

## Correction

# Correction: Comparison of Hepatocellular Carcinoma miRNA Expression Profiling as Evaluated by Next Generation Sequencing and Microarray

## The PLOS ONE Staff

In Table 2, the headings Tumor and Control are incorrectly swapped in the second column and in the second row. Please see the corrected Table 2 here.

**Table 2.** Performance of discrimination between 14 HCC samples and 6 normal tissue samples using miRNA expression obtained by NGS analysis.

		Result	
		Tumor	Control
Prediction	Tumor	12	0
	Control	2	6

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## Reference

1. Murakami Y, Tanahashi T, Okada R, Toyoda H, Kumada T, et al. (2014) Comparison of Hepatocellular Carcinoma miRNA Expression Profiling as Evaluated by Next Generation Sequencing and Microarray. PLoS ONE 9(9): e106314. doi:10.1371/journal.pone.0106314

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