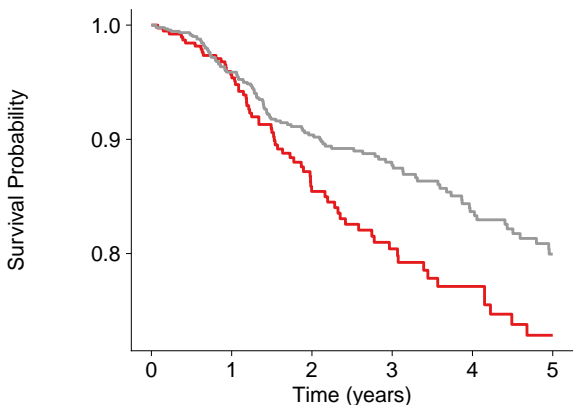


# DFI TCGA-BRCA-LUAD-COAD AMH fully active



## High

At Risk	400	335	196	140	97	71
Events	0	16	45	55	60	65

## Low

At Risk	952	786	476	336	239	173
Events	0	36	74	86	99	108