



Correction to: α 2,6-Sialylation promotes immune escape in hepatocarcinoma cells by regulating T cell functions and CD147/MMP signaling

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Subsequently to the publication of this article, the authors have noticed that the published version of Fig. 4H contained incorrect data showing the migration of the Mock cells (left panel, Hepa1-6 cells transfected with pcDNA3.1). The correct Fig. 4 and legend are given below. This mistake did not affect the conclusions of this work. The authors apologize to the readers for any inconvenience caused for this error.

Fig. 4 Upregulation of ST6Gal-I promotes tumor cell proliferation and migration in murine hepatoma carcinoma cells. Overexpression of ST6Gal-I in Hepa1-6 and Hca-P cells were confirmed by real-time PCR (**a, d**), Western-blot assays (**b, e**), and SNA lectin blot (**c, f**). **g** Cell proliferation ability measured by a Cell Counting Kit-8 assay. **h** Transwell migration assay was used to determine the migration ability of cells. Results are the mean of three independent experiments \pm SD ($*P < 0.05$, $**P < 0.01$). Hepa1-6/St or Hca-P/St, pcDNA3.1/ST6Gal-I vector transfected stable clones

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