

Hazard ratio

mean_highos_genes_z (N=180)

0.48
(0.31 - 0.74)



0.001 **

mean_lowos_genes_z (N=180)

1.51
(0.97 - 2.33)



0.066

Events: 86; Global p-value (Log-Rank): 0.0028769

AIC: 802.66; Concordance Index: 0.6