



Supplementary material 5. Chromosomal breaks in ten carcinoma cell lines.

A. Number of breaks per an average chromosome detected by metaphase CGH, multicolour karyotyping (M-FISH) and three chromosome specific painting probes (chr2, 3 and 12). "nd": not done.

B. Number of breaks per chr3 detected by metaphase CGH, M-FISH, chr3 painting and multipoint FISH (mpFISH). "nd": not done.

C. Number of breaks within pericentromeric region (pericen), fragile sites (FRA), newly identified tumor break-prone regions (TBR) and other sites (others).