CORRECTION





Correction to: High-yield production of L-serine from glycerol by engineered *Escherichia coli*

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Correction to:

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Unfortunately, the order of the Figs. 1–4 has been positioned wrongly in the print published article. Following are the corrections: Fig. 1 of this article should be placed in the position of Fig. 2, Fig. 2 should be placed in the position of Fig. 3, Fig. 3 should be placed in the position of Fig. 4, and Fig. 4 should be placed in the position of Fig. 1. The correct figures (Figs. 1, 2, 3, 4) are given below.

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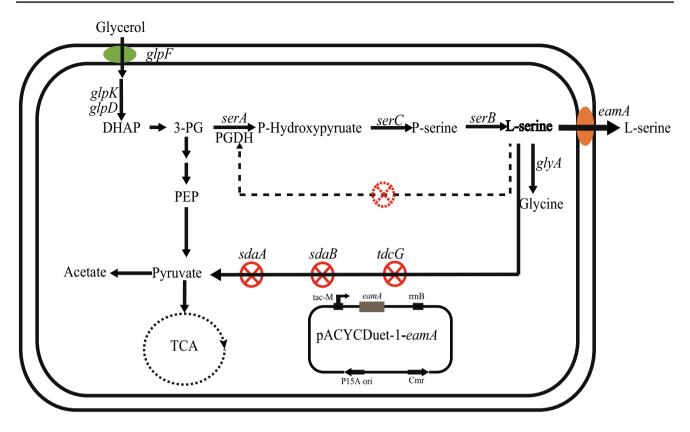
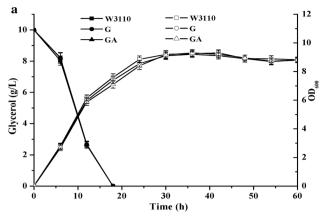


Fig. 1 L-serine biosynthesis pathways from glycerol in *E. coli*. Genes and enzymes: glycerol facilitator (*glpF*); glycerol kinase (*glpK*); glycerol 3-phosphate dehydrogenase (*glpD*); 3-phosphoglycerate dehydrogenase (*serA*); phosphoserine aminotransferase (*serC*); phosphoserine phosphatase (*serB*); cysteine/acetyl serine transporter (*eamA*); serine hydroxymethyltransferase (*glyA*); L-serine deaminases (*sdaA*, *sdaB*,

and tdcG). Intermediates: dihydroxyacetone phosphate (DHAP); phosphoenolpyruvate (PEP); tricarboxylic acid cycle (TCA). Highlighted arrows indicate pathways in which genes were upregulated by overexpression eamA. Red crosses on solid lines (\bigotimes) indicate genes that were deleted. Crosses on dashed lines (\bigotimes) indicate the removal of feedback inhibition (color figure online)





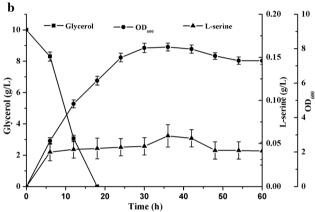


Fig. 2 Comparison of glycerol consumption, cell growth, and L-serine production of strains W3110, G, GA and GAA. a Profiles of residual glycerol and cell density with strains W3110, G, and GA. Squares represent W3110, circles represent G, triangles represent GA, open symbols represent OD₆₀₀, closed symbols represent residual glycerol. b Profiles of glycerol consumption, cell growth and L-serine production in GAA. Squares represent residual glycerol, circles represent cell growth, triangles represent L-serine. Values denote the average of three independent experiments, and error bars indicate standard deviation

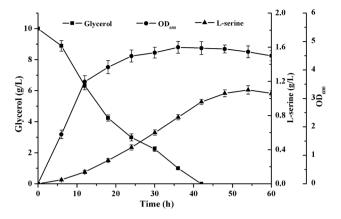


Fig. 3 Time course of L-serine production in the 4W strain during batch fermentation. Squares represent residual glycerol, circles represent cell growth, triangles represent L-serine. Values denote the average of three independent experiments, and error bars indicate standard deviation

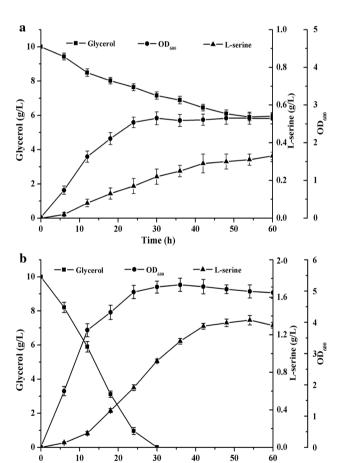


Fig. 4 Comparison of glycerol consumption, cell growth and L-serine production in strains 4W-glpD-glpK and 4WA. **a** Profiles of 4W-glpD-glpK. **b** Profiles 4WA. Squares represent residual glycerol, circles represent cell growth, triangles represent L-serine. Values denote the average of three independent experiments, and error bars indicate standard deviation

Time (h)

