

Supplementary Table 2. Chromosome alterations in matched ACC samples (Cochin cohort, 14 patients)

LOH, loss of heterozygosity

Patient	Sample	Tumor cells (%)	Altered genome (%)	Number of selected chromosome arms with complete LOH	Chromosome alteration profile
1	P	71	26.6	0	Noisy
	M	56	16.1	1	Noisy
2	P	67	38.8	2	Noisy
	R	74	38.1	2	Noisy
3	P	95	20.7	1	Noisy
	M	92	24.1	1	Noisy
4	P	92	64.4	1	Noisy
	M	79	66.6	0	Noisy
5	P	95	25.7	3	Noisy
	M	84	31	3	Noisy
6	P	72	40	2	Noisy
	M	88	61	4	Noisy
7	R	40	62.5	9	Chromosomal
	M	46	83.8	9	Chromosomal
8	R	60	77.2	9	Chromosomal
	M	92	79.9	9	Chromosomal
9	M1	NA	57.3	0	Noisy
	M2	91	57.4	0	Noisy
10	P1	67	63	9	Chromosomal
	P2	80	80.8	9	Chromosomal
11	P	88	68.6	9	Chromosomal
	M	100	53.2	9	Chromosomal
12	P	82	21.8	2	Noisy
	M	80	56.9	2	Noisy
13	P	51	74.3	9	Chromosomal
	M	80	88.2	9	Chromosomal
14	P1	73	11.6	1	Undetermined
	P2	88	11.7	1	Undetermined