Erratum to: Loss of heterozygosity analysis in malignant gliomas

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Unfortunately, in Table 1 of the article cited above, the 3rd line in the first column appeared incorrectly.

Table 1 should be correctly shown as follows:

Table 1 Comparison of common molecular assays for LOH detection

	PCR-based assay	FISH	CGH (aCGH)	SNP array
Allelic imbalance	○	×	×	○
Copy number alterations	×	○	○	○
Copy neutral LOH (UPD) (mitotic recombination)	○	×	×	○
Homozygous deletions	×	○	○	○
Matching DNA required	○	×	○	Δ ^a

^a High resolution SNP array can detect LOH regions without matching DNA

The online version of the original article can be found under doi:[10.1007/s10014-011-0038-0](https://doi.org/10.1007/s10014-011-0038-0).

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