

Development of an amino acid-based 1-propanol production pathway via metabolic engineering in *Escherichia coli*

_____, _____, Jin hwan Park, Tae Yong Kim, _____^{*}
KAIST
(leaves4@kaist.ac.kr^{*})

1-propanol has been considered as a next generation biofuel that can be used as a gasoline substitute. Here we suggest a novel biosynthetic pathway for 1-propanol production using an amino acid overproducing *Escherichia coli* strain. The highest titer of 1-propanol (3.5 g/L), performed by Atsumi et al., has been produced in metabolically engineered *Escherichia coli* harboring *Methanococcus jannaschii* citramalate synthase (CimA), which converts pyruvate directly into 2-ketobutyrate. The previously reported L-threonine overproducing *E. coli* TH20 strain was used in this study. Deletion of competing pathways and *in silico* flux response analysis for the carbon source optimization successfully generated an *E. coli* strain producing 1-propanol. Additional metabolic engineering of the resulting strain further improved the titer. [This work was supported by the Advanced Biomass R&D Center of Korea (NRF-2010-0029799) through the Global Frontier Research Program of the Ministry of Education, Science and Technology (MEST). Further supports by BioFuelChem, EEWS program of KAIST.].