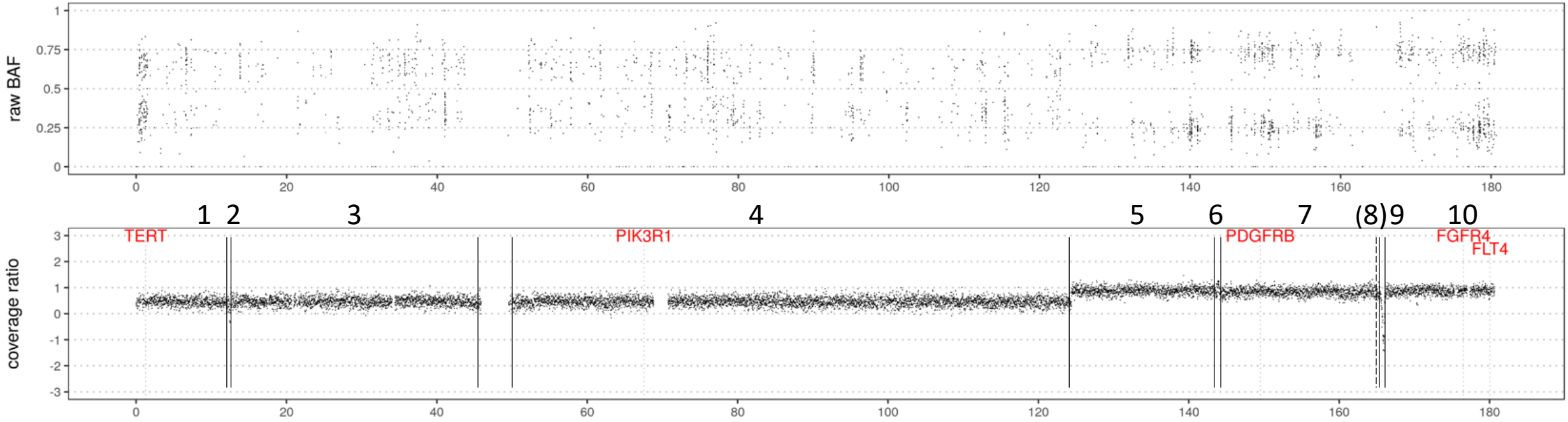


chr5 INF\_R\_652\_relapse2\_SARCOMA\_MPNST



9-10 segments judging from CN-plot

14 segments according to output file

6 segments, when merging segments next to each other and above a certain cut-off

	chr	start	end	Mb	ratio	zscore	
1	5	50001	12450000	12.40	0.47	32.86	1
2	5	12450001	12550000	0.10	-0.14	-2.99	2
3	5	12550001	34050000	21.50	0.50	38.73	3
4	5	34050001	40800000	6.75	0.54	63.52	
5	5	40800001	45900000	5.10	0.49	59.23	
6	5	49500001	124400000	74.90	0.50	151.25	4
7	5	124400001	143800000	19.40	0.92	219.17	
8	5	143800001	144050000	0.25	1.17	39.19	
9	5	144050001	159900000	15.85	0.90	153.09	
10	5	159900001	162700000	2.80	0.83	77.39	5
11	5	162700001	165450000	2.75	0.90	96.58	
12	5	165450001	165650000	0.20	0.62	19.53	6
13	5	165650001	166050000	0.40	-0.46	-15.07	
14	5	166050001	180750000	14.70	0.90	235.99	