

Metabolic engineering of *C. glutamicum* for L-arginine production

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L-arginine was produced by metabolically engineered *Corynebacterium glutamicum*. Random mutagenesis was first done on *C. glutamicum* to increase tolerance to L-arginine. The PPP flux was strengthened by downregulating the *pgi* gene and overexpressing the *opcA*, *pgl*, *tal*, *tkt*, and *zwf* genes. Next, the *Ncgl1221* gene encoding L-glutamate exporter was inactivated to channel L-glutamate to L-arginine formation. Also, the expression levels of the *argF* and *carAB* genes were optimized to convert L-ornithine to L-citrulline effectively. Finally, the *argGH* operon was overexpressed. Fed-batch fermentation of the final strain was performed in a 1,500 L bioreactor produced 81 g/L of L-arginine. The approaches described here will be useful in developing strains of Corynebacteria regarding the production of arginine and its derivatives. [This work was supported by the Technology Development Program to Solve Climate Changes on Systems Metabolic Engineering for Biorefineries from the Ministry of Science, ICT and Future Planning (MSIP) through the National Research Foundation (NRF) of Korea (NRF-2012M1A2A2026556 and NRF-2012M1A2A2026557)]