


Merged Cohort of LGG and GBM (TCGA, Cell 2016) (/study?id=lgggbm_tcga_pub)


Groups: (drag to reorder) (A) Age<=35 (200) (B) Age>60 (398) Select all | Deselect all

Overlap(overlap?sessionId=5ed9b11fe4b030a3bfd0cc1a)

Survival(survival?sessionId=5ed9b11fe4b030a3bfd0cc1a)

Clinical(clinical?sessionId=5ed9b11fe4b030a3bfd0cc1a)





Plot Type

100% stacked bar chart

☐ Swap Axes

☐ Horizontal Bars

Clinical Attribute	Attribute Type	Statistical Test	p-Value	q-Value ▲
ATRX status	Sample	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
RPPA Cluster	Sample	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
Overall Survival Status	Patient	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
Supervised DNA Methylation Cluster	Sample	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
Pan-Glioma DNA Methylation Cluster	Sample	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
Study	Patient	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
IDH status	Sample	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
IDH/codel subtype	Sample	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
IDH-specific DNA Methylation Cluster	Sample	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
Pan-Glioma RNA Expression Cluster	Sample	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
U133a	Sample	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
Transcriptome Subtype	Sample	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
Original Subtype	Sample	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
IDH-specific RNA Expression Cluster	Sample	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
Chr 7 gain/Chr 10 loss	Sample	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
Neoplasm Histologic Grade	Sample	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
Neoplasm Histologic Type Name	Patient	Chi-squared Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
Mutation Count	Sample	Kruskal Wallis Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
Diagnosis Age	Patient	Kruskal Wallis Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰
Karnofsky Performance Score	Patient	Kruskal Wallis Test	< 10 ⁻¹⁰	< 10 ⁻¹⁰

Showing 1-20 of 50

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Show more

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