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Development of a system for integrative and stable transformation of the zygomycete *Rhizopus oryzae* by *Agrobacterium*-mediated DNA transfer

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Owing to a technical error, the legend to Fig. 1 was omitted. The figure and the legend now appear below.

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Fig. 1 A–C Analysis of transformants by Southern hybridization. **A** Southern analysis of [pRNU54] transformants. Chromosomal DNA was digested with *Eco*RI and probed with pRNU54. WT, undigested chromosomal DNA of COM1247. The arrowheads indicate the expected fragments. **B** Southern analysis of [pRNU54-fragment] transformants. Chromosomal DNA was digested with *Eco*RI and probed with the 5.6-kb *Sph*I fragment of pRNU54. Transformants obtained after introduction of the linear 5.6-kb *Sph*I fragment of pRNU54, corresponding to the *pyr4* marker, are designated as lin. Transformants obtained after introduction of the *Sph*I fragment pre-treated with DNA-ligase are designated as lig. The arrowheads indicate the expected fragments. **C** Southern analysis of [pRN1A *pyr4*] transformants ([pRN1A *pyr4*]#4A and [pRN1A *pyr4*]#9A), which were able to grow on minimal medium lacking uridine and on acetamide as the sole nitrogen source. Chromosomal DNA was digested with *Bam*HI and probed with pRN1A *pyr4*. The arrowheads indicate the expected fragments