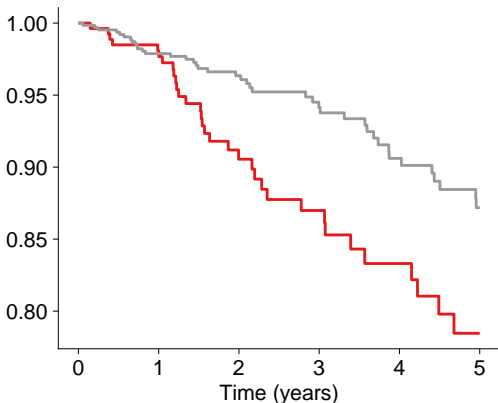


DFI TCGA-BRCA AMH fully active signature

Survival Probability



— High — Low

Meta-analysis HR(95%CI)=0.49(0.34-0.7);P=9e-05

High

At Risk	280	238	141	107	75	55
Events	0	5	20	25	29	33

Low

At Risk	670	551	354	255	189	139
Events	0	13	20	27	35	41