



Corrigendum: Cystathionine β -Synthase Is Necessary for Axis Development *in vivo*

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A Corrigendum on

Cystathionine β -Synthase Is Necessary for Axis Development *in vivo*

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In the original article, there was an error in **Figure 2C** as published. The sequence of the *cbasa splice 1* (CBSA-S1) morpholino was incorrectly typed as

TACCTGCACAAAGTGAACACAACCA

The correct sequence is

TACCTGCACAAAGTGAACACAACCA

The name of the morpholino was changed in the figure from *cbasa splice* (CBSA-S1) to *cbasa splice 1* (CBSA-S1) to match the legend.

The corrected **Figure 2** appears below. The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way.

The original article has been updated.

Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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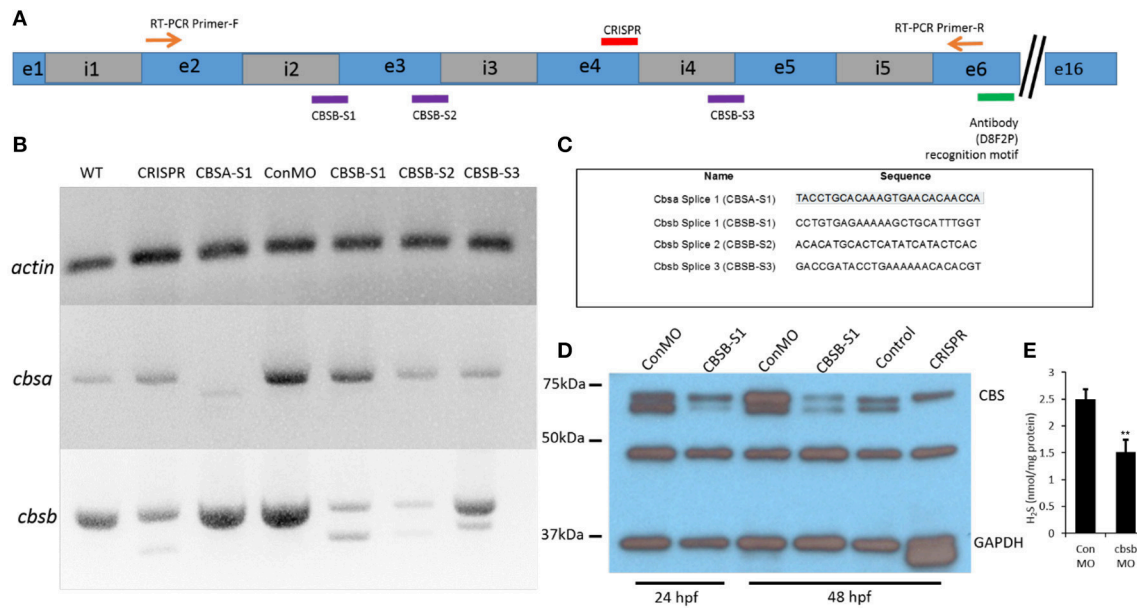


FIGURE 2 | Loss-of-function efficacy studies. Panel **(A)** shows a “partial” cartoon representation of the *cbsb* genomic site with the location of the MO sites (purple rectangles) at appropriate intron (i) and exon (e) junctions, CRISPR-targeted site (red rectangle), site of RT-PCR forward (F) and reverse (R) primers, and the antibody recognition site. Panel **(B)** shows RT-PCR for three genes (*cbsb*, *cbsa*, *actin*) in total RNA from injected embryos (~24 hpf) (left to right): wild type (WT) control and *cbsb* CRISPR-injected fish, *cbsa* splice 1 (CBSA-S1), control morpholino (ConMO), *cbsb* splice1 (CBSB-S1), *cbsb* splice 2 (CBSB-S2), *cbsb* splice 3 (CBSB-S3). Panel **(C)** shows the sequence of the morpholinos used in this study. Panel **(D)** shows CBS and GAPDH western blots for ConMO, CBSB-S1 at 24 and 48 hpf along with control and *cbsb* CRISPR fish at 48 hpf. Panel **(E)** shows the comparison between CBSB-S1 MO and ConMO-injected embryos for hydrogen sulfide production. $n = 3$ for both groups (data from three experiments). Twenty embryos in each group in each experiment. $**P < 0.01$.