

CORRECTION

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# Correction to: Proximity ligation scaffolding and comparison of two *Trichoderma reesei* strains genomes

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## Correction to: *Biotechnol Biofuels* (2017) 10:151

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Following publication of the original article [1], the authors reported a problem in the drawing of Rut-C30 chromosome III in Fig. 2b of the original article [1]. The two fragments of chromosome I inserted inside chromosome III should be swapped, and the direction of the fragment containing rim101 and cella genes should be inverted. This reversed insertion indicates that at least 2 rearrangements occurred simultaneously, so the possible scenario proposed in Fig. 2c of the original article was inaccurate. The corrected Fig. 2 with modified panels b and c is available in this erratum. The detailed description

of Rut-C30 assembly in the Additional file 5 of the original article is correct.

The authors also noticed two mistakes in chromosome numbering in the description of these translocations. The correct description is

“The three translocations resulted finally in the right arm of chromosome I (3' end of scaffold 48 and main fragment of scaffold 5: 1.63 Mb and 442 genes in total), to be swapped with the right arm of chromosome VI (3' end of scaffold 22: 402 kb and 114 genes). But also in two fragments of chromosome I (one with a fragment of scaffold 4, and the other one

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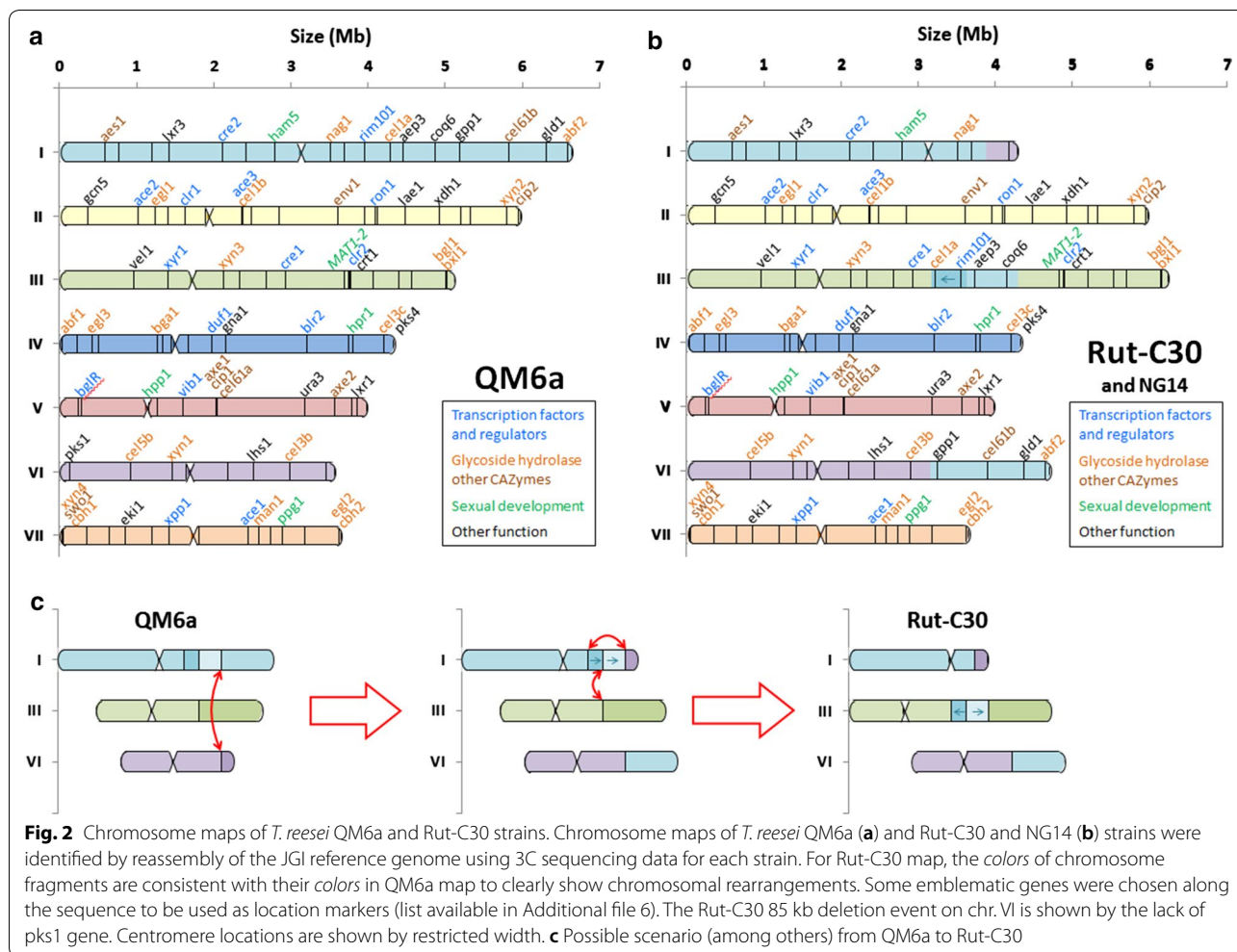
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with another fragment of scaffold 4, scaffold 49 and a small fragment of scaffold 48) to be inserted head to foot in the middle of the chromosome III (1.13 Mb and 310 genes in total for both fragments).”

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1. Jourdiere E, Baudry L, Poggi-Parodi D, Vicq Y, Koszul R, Margeot A, Marbouty M, Bidard F. Proximity ligation scaffolding and comparison of two *Trichoderma reesei* strains genomes. *Biotechnol Biofuels*. 2017;10:151. <https://doi.org/10.1186/s13068-017-0837-6>.