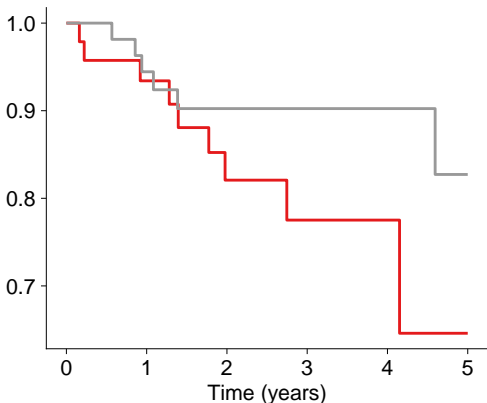


DFI TCGA-COAD AMH fully active signature

Survival Probability



High

At Risk	47	40	26	11	7	4
Events	0	3	7	8	8	9

Low

At Risk	55	51	31	20	13	9
Events	0	3	5	5	5	6