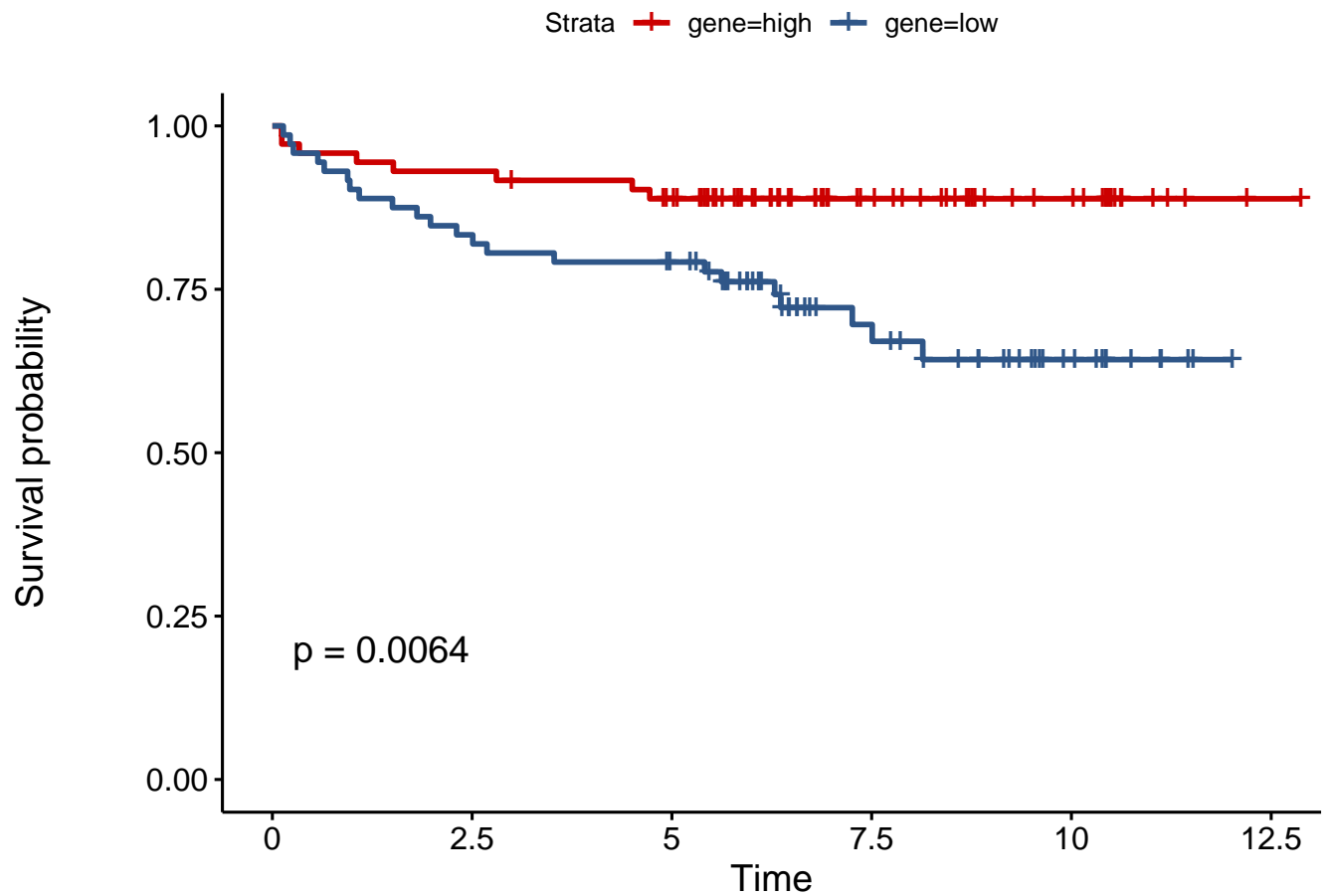


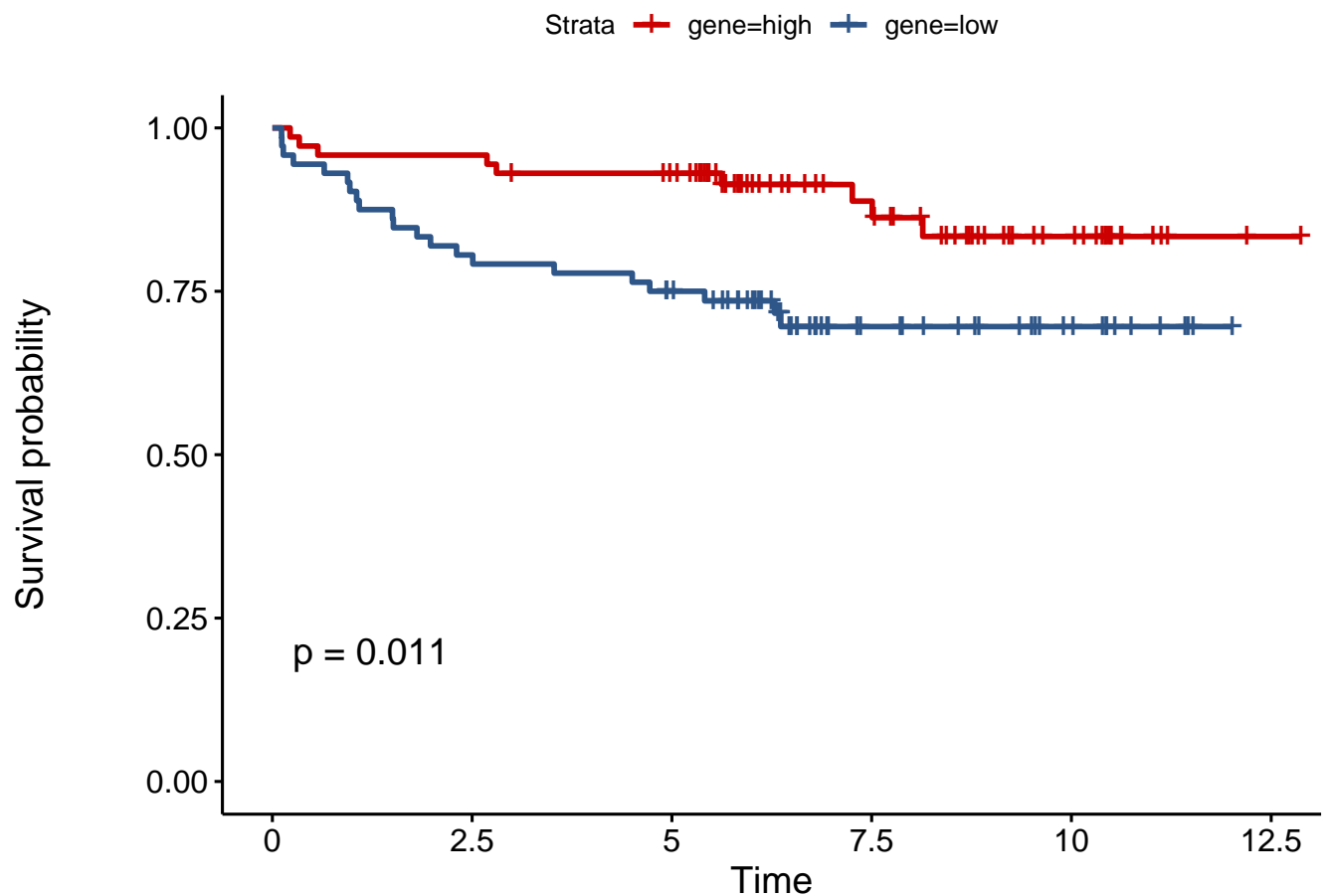
circ_30029



circ_30029

Strata	gene=high	72	67	61	30	16	1
	gene=low	72	60	55	27	11	0
		0	2.5	5	7.5	10	12.5
		Time					

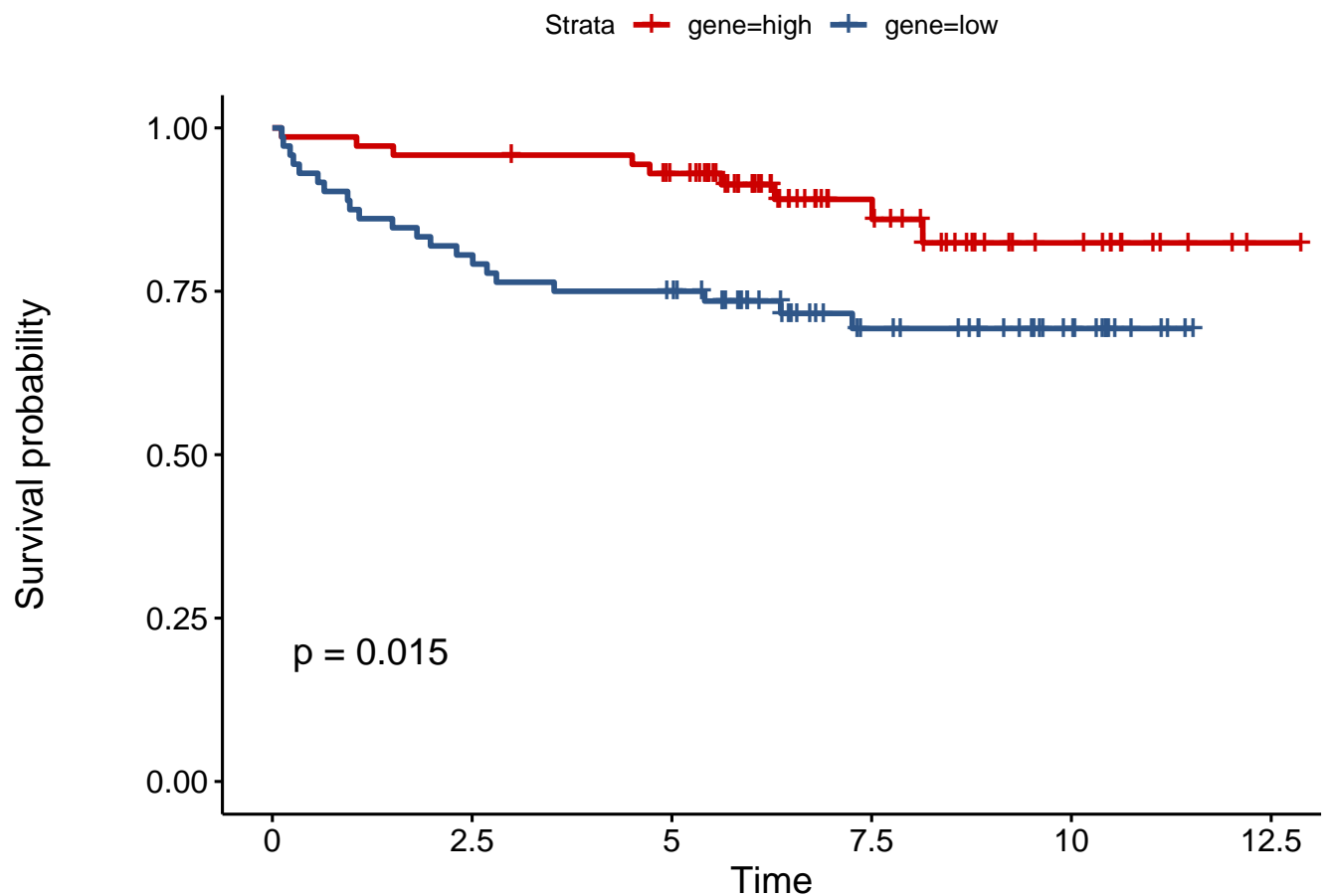
circ_117300



circ_117300

Strata	gene=high	72	69	64	35	16	1
	gene=low	72	58	52	22	11	0
		0	2.5	5	7.5	10	12.5
		Time					

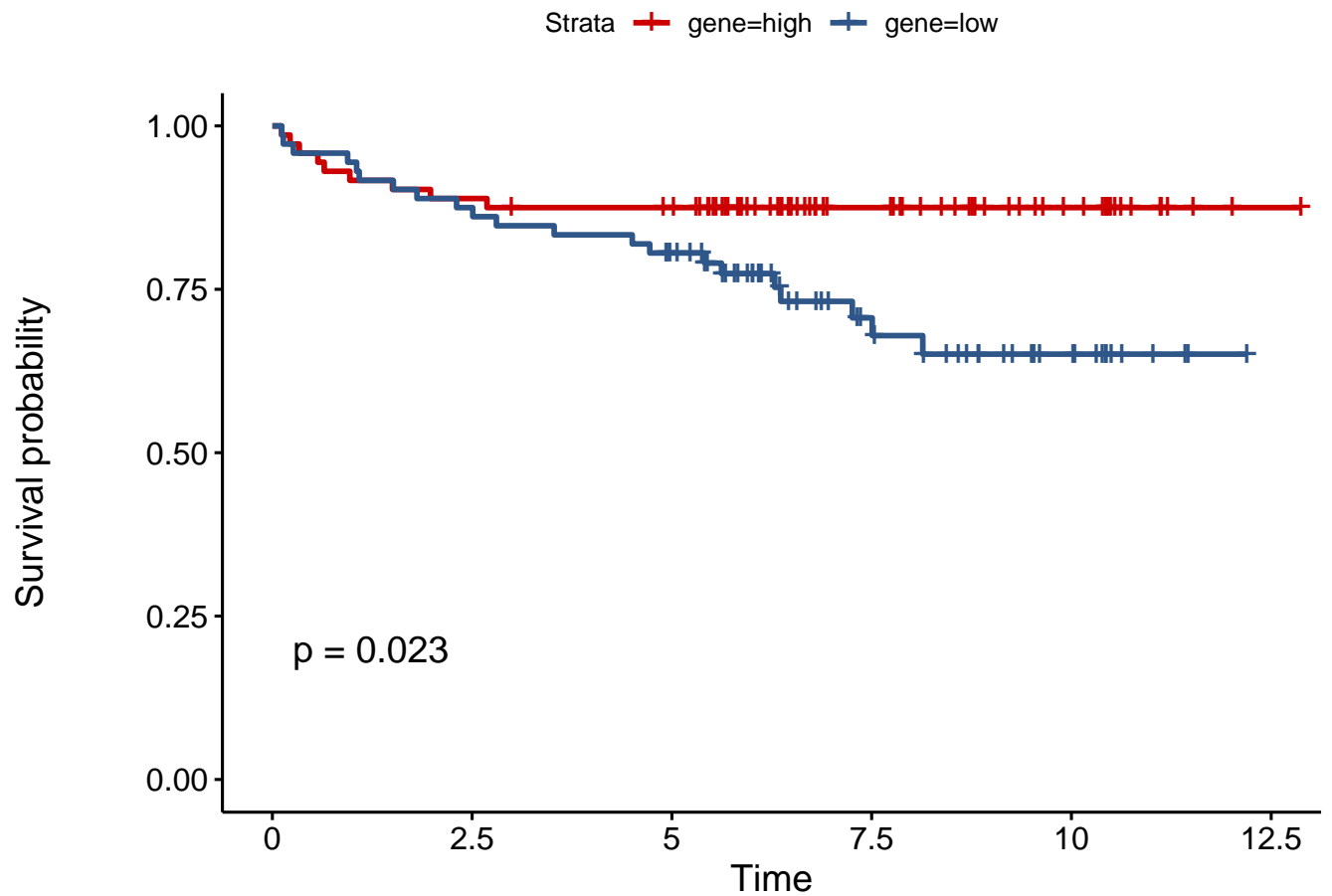
circ_176436



circ_176436

Strata	gene=high	72	69	63	29	12	1
	gene=low	72	58	53	28	15	0
		0	2.5	5	7.5	10	12.5
		Time					

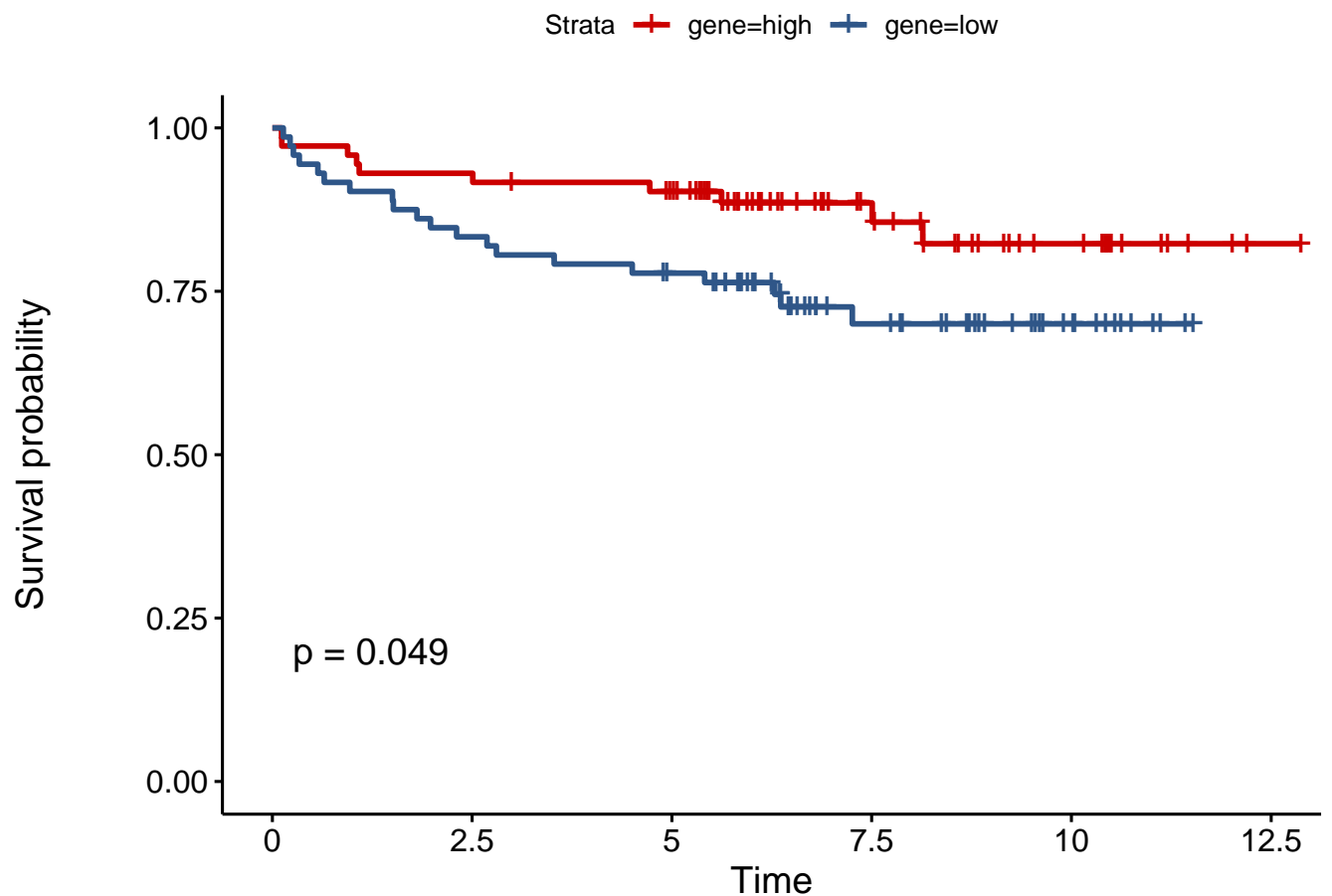
circ_112897



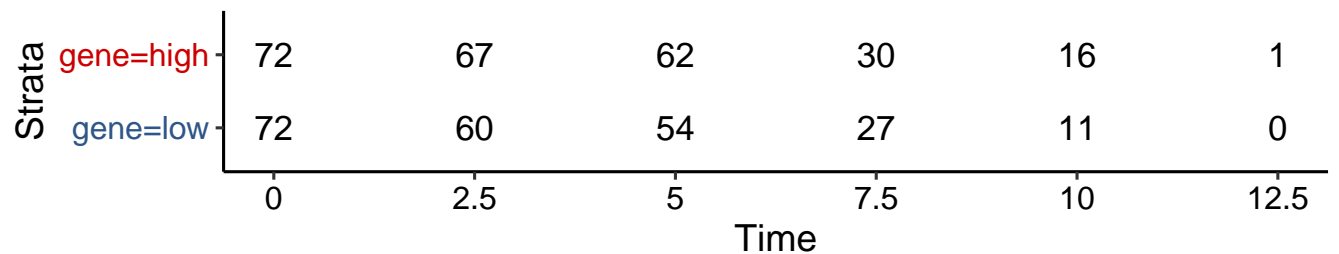
circ_112897

Strata						
gene=high	72	64	61	31	15	1
gene=low	72	63	55	26	12	0
	0	2.5	5	7.5	10	12.5
	Time					

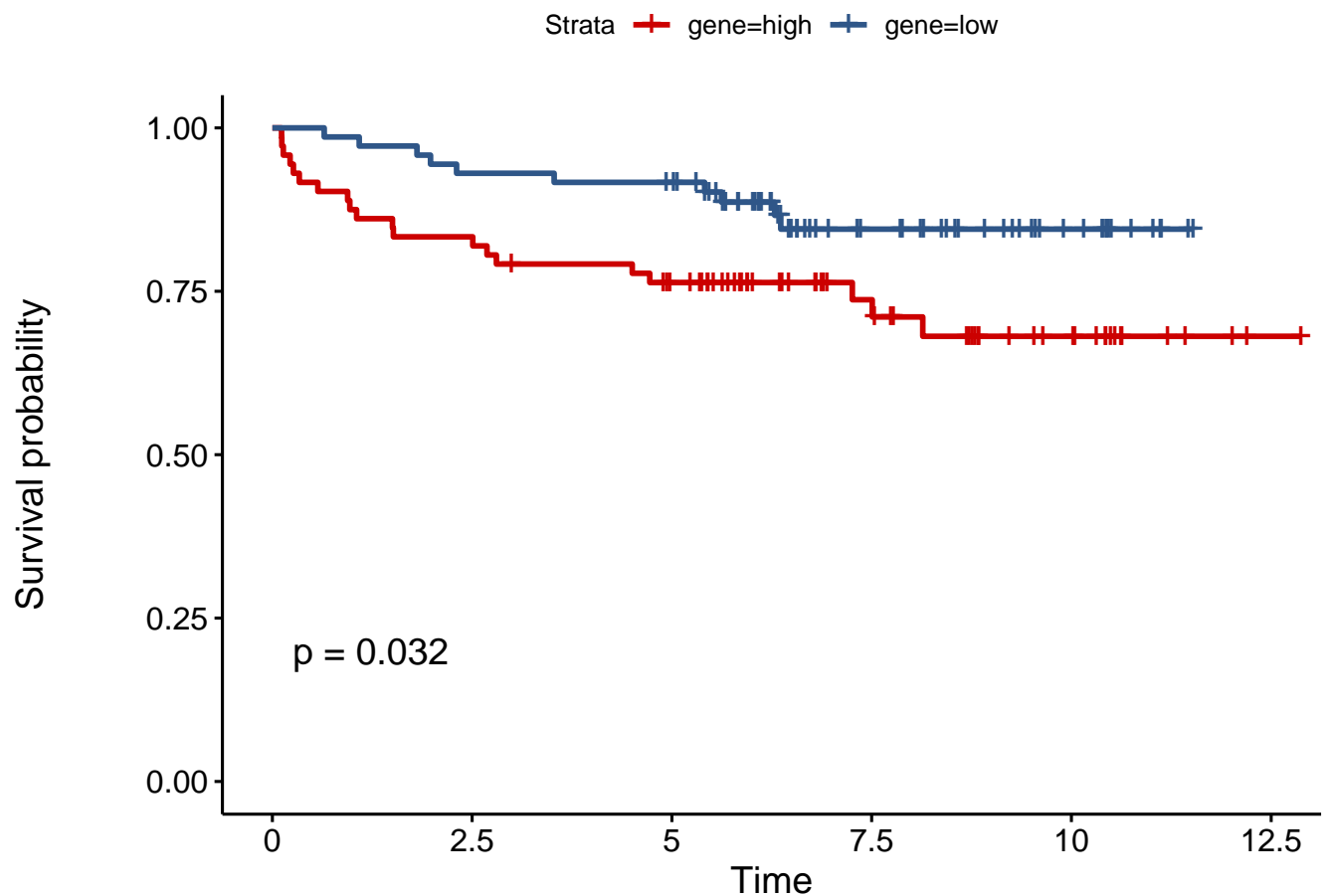
circ_17720



circ_17720



circ_178252



circ_178252

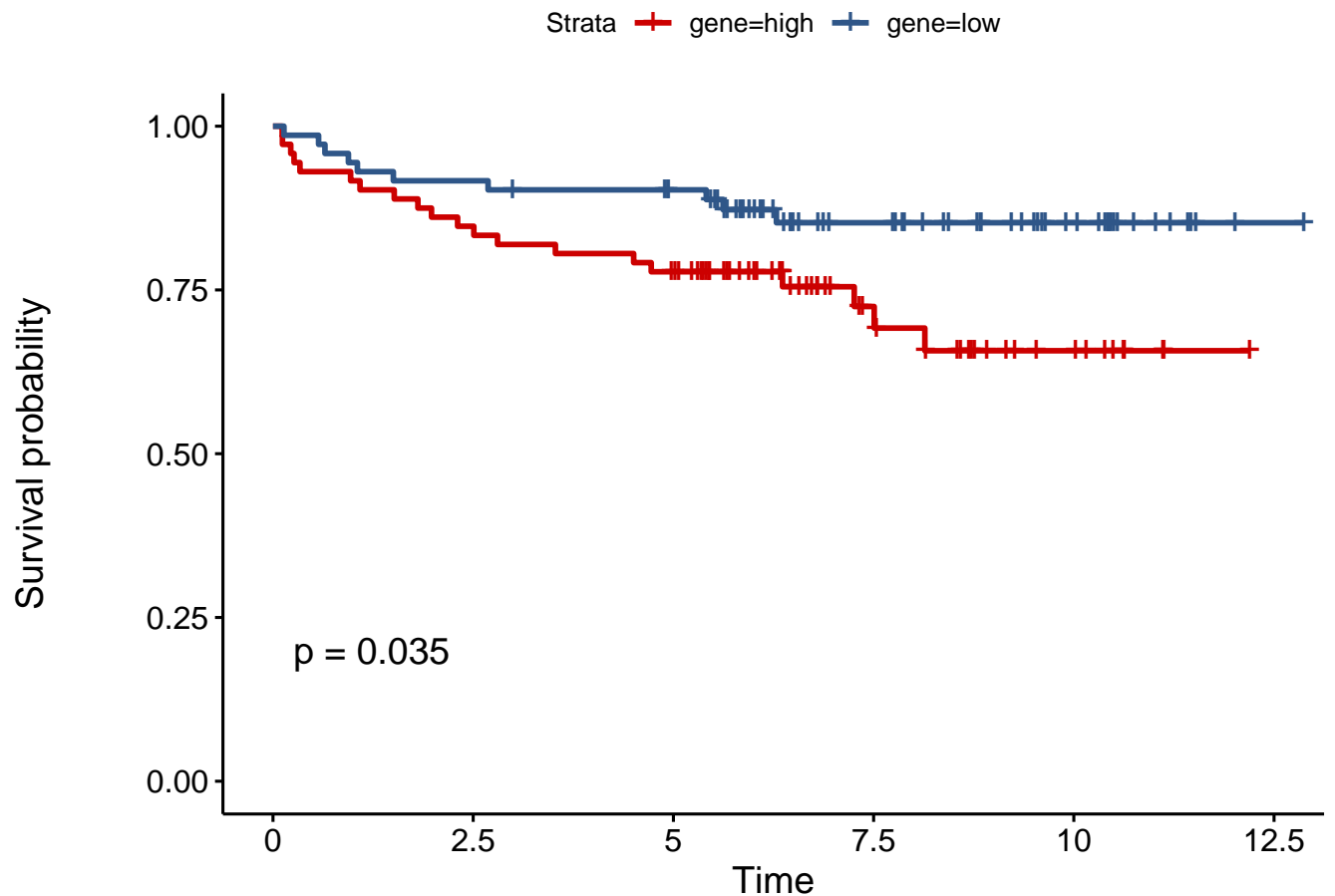
Strata

gene=high	72	60	51	28	14	1
gene=low	72	67	65	29	13	0

Time

Detailed description: This table shows the number of subjects at risk for each strata at different time points. The columns correspond to time points 0, 2.5, 5, 7.5, 10, and 12.5. The rows correspond to the 'gene=high' and 'gene=low' strata.

circ_115617



circ_115617

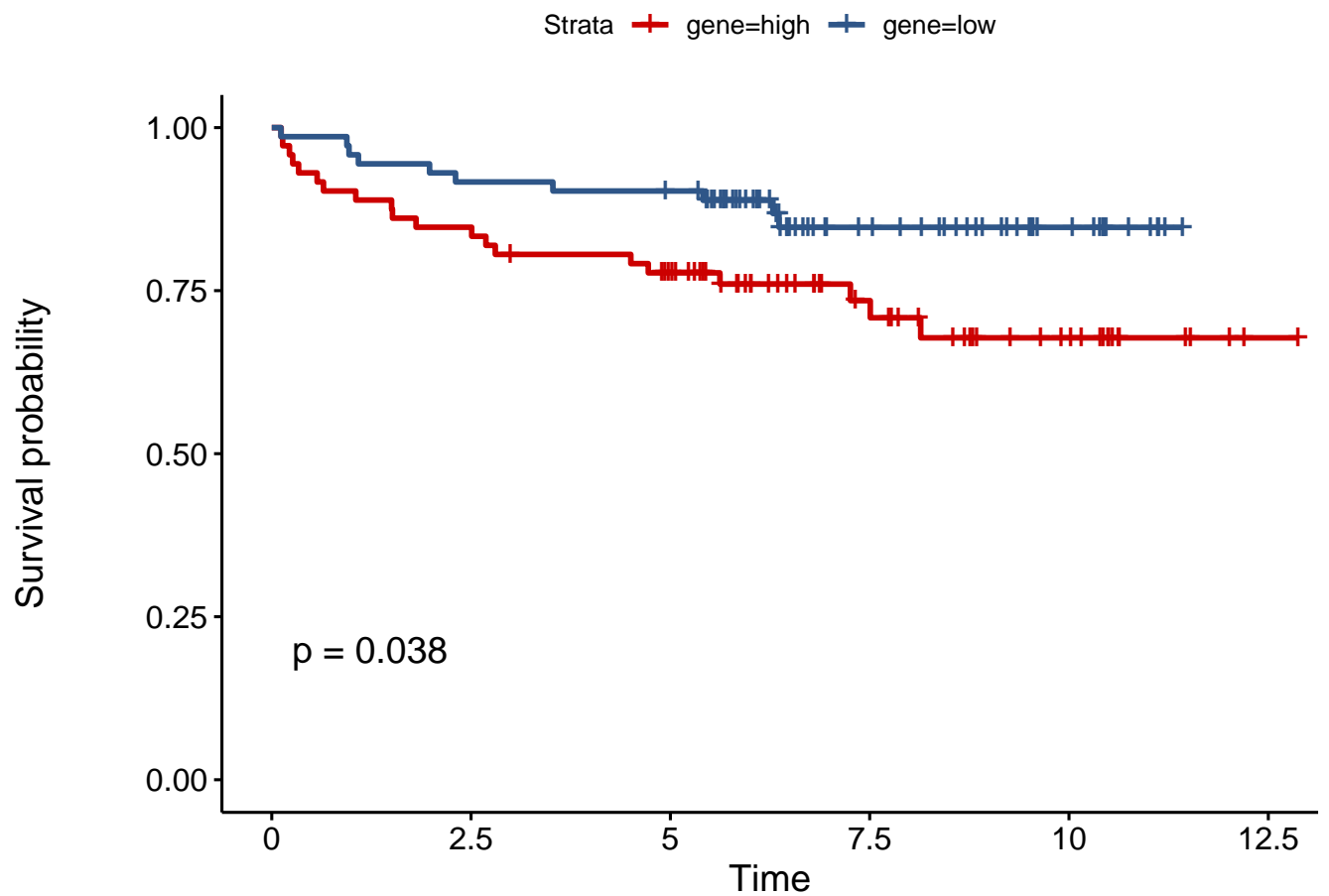
Strata gene=high gene=low

gene=high	72	61	55	22	9	0
gene=low	72	66	61	35	18	1
	0	2.5	5	7.5	10	12.5

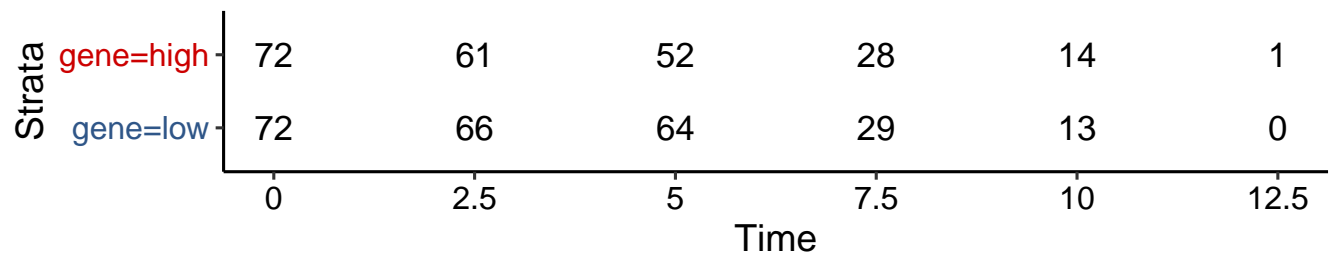
Time

Detailed description: This table shows the number of subjects at risk for each stratum at various time points. The columns represent time points: 0, 2.5, 5, 7.5, 10, and 12.5. The rows represent the strata: gene=high and gene=low.

circ_14736



circ_14736



gene=high	72	61	52	28	14	1
gene=low	72	66	64	29	13	0
	0	2.5	5	7.5	10	12.5