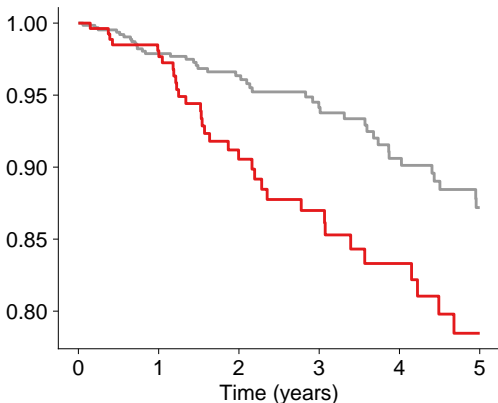


# DFI TCGA-BRCA AMH fully active signature

Survival Probability



## Low

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

## High

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