

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

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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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