Author Correction: Musashi 1 regulates the timing and extent of meiotic mRNA translational activation by promoting the use of specific CPEs

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Correction to: *Nature Structural & Molecular Biology* <https://www.nature.com/articles/nsmb.3434>, published online 17 July 2017.

In the version of this article originally published, the legend to Supplementary Fig. 3 was incorrect (included text describing a panel not presented there (cited as b) and cited the wrong panel letter (c) for the text describing panel b). The correct legend is as follows: “Immature oocytes were injected with the indicated RNA constructs, then overexpressed HA-Msi1 or endogenous CPEB1 were immunoprecipitated. The IPs were analyzed by western blot and by RT qPCR. Anti-HA IP without HA-Msi1 overexpression and IgG IP were used as controls. (a) Left panel, IP HA-Msi1 analyzed by western blot. Right panel, qPCRs of the indicated reporters performed using reversed transcribed immunoprecipitated mRNAs. Input was normalized against endogenous *GAPDH* mRNA, and IP efficiency, against endogenous *c-mos* mRNA. (b) Translational regulation of Mos (+12) and non-consensus CPE containing reporters. Synthetic mRNAs containing the firefly luciferase coding sequence fused to the indicated 3′ UTRs were co-injected with control renilla luciferase mRNA. Oocytes were collected 2 h after GVBD to determine luciferase activity. Schematic representations of each 3′ UTR are shown to the left, with consensus CPEs (C) depicted as blue boxes; non-consensus CPEs, green boxes; MBEs (M), orange circles; PAS (H), red hexagons; and mutated CPE, MBE or PAS, as an empty box, circle or hexagon, respectively; distance between elements is given in nucleotides.”

Also, the directions (left, right) in the legend for Supplementary Fig. 4a were incorrect; the correct panel legend is as follows: “(a) Effects of MBEb mutation on CPEB1 binding. Immature oocytes were injected with the indicated constructs. Upper panel, IP against CPEB1 analyzed by western blot; lower panel, qPCR of the indicated reporter after reverse transcription of the immunoprecipitated mRNAs (data, mean $\pm$ SEM; $n = 3$).”

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