

Table S2. The result of allele-specific copy number analyses.

Sample	Chromosome	Start	End	Total copy number	Major allele copy number	Minor allele copy number	Representative cancer-associated genes
C	3	51403500	90242000	1	1	0	
C	3	93874275	197960500	4	4	0	<i>PIK3CA</i>
C	5	140417	43175600	3	2	1	
C	8	232934	36121670	1	1	0	
C	8	46841213	117441100	4	3	1	
C	8	117465900	141168395	5	3	2	<i>MYC</i>
C	8	141175174	145054000	3	3	0	
C	9	154800	138175800	1	1	0	<i>SMARCA2, CDKN2A, CDKN2B, NOTCH1</i>
C	10	179900	38383216	1	1	0	
C	11	67522513	102955600	3	2	1	
C	11	103062300	134716165	1	1	0	<i>ATM</i>
C	13	19026389	22897511	1	1	0	
C	17	145200	26982000	1	1	0	
C	17	29204400	74381000	3	2	1	<i>HNF1B, ERBB2, STAT3</i>
C	17	74432100	81194100	1	1	0	<i>TP53</i>
C	X	288706	155238800	1	1	0	
AE	5	173523613	180902400	1	1	0	
AE	8	71962867	145054000	3	2	1	<i>MYC</i>
AE	11	68347900	68449100	2	2	0	
AE	11	68537800	103062300	2	2	0	
AE	11	103062500	134716165	2	2	0	<i>ATM</i>
AE	22	41599452	50780800	2	2	0	

Abbreviations in 'Sample' column are as follows. C: clear cell carcinoma, AE: adjacent endometriosis.