Erratum to: FGFR1 expression and gene copy numbers in human lung cancer

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There was an error in the original version of Fig. 7 representing the DNA imbalances of the 4 lung cancer subtypes as detected by conventional CGH. For large cell lung cancer (LCLC), the ideogram of chromosome

9 together with the histograms of DNA gains and losses was shown and not chromosome 8 harbouring the FGFR1 locus.

The corrected version with the chromosome 8 histogram (see new figure), however, indicates an identical finding, i.e. that chromosome 8p is more often affected by deletions than DNA gains.

The online version of the original article can be found at <http://dx.doi.org/10.1007/s00428-012-1250-y>.

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Fig. 7 DNA imbalances of lung cancer subtypes according to conventional comparative genomic hybridization/CGH (compiled from data of Petersen I in Travis et al. 2004, [30]). Cases with DNA gains were represented by the histogram on the *right side* of the chromosome

ideogram, cases with DNA loss on the *left side*. The data indicated that 8p deletions are at least as frequent as DNA gains with pronounced gains (indicated by the *red colour*) as putative correlates of an amplification being undetectable in the NSCLC subtypes