

AMH signature

TCGA-BRCA

TCGA-LUAD

TCGA-COAD

Meta

TCGA-BRCA

TCGA-LUAD

TCGA-COAD

Meta

Active

Loss

Weight (%)

- 30
- 60
- 90
- 120

Signature

- Active
- Loss

- DFI
- DSS

0.2 0.5 1 2 4 10

HR

estimates with a standard error > 10 were removed from the plot