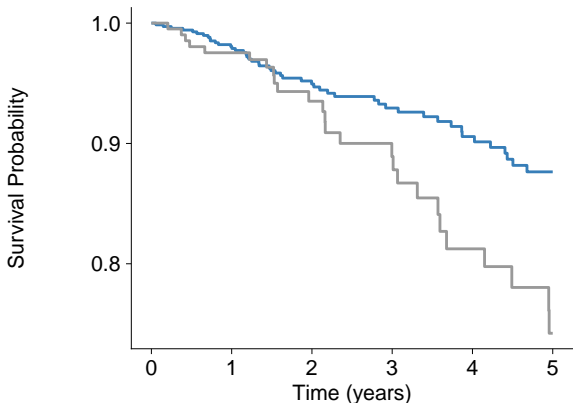


# DFI TCGA-BRCA AMH loss signature (TME)



## High

At Risk	726	605	382	281	209	155
Events	0	13	29	36	42	48

## Low

At Risk	224	184	113	81	55	39
Events	0	5	11	16	22	26