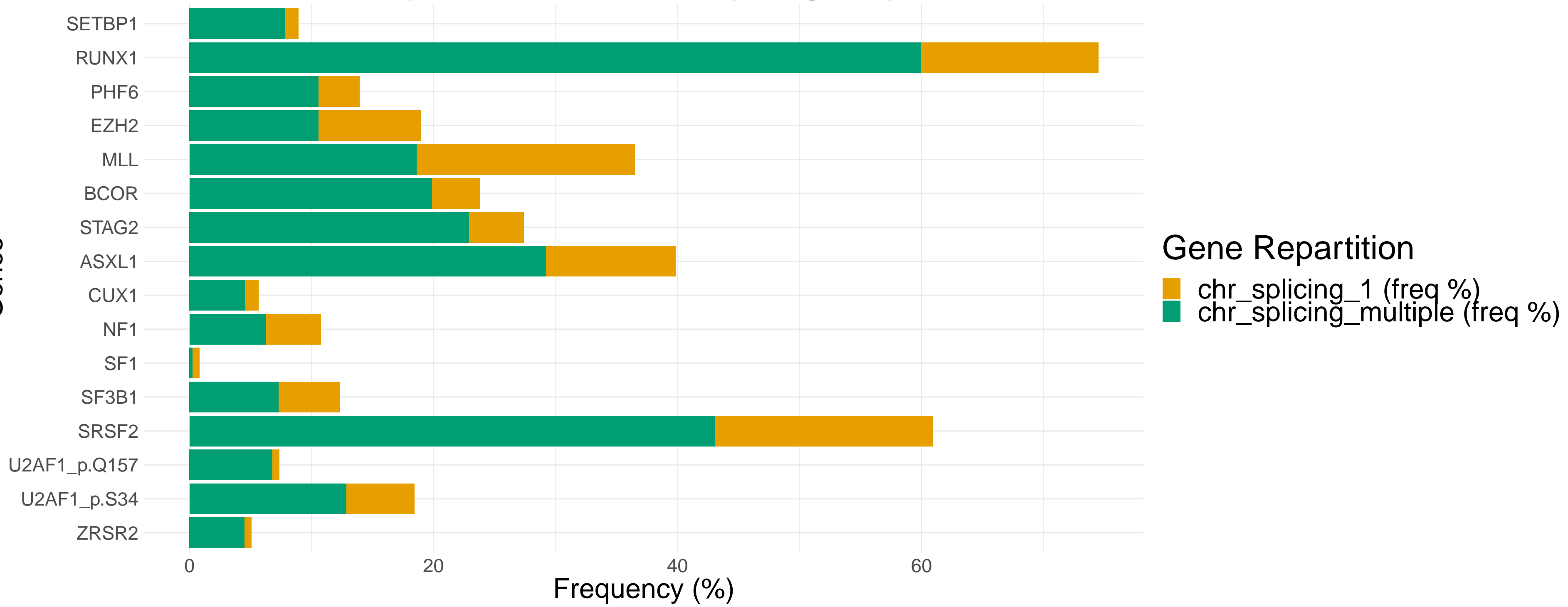
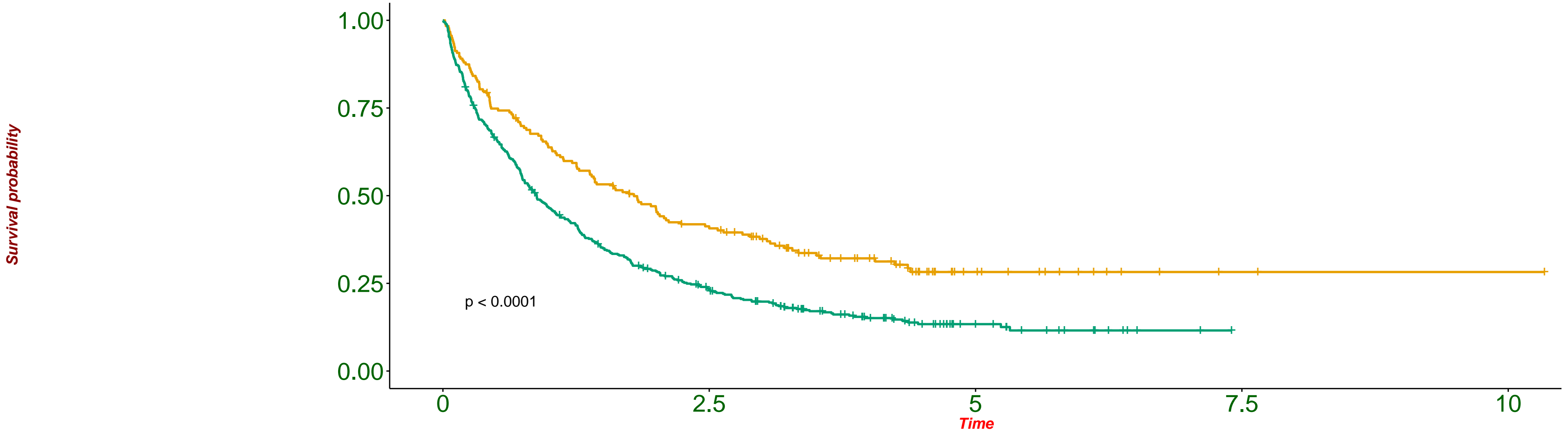


# Gene Repartition for Chromatin splicing components



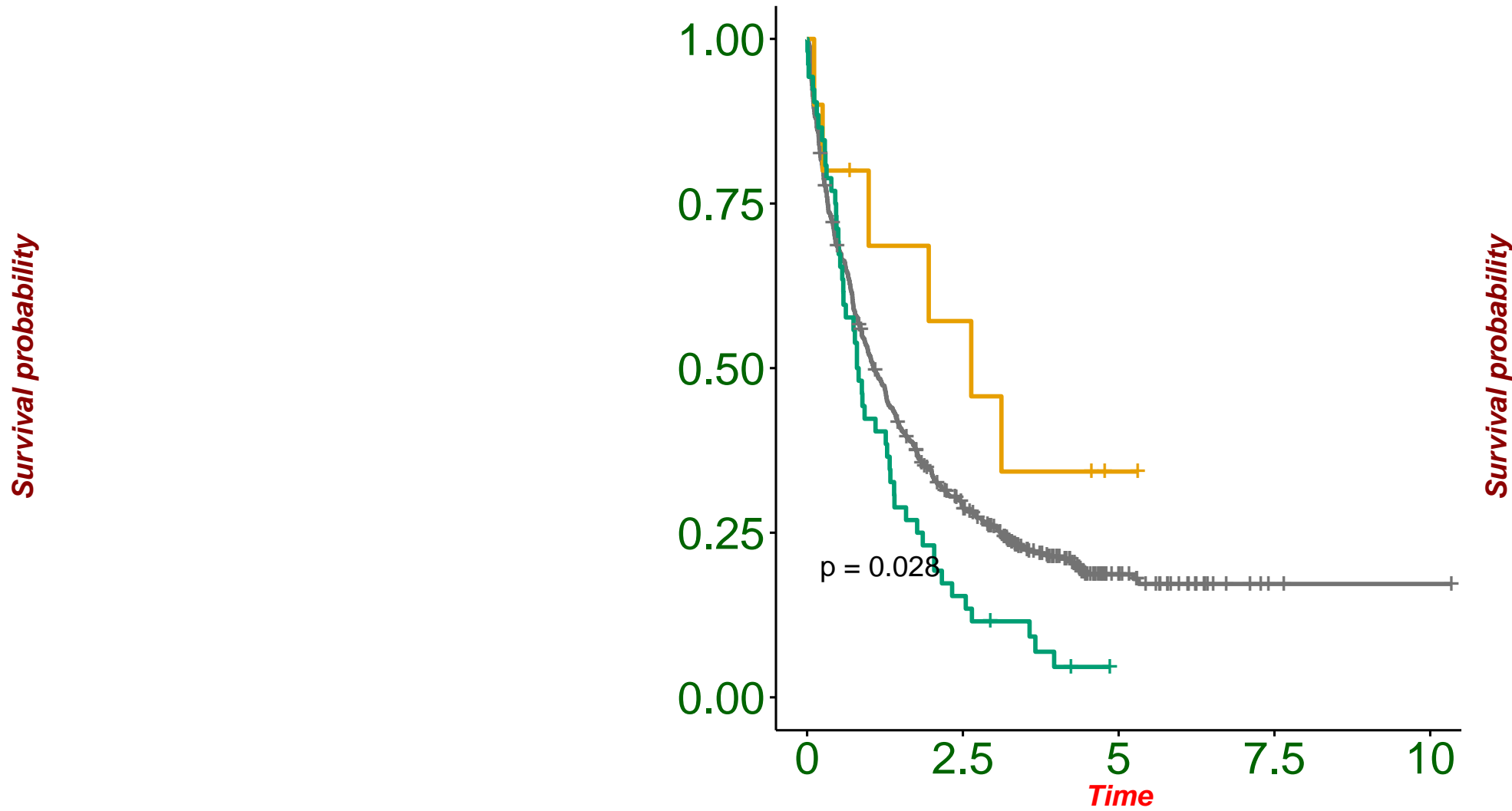
Kaplan–Meier estimates



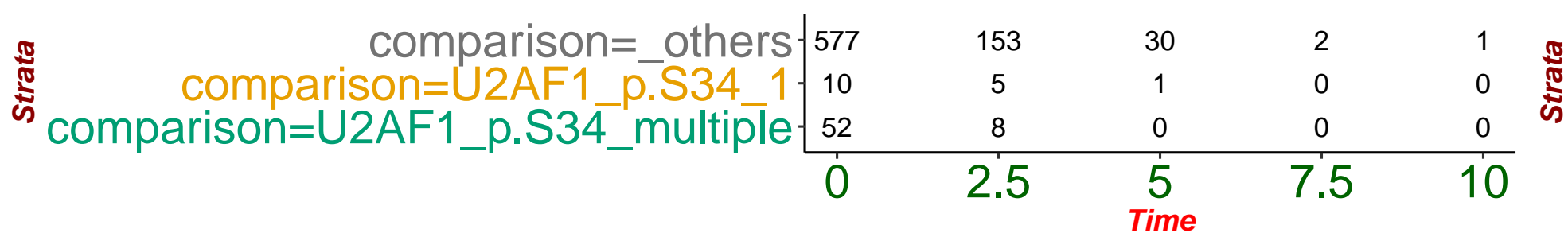
Number at risk



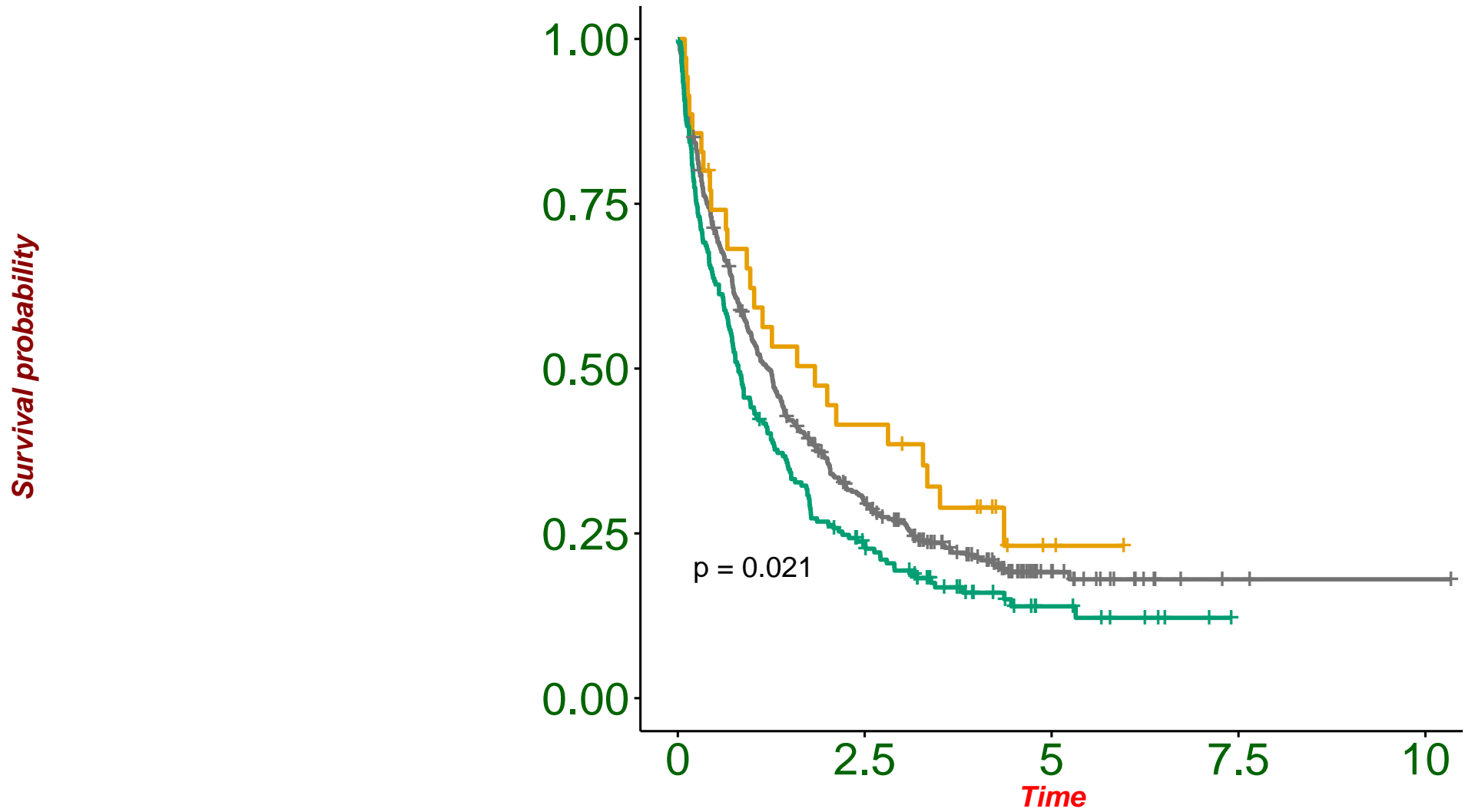
Kaplan–Meier estimates



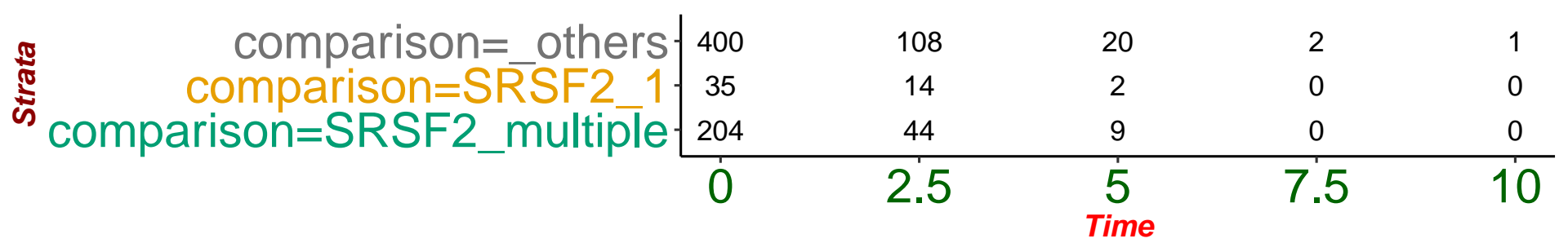
Number at risk

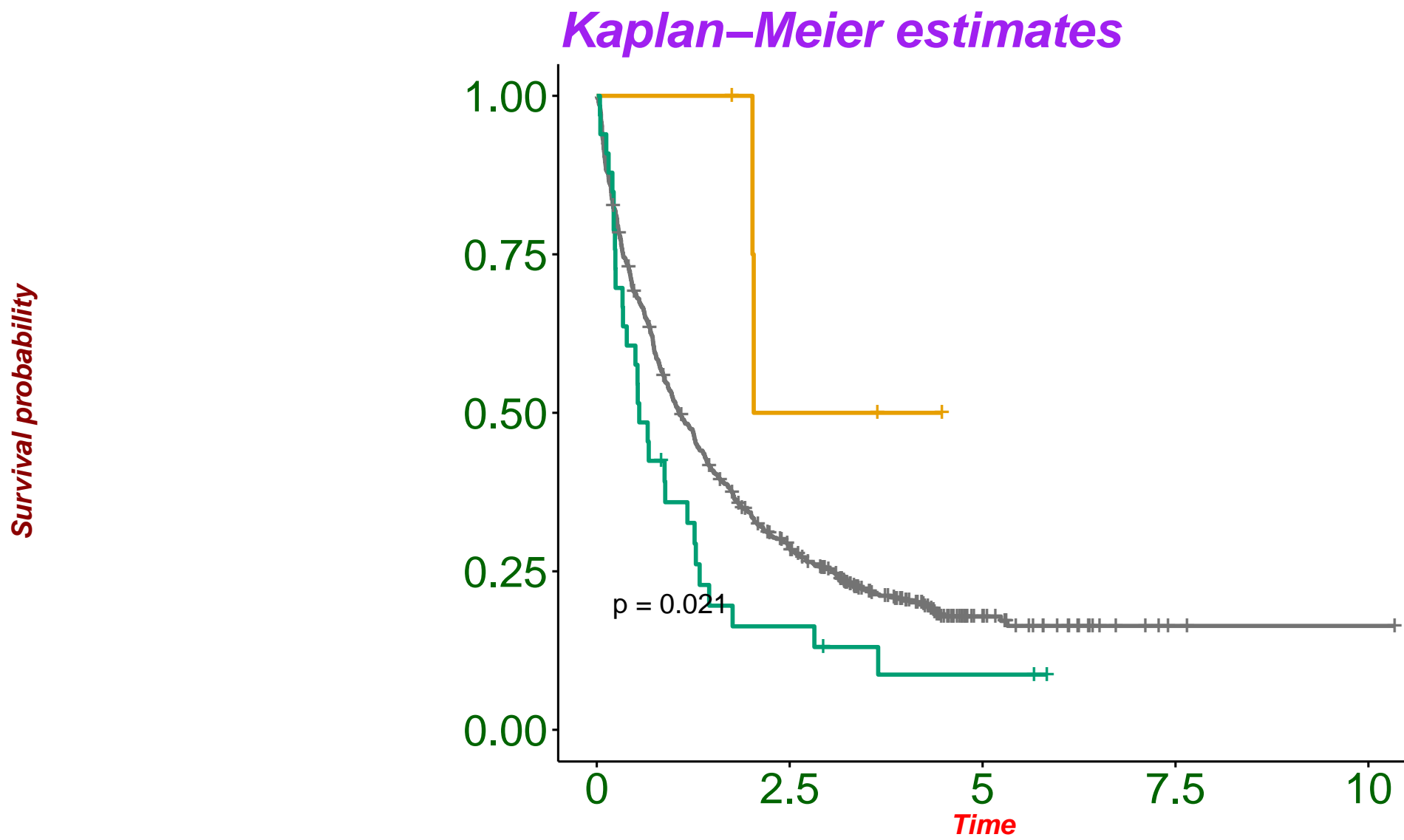


Kaplan–Meier estimates



Number at risk

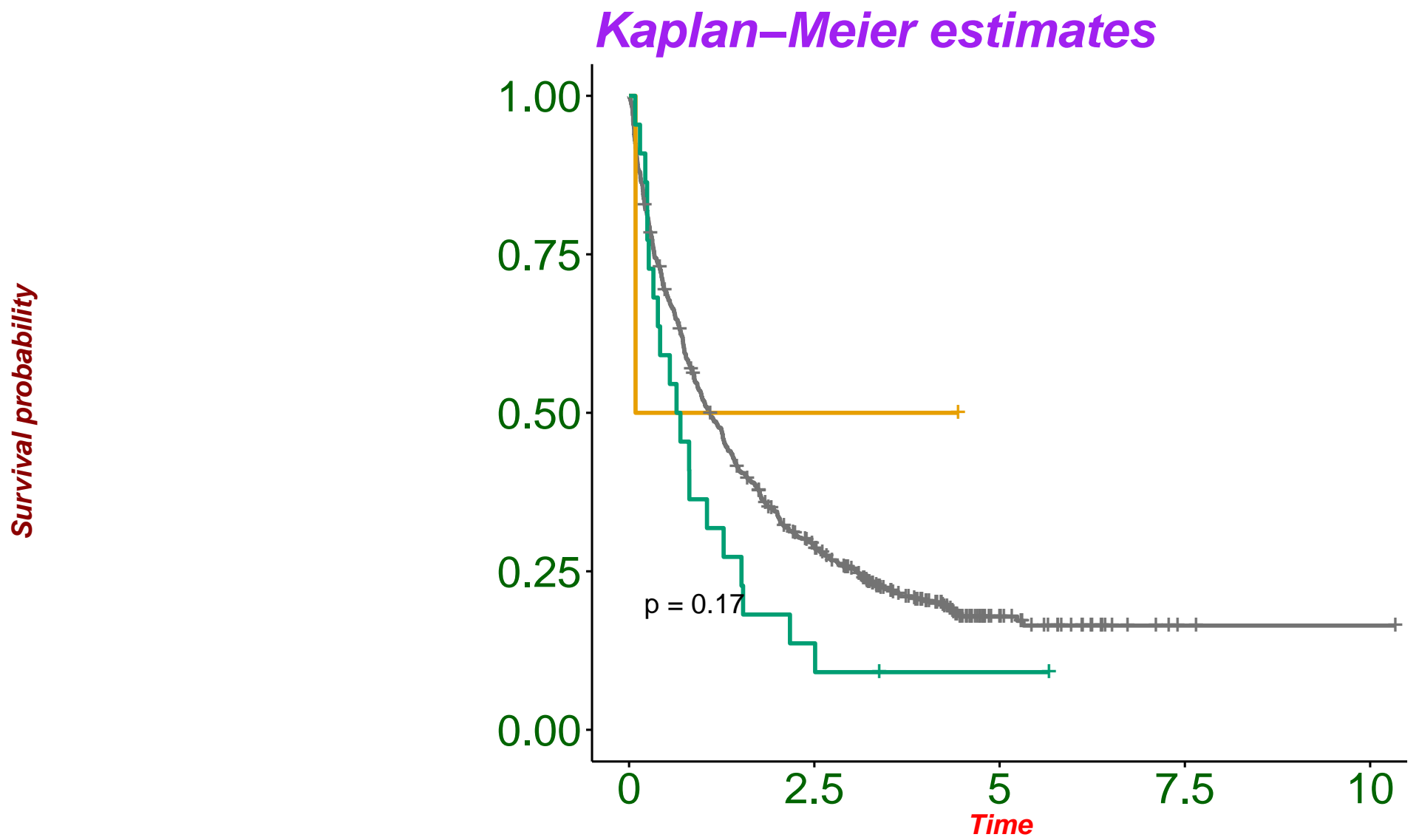




### Number at risk

comparison=_others	601	159	29	2	1
comparison=NF1_1	5	2	0	0	0
comparison=NF1_multiple	33	5	2	0	0

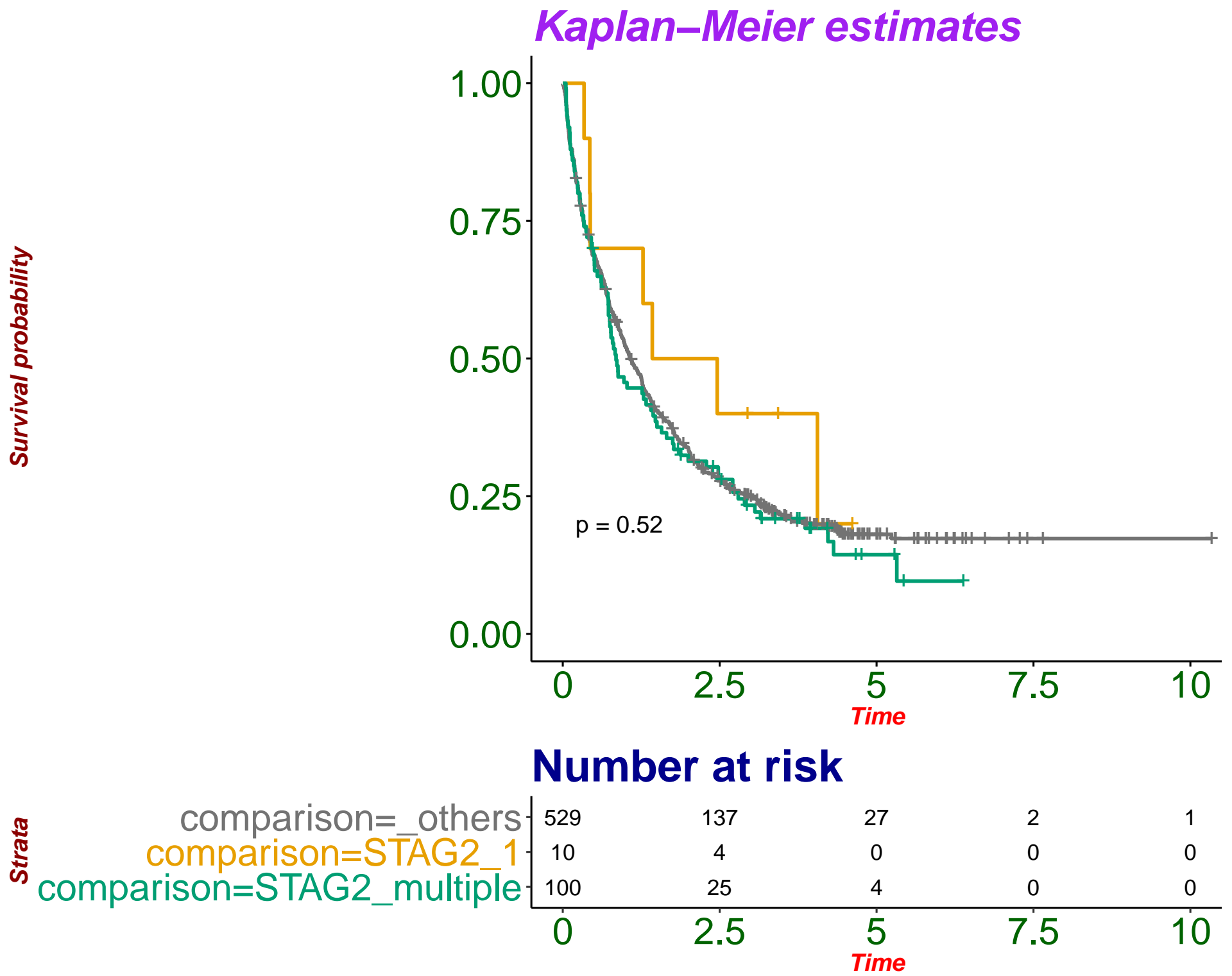
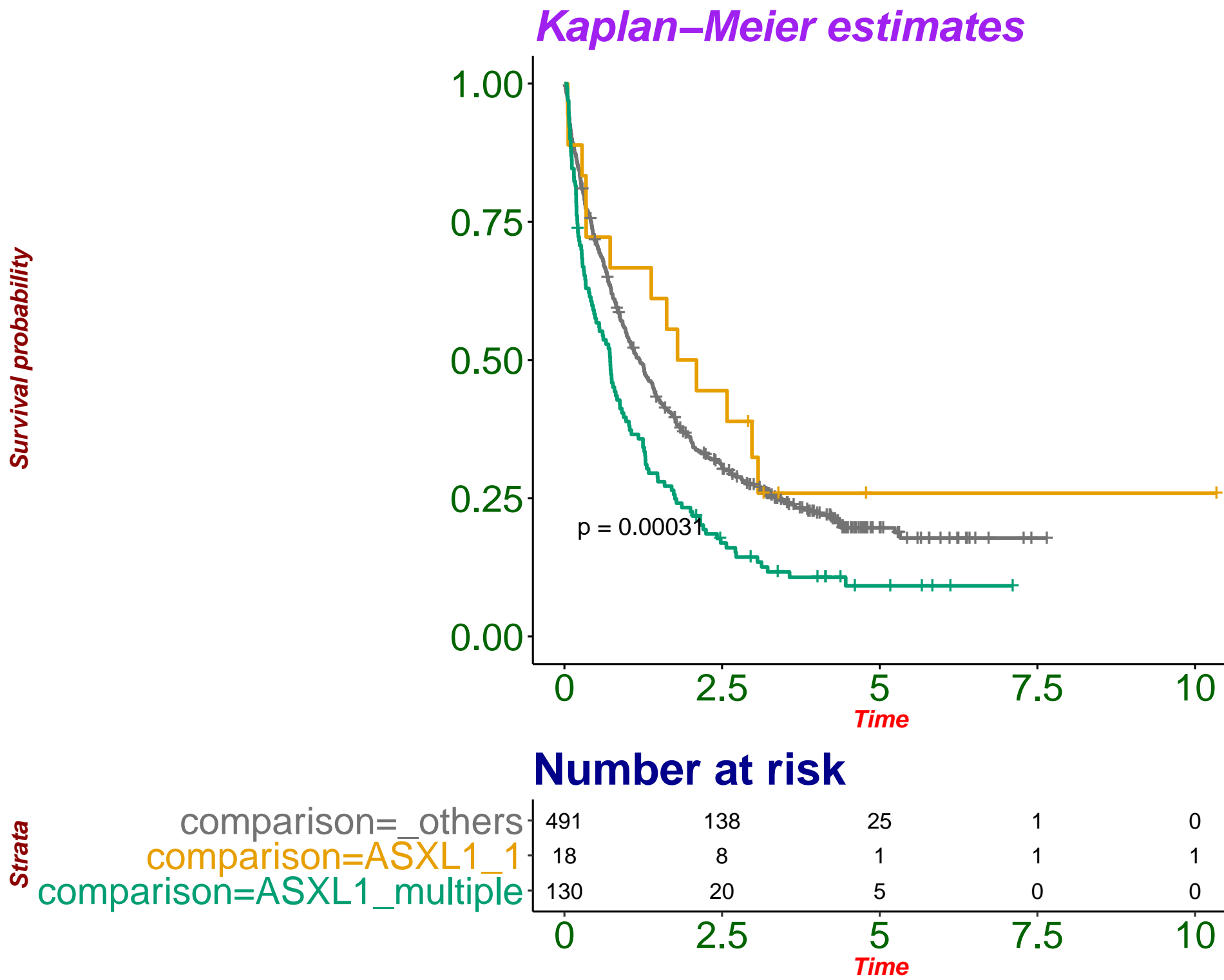
Time



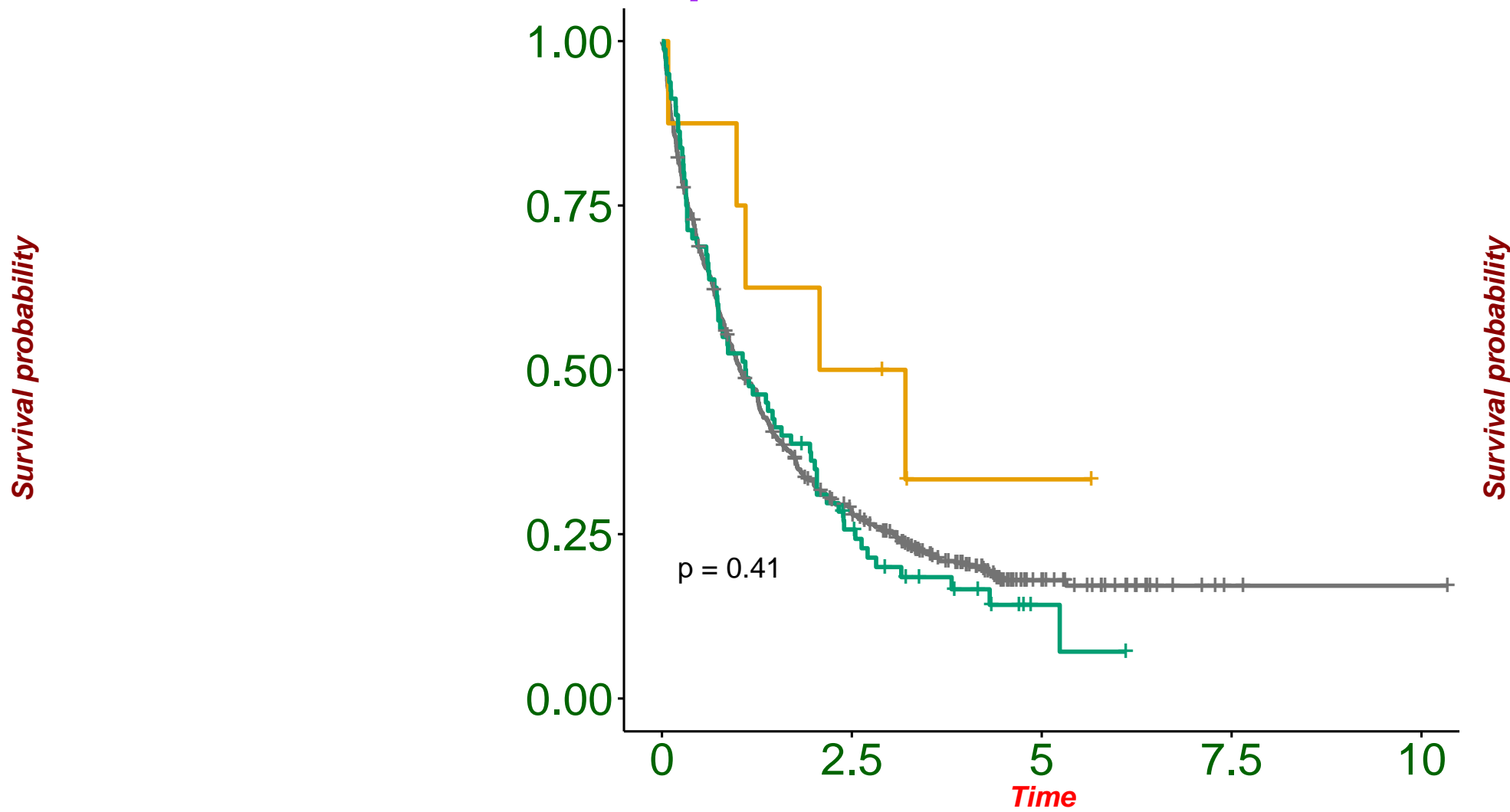
### Number at risk

comparison=_others	615	162	30	2	1
comparison=CUX1_1	2	1	0	0	0
comparison=CUX1_multiple	22	3	1	0	0

Time



Kaplan–Meier estimates



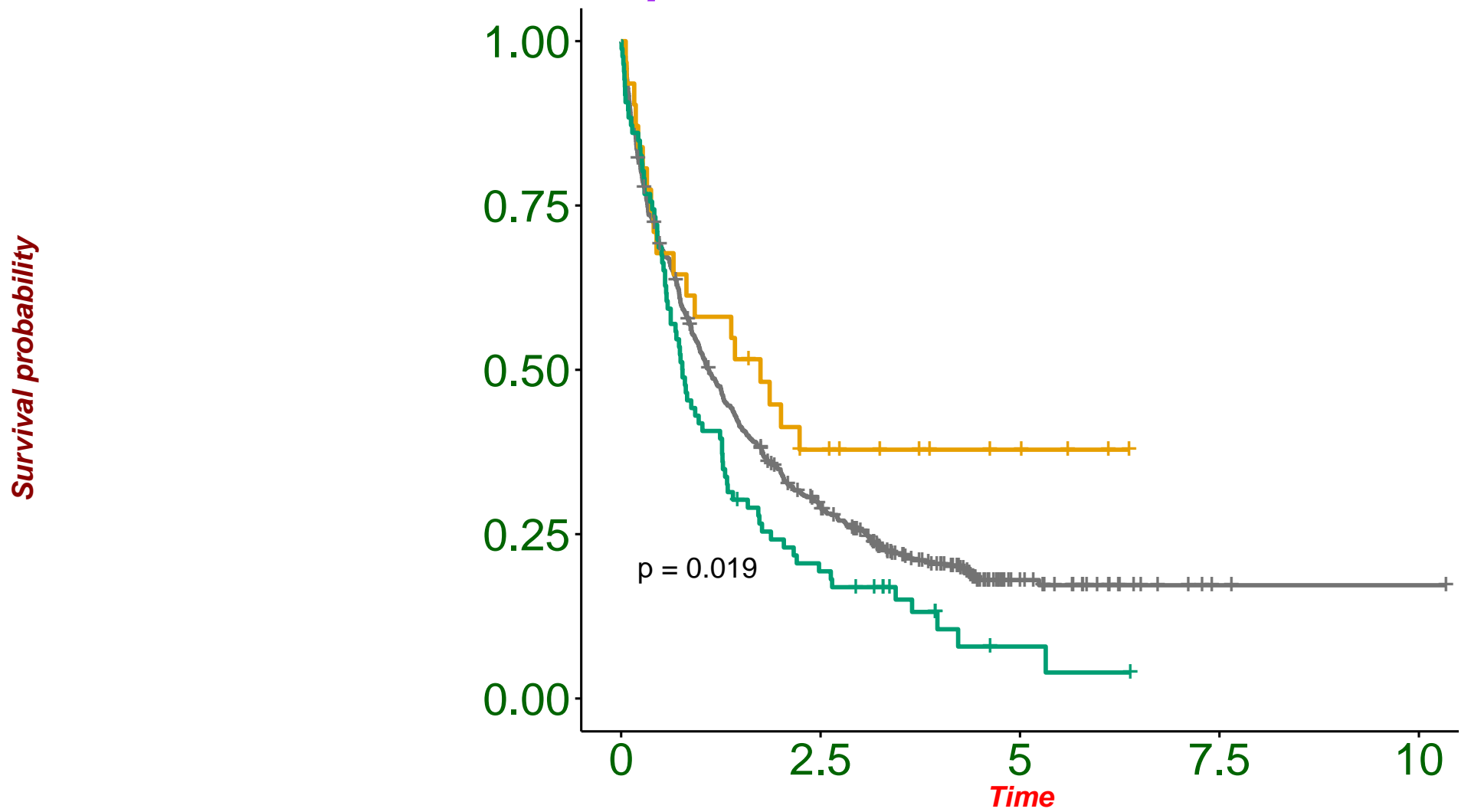
Number at risk

Strata

comparison=_others	551	143	28	2	1
comparison=BCOR_1	8	4	1	0	0
comparison=BCOR_multiple	80	19	2	0	0

Time

Kaplan–Meier estimates



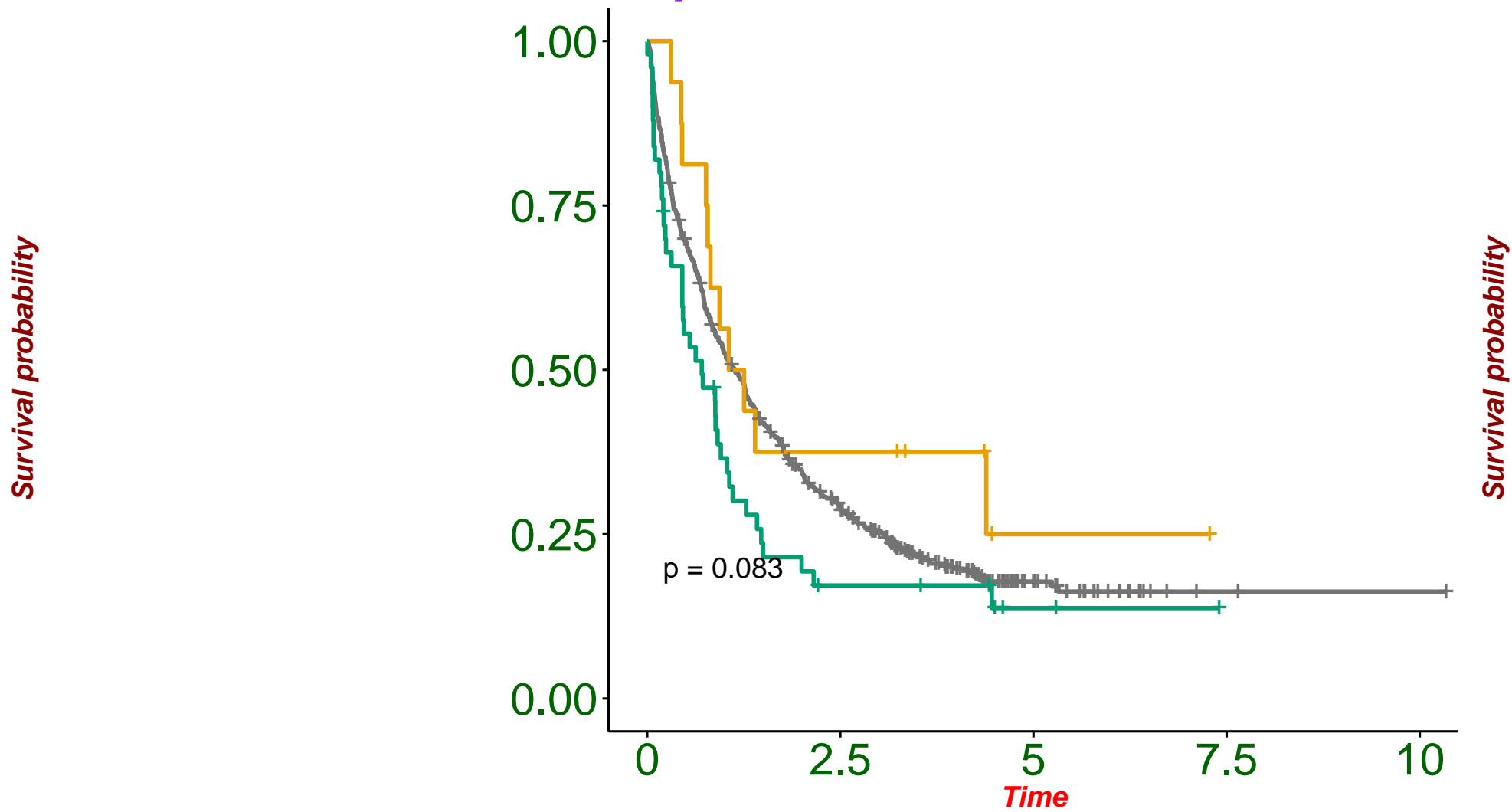
Number at risk

Strata

comparison=_others	522	140	25	2	1
comparison=MLL_1	31	10	4	0	0
comparison=MLL_multiple	86	16	2	0	0

Time

*Kaplan–Meier estimates*

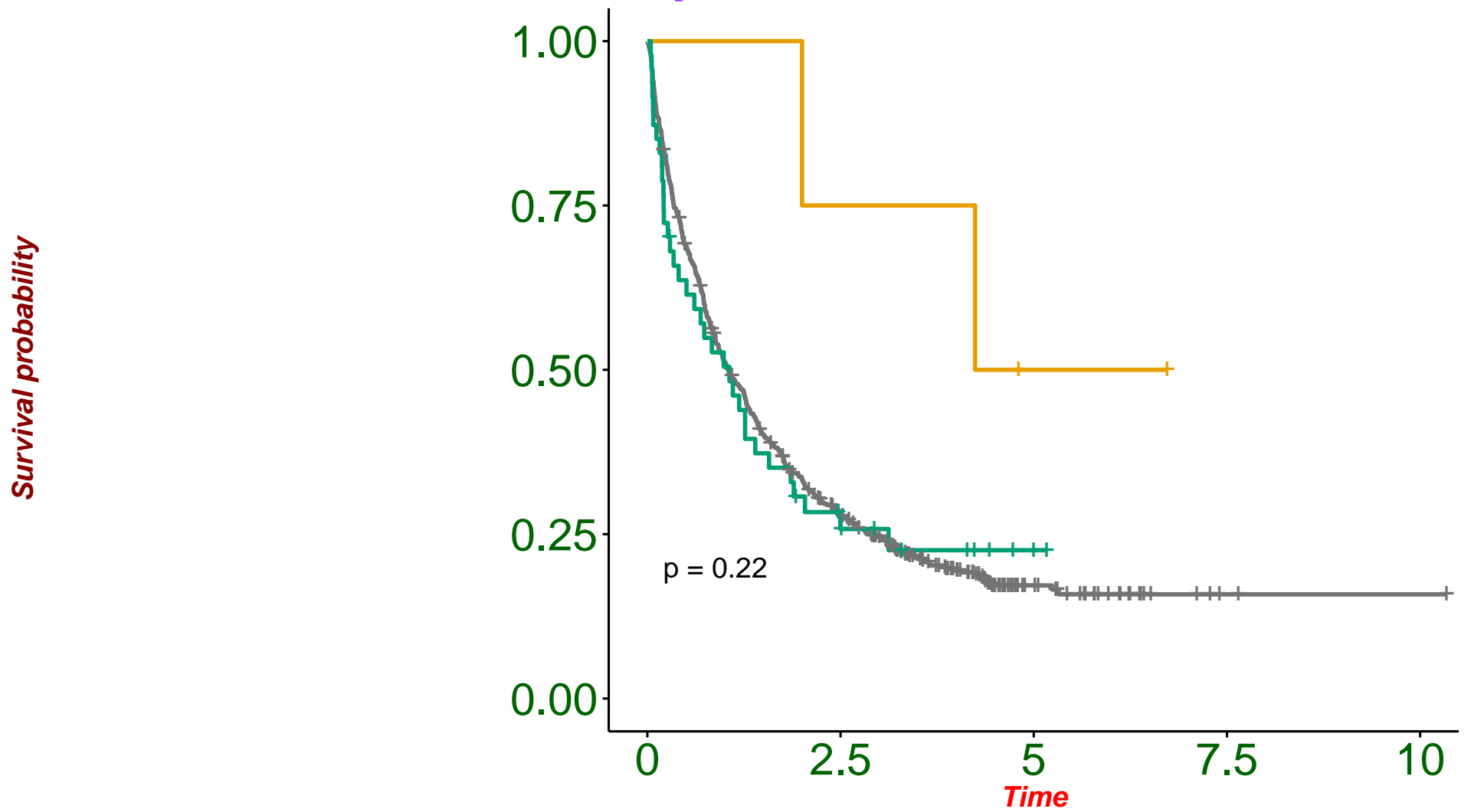


**Number at risk**

Strata	0	2.5	5	7.5	10
comparison=_others	573	153	28	2	1
comparison=EZH2_1	16	6	1	0	0
comparison=EZH2_multiple	50	7	2	0	0

Time

*Kaplan–Meier estimates*



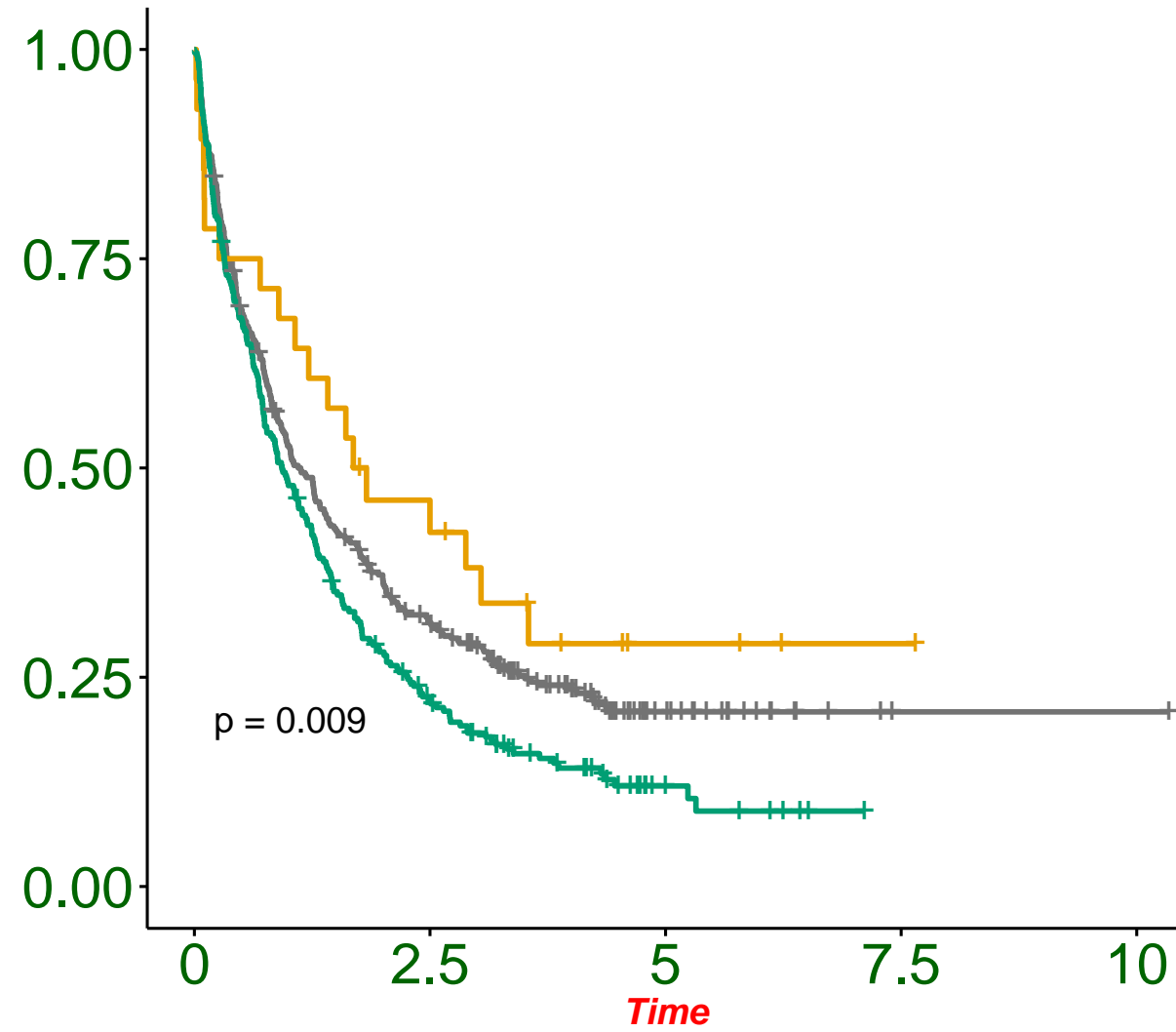
**Number at risk**

Strata	0	2.5	5	7.5	10
comparison=_others	588	153	29	2	1
comparison=PHF6_1	4	3	1	0	0
comparison=PHF6_multiple	47	10	1	0	0

Time

# Kaplan–Meier estimates

Survival probability



## Number at risk

Strata

comparison=_others	355	103	20	1	1
comparison=RUNX1_1	28	11	3	1	0
comparison=RUNX1_multiple	256	52	8	0	0
	0	2.5	5	7.5	10