ERRATUM

Erratum to: Identification and characterization of *EYK1*, a key gene for erythritol catabolism in *Yarrowia lipolytica*

F. Carly^{1,2} · H. Gamboa-Melendez² · M. Vandermies³ · C. Damblon⁴ · J. M. Nicaud² · P. Fickers³

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The original article was updated because several pieces of information were omitted. The following paragraph has been added in section Material and methods:

Sequences analysis

Genome sequences of *Yarrowia* species were assembled and annotated by Cécile Neuveglise, Hugo Devillers and coworkers (to be published). Homologs of *EYK1* in *Yarrowia* species were identified by Blast on the private site of GRYC (Genome Resources for Yeast Chromosomes;

<http://gryc.inra.fr>) using *EYK1* gene as template and retrieved using the download functionality developed by H. Devillers.

The legend of Fig. S1 should be read as follows:

Multiple alignments of *YALY0F01606g* genes with the *DAK1*, *DAK2* and *DAK3* in the *lipolytica* clade. Homologs of *EYK1* in *Yarrowia* species were identified by Blast on the private site of GRYC using *EYK1* gene as template and retrieved using the download functionality. Color code: red, similar in all gene; green, similar in two of the three genes; yellow highlighted, different in *DAK3* group compared to *DAK1* and *DAK2* gene. Sequences are from strains YALI: *Yarrowia lipolytica* CLIB122; YAGA: *Yarrowia galli* CBS 9722 (96.77%); YAYA: *Yarrowia yakushimensis* CBS 10253 (91.62%); YAAL: *Yarrowia alimentaria* CBS 10151 (87.22%) and YAPH: *Yarrowia phangngensis* CBS 10407 (85.01%). Maximal identities with *Yarrowia lipolytica* *EYK1* (YALI0F01606g) are indicated in bracket.

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✉ P. Fickers
pfickers@ulg.ac.be

¹ Unité de Biotechnologies et Bioprocédés, Université Libre de Bruxelles, Brussels, Belgium

² Micalis Institute, INRA, AgroParisTech, Université Paris-Saclay, 78350 Jouy-en-Josas, France

³ Microbial Processes and Interactions, TERRATEaching and Research Centre, University of Liège—Gembloux Agro-Bio Tech, Gembloux, Belgium

⁴ Laboratoire de Chimie Biologique Structurale, Département de Chimie, Université de Liège, Liège, Belgium

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