(a) BAC array and SNP array showing density of partial identity scores. Numbers indicate the number of aberrations:
- BAC array: 1237, 684, 5
- SNP array: 4, 3, 1, 2.

(b) Diagram depicting the relationship between normal tissue, primary tumor, and recurrence, with specific genetic alterations (+1q,-6p25.3,-6q24.2,-16q12.1-q24.3,-2q24.1-q32.3,-18p11.21-q23,-17p13.3-q11.2).

(c) Box plots comparing the number of aberrations in true recurrences (IR) and new primary tumors (PT) with CP.