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Corrigendum

Corrigendum to “Tracing the tumor lineage” [Mol. Oncol. 4 (2010) 267–283]

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On page 276, right column, Figure 5c should read Figure 6c in line 8 and 10 from the bottom.

Due to an unfortunate error a wrong profile was displayed in the upper panel of Figure 6b. The text in the lower panel of

Figure 6c should read chr11p instead of chr10p. The corrected figure and legend are given below.

Figure 6 – Genomic progression in a metastatic breast cancer patient. (a) A frozen primary breast tumor and metastatic liver tumor resected from a single patient. (b) CGH profiles measured by Representational Oligonucleotide Microarray Analysis (ROMA). The segmented breast tumor profile is plotted in green and liver profile is plotted in red. (c) (Upper Panel) Segmented breast and liver profiles are plotted and a high Pearson's correlation was calculated ($c = 0.96$) between the copy number events. (Lower Panel) An enlarged region of Chromosome 11p is plotted showing the high similarity between complex rearrangements in the primary and metastatic tumors.

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