

## Hazard ratio

**mean\_highos\_genes\_z** (N=153)

0.24  
(0.15 – 0.39)



<0.001 \*\*\*

**mean\_lowos\_genes\_z** (N=153)

2.93  
(1.98 – 4.34)



<0.001 \*\*\*

# Events: 92; Global p-value (Log-Rank): 5.4449e-11

AIC: 787.72; Concordance Index: 0.71