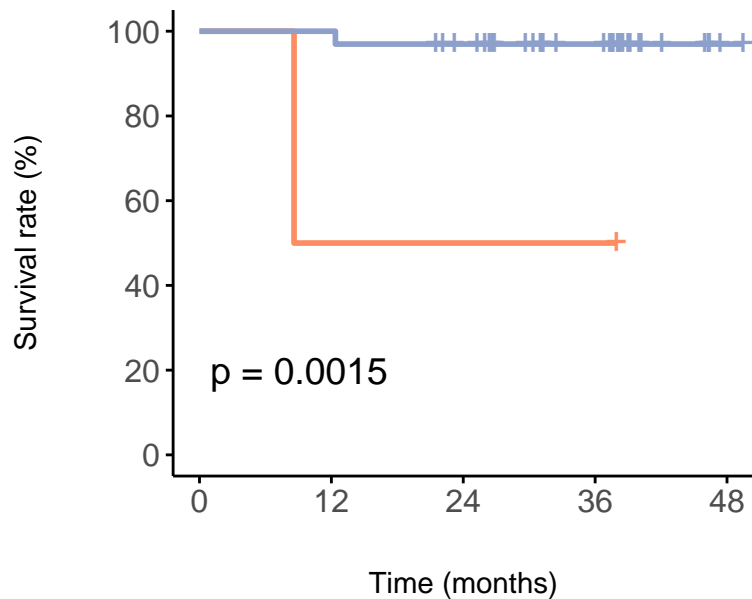


Number at risk

2	1	1	1	0
33	33	29	18	1

Pre-treatment ctDNA
Week 1 Vol

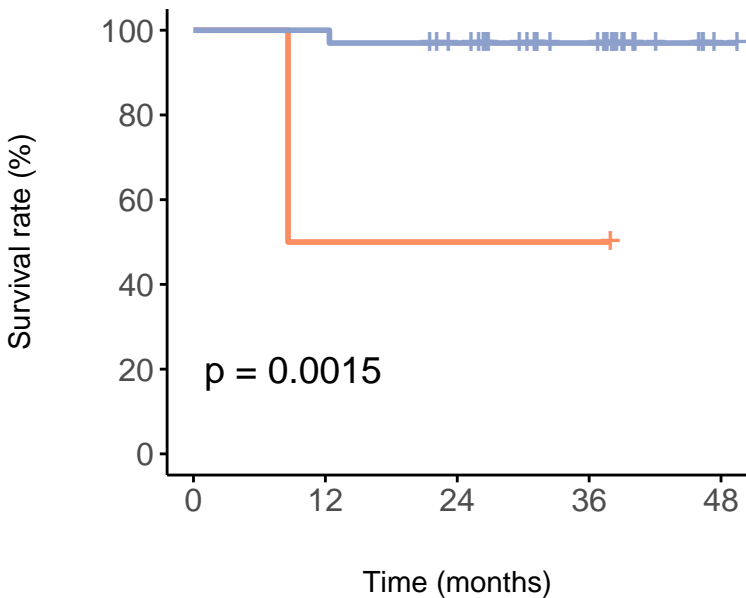


value=high value=low

Number at risk

2	1	1	1	0
33	33	29	18	1

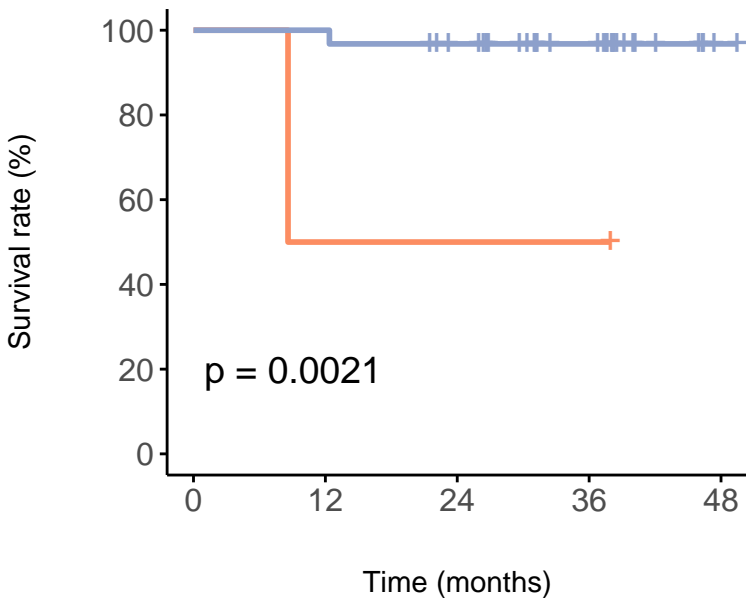
Pre-treatment ctDNA
Week 2 Vol



Number at risk

2	1	1	1	0
33	33	29	18	1

Pre-treatment ctDNA
Week 3 Vol

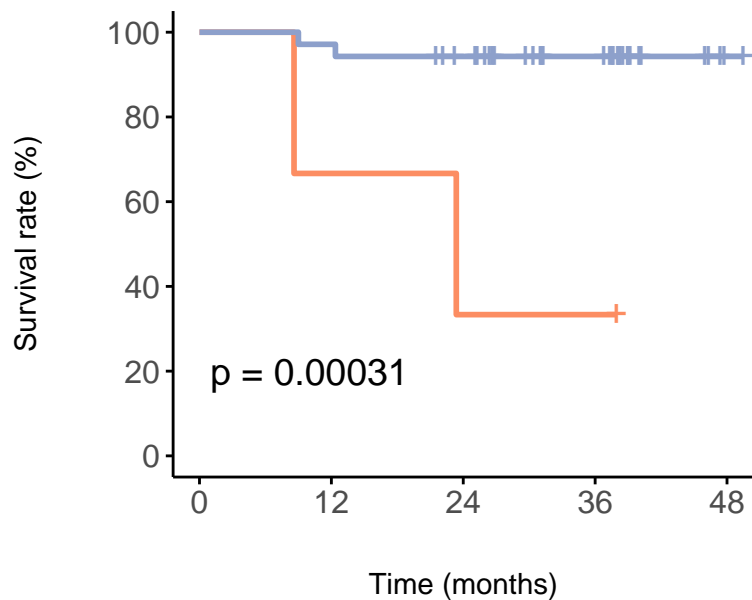


+ value=high + value=low

Number at risk

—	2	1	1	1	0
—	31	31	27	17	1

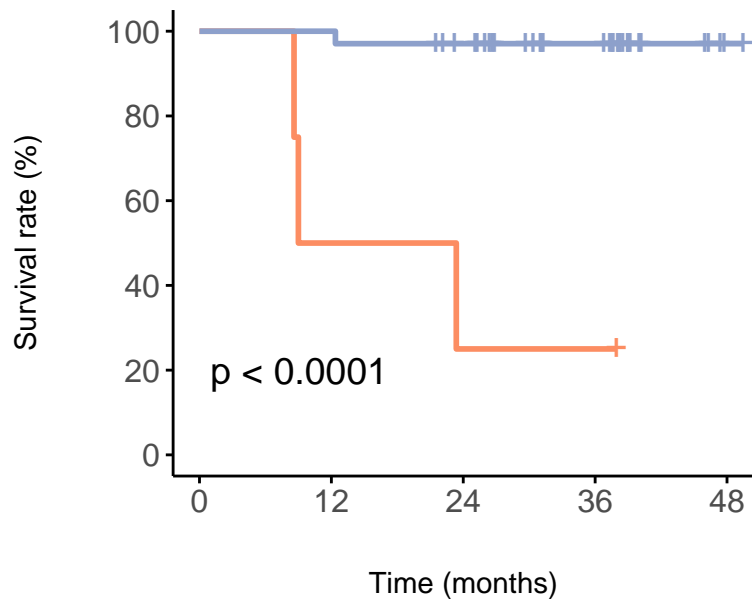
Week 1 ctDNA
Pre-treatment Vol



Number at risk

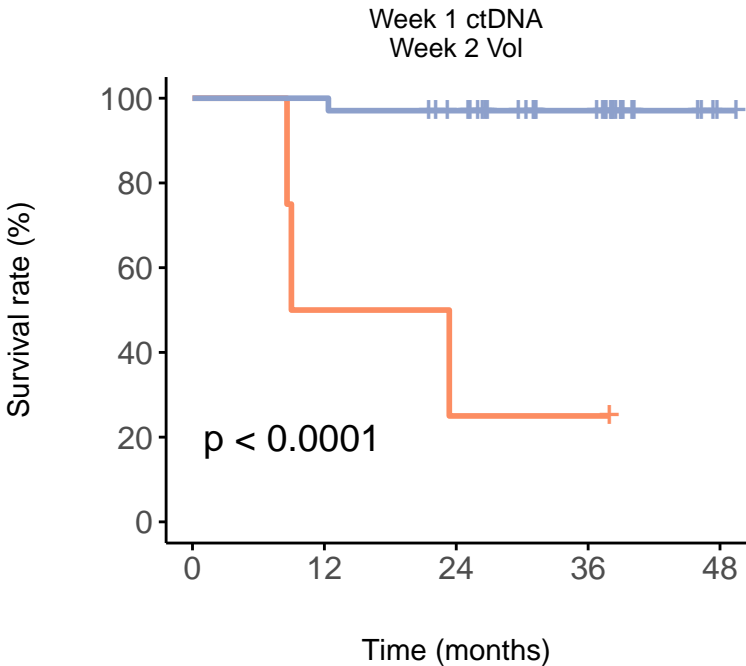
3	2	1	1	0
35	34	30	19	2

Week 1 ctDNA
Week 1 Vol



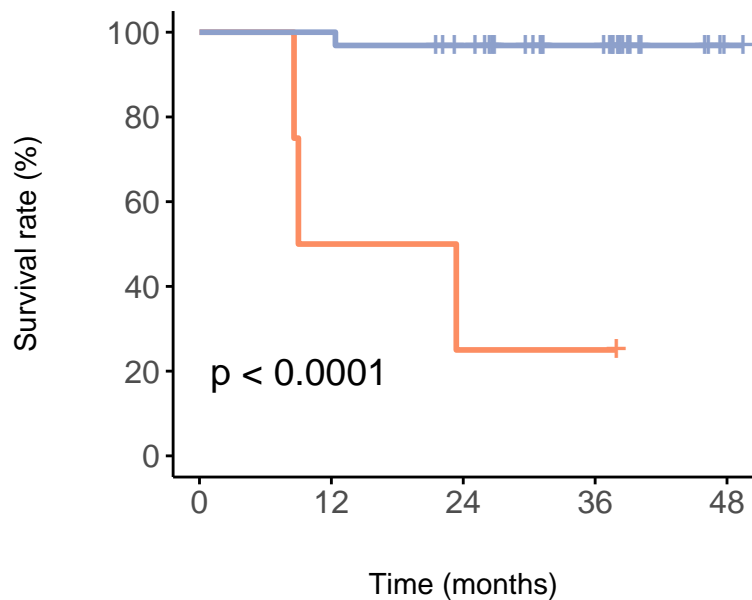
Number at risk

4	2	1	1	0
34	34	30	19	2



	0	12	24	36	48
value=high	4	2	1	1	0
value=low	34	34	30	19	2

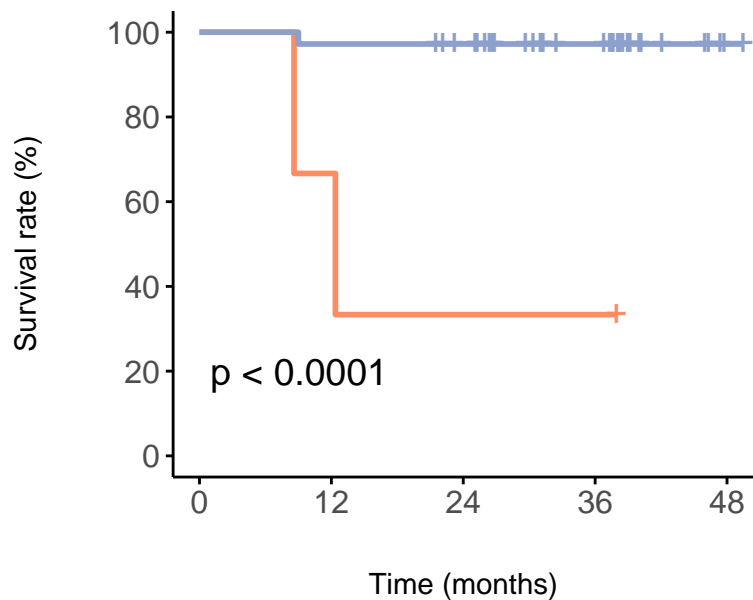
Week 1 ctDNA
Week 3 Vol



Number at risk

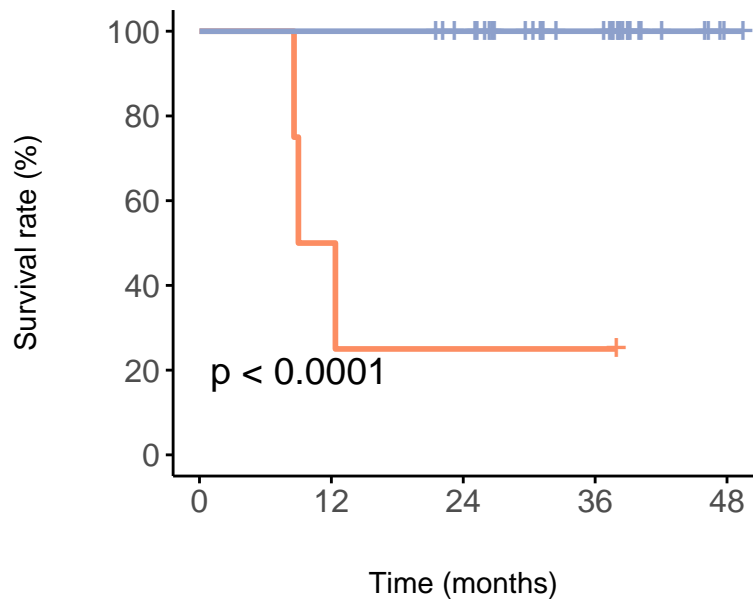
4	2	1	1	0
32	32	28	18	2

Week 2 ctDNA
Pre-treatment Vol



Number at risk					
value=high	3	2	1	1	0
value=low	36	35	32	20	2

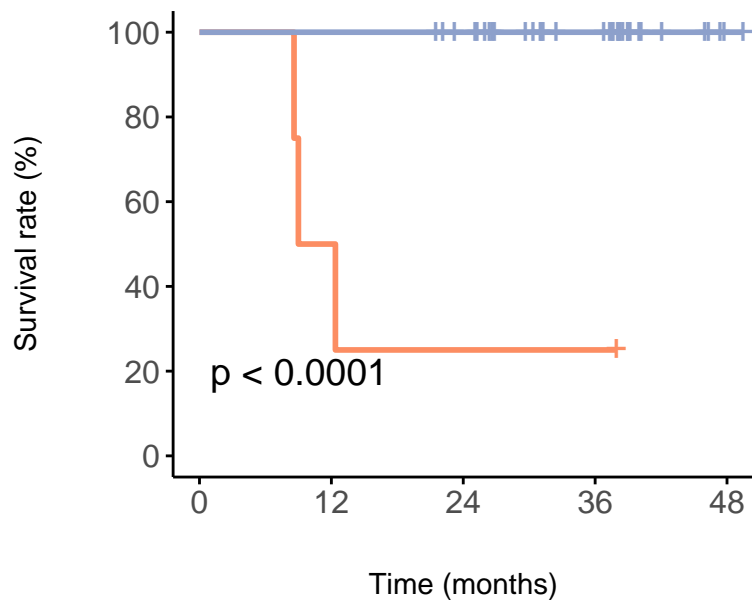
Week 2 ctDNA
Week 1 Vol



Number at risk

4	2	1	1	0
35	35	32	20	2

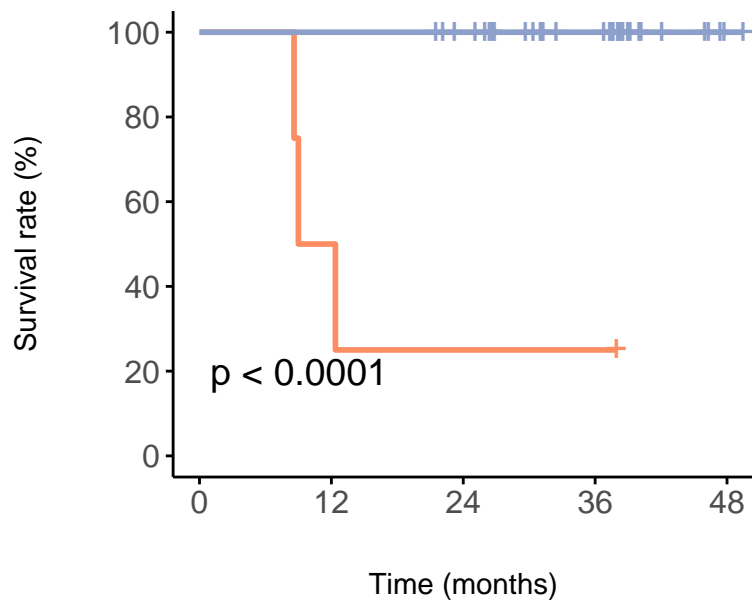
Week 2 ctDNA
Week 2 Vol



Number at risk

4	2	1	1	0
35	35	32	20	2

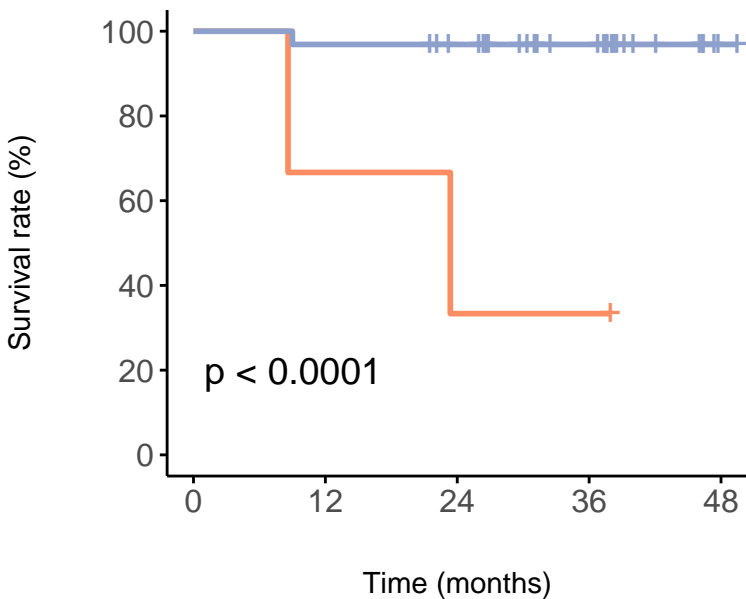
Week 2 ctDNA
Week 3 Vol



Number at risk

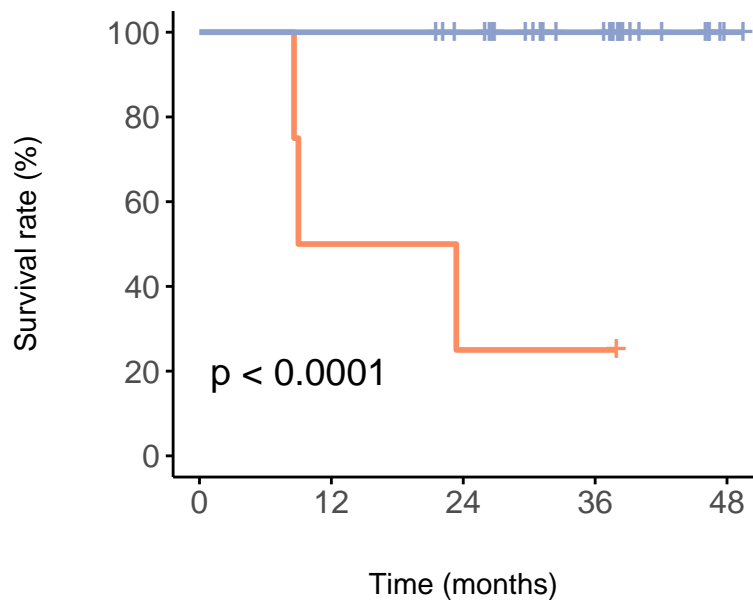
4	2	1	1	0
33	33	30	19	2

Week 3 ctDNA
Pre-treatment Vol



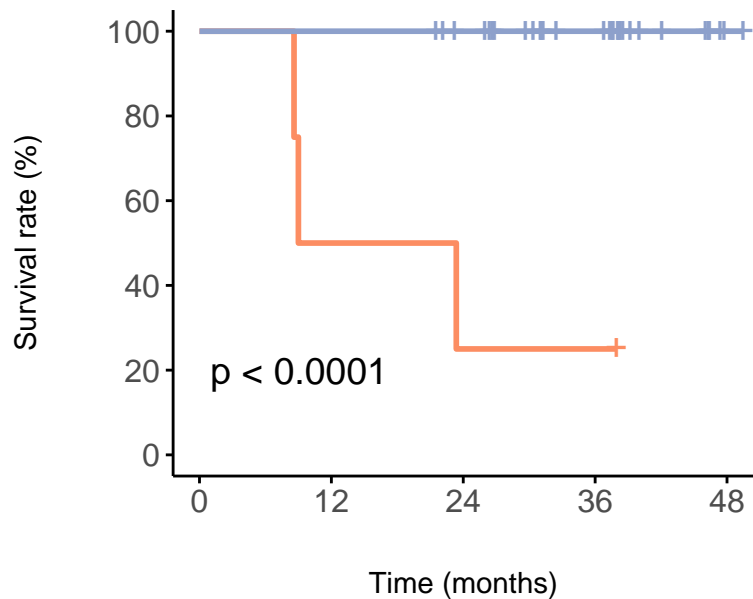
	Number at risk				
	0	12	24	36	48
value=high	3	2	1	1	0
value=low	32	31	28	18	2

Week 3 ctDNA
Week 1 Vol



	0	8	24	36	48
value=high	4	2	1	1	0
value=low	31	31	28	18	2

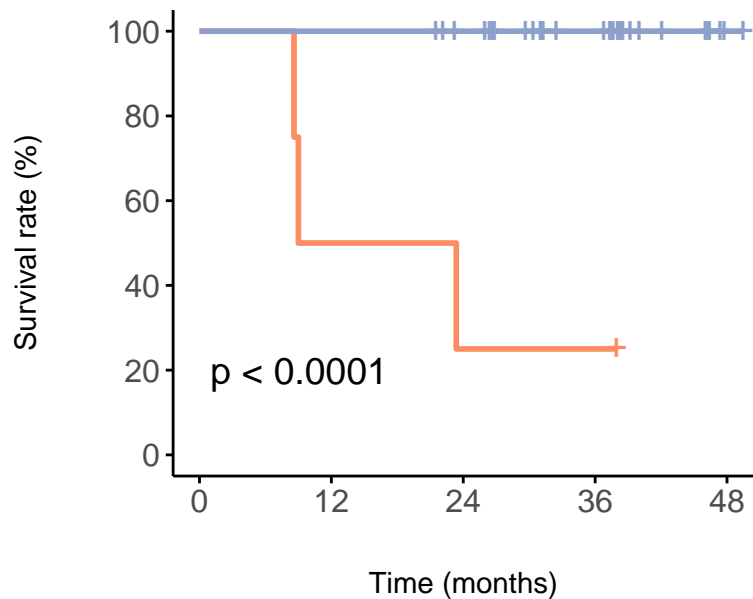
Week 3 ctDNA
Week 2 Vol



Number at risk

4	2	1	1	0
31	31	28	18	2

Week 3 ctDNA
Week 3 Vol



Number at risk

4	2	1	1	0
31	31	28	18	2