

# Corrigendum: MicroRNA122 is a key regulator of $\alpha$ -fetoprotein expression and influences the aggressiveness of hepatocellular carcinoma

Kentaro Kojima, Akemi Takata, Charles Vadnais, Motoyuki Otsuka, Takeshi Yoshikawa, Masao Akanuma, Yuji Kondo, Young Jun Kang, Takahiro Kishikawa, Naoya Kato, Zhifang Xie, Weiping J. Zhang, Haruhiko Yoshida, Masao Omata, Alain Nepveu & Kazuhiko Koike

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This Article contains errors in the numbering of several papers in the reference list; reference 34 is incorrectly listed as reference 44 and references 35 to 44 are incorrectly listed as 34 to 43. The correct numbering is as follows.

34. Cadieux, C. *et al.* Mouse mammary tumor virus p75 and p110 CUX1 transgenic mice develop mammary tumors of various histologic types. *Cancer Res.* **69**, 7188–7197 (2009).
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36. Michl, P. *et al.* CUTL1 is a target of TGF( $\beta$ ) signaling that enhances cancer cell motility and invasiveness. *Cancer Cell* **7**, 521–532 (2005).
37. Aleksic, T. *et al.* CUTL1 promotes tumor cell migration by decreasing proteasome-mediated Src degradation. *Oncogene* **26**, 5939–5949 (2007).
38. Kunath, T. *et al.* Transgenic RNA interference in ES cell-derived embryos recapitulates a genetic null phenotype. *Nat. Biotechnol.* **21**, 559–561 (2003).
39. Shouval, D. *et al.* Tumorigenicity in nude mice of a human hepatoma cell line containing hepatitis B virus DNA. *Cancer Res.* **41**, 1342–1350 (1981).
40. Nomura, F., Ohnishi, K. & Tanabe, Y. Clinical features and prognosis of hepatocellular carcinoma with reference to serum alpha-fetoprotein levels. Analysis of 606 patients. *Cancer* **64**, 1700–1707 (1989).
41. Johnson, P., Melia, W., Palmer, M., Portmann, B. & Williams, R. Relationship between serum alpha-fetoprotein, cirrhosis and survival in hepatocellular carcinoma. *Br. J. Cancer* **44**, 502–505 (1981).
42. Xu, H. *et al.* Liver-enriched transcription factors regulate microRNA-122 that targets CUTL1 during liver development. *Hepatology* **52**, 1431–1442 (2010).
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