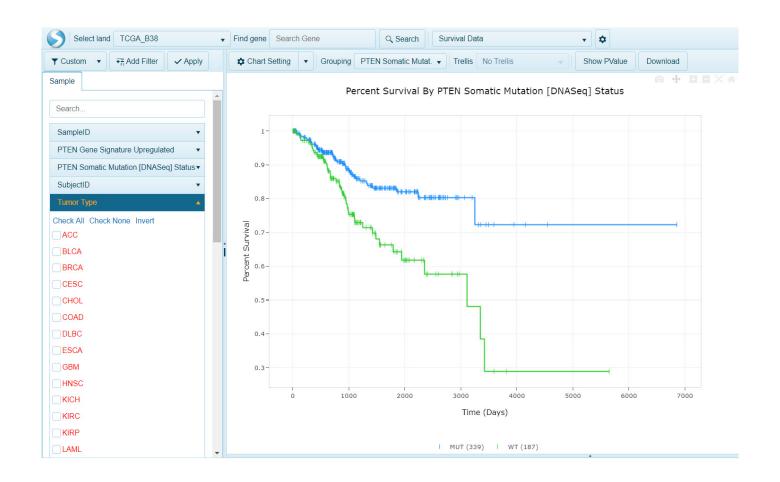


Recreate previous survival plot with UCEC patients only



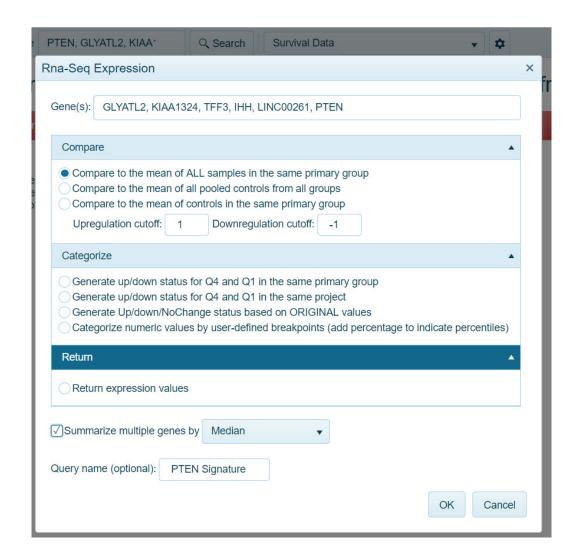


Survival plot for UCEC patients, grouped by PTEN WT or MUT



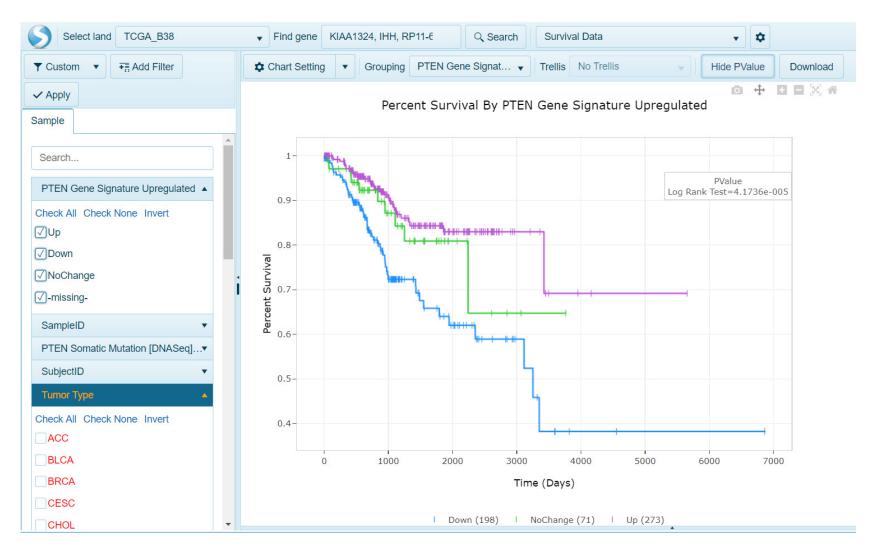


Create custom query for gene set expression of 6 genes





Group survival data by gene set expression



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Land explorer provides an opportunity to simply and rapidly browse, filter, and visualise the millions of data points and annotations in the Lands

- Found all TCGA expression data for PTEN across different cancer types
- Visualised the distribution of PTEN somatic mutations across TCGA samples
- Profiled the landscape of PTEN mutations across endometrial carcinoma samples
- Examined Kaplan-Meier survival plots for endometrial carcinoma separated by PTEN mutation status
- Browsed and filtered the prebuilt comparisons for expression data comparing endometrial carcinoma samples with- or without somatic mutations in PTEN
- Selected genes from the volcano plot visualisation and created an RNA-Seq gene set
- Used the gene set expression characteristics as a surrogate for PTEN mutation status as a prognostic indicator in endometrial carcinoma



Thank you for your attention!



Questions?

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omicsoft.support@qiagen.com



OmicSoft Land Explorer: Free trial throughout March!

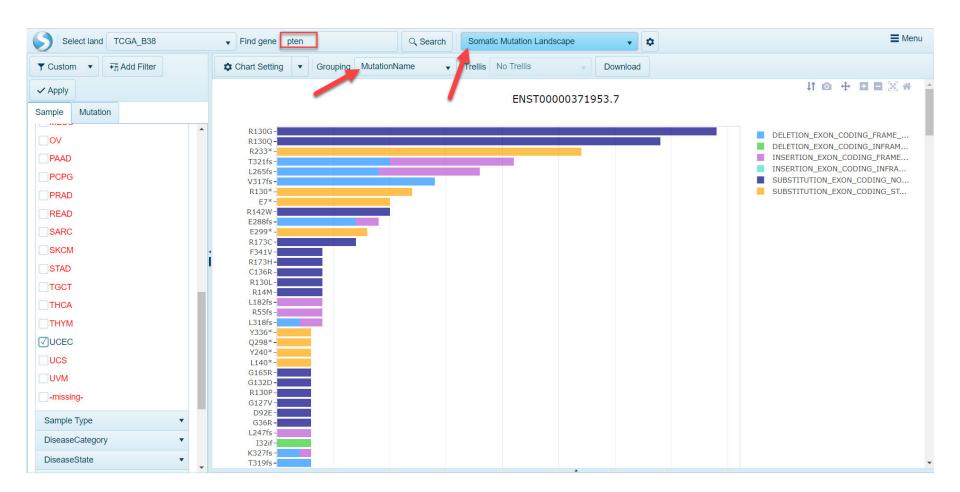


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Switch to Somatic Mutation Landscape view to view specific amino acid changes in PTEN mutations



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