

## Metabolic engineering for the production of short chain alkane

천승우, 최용준\*, 이상엽

KAIST

(leaves4@kaist.ac.kr\*)

The  $\beta$ -oxidation was blocked by *fadE* gene deletion preventing degradation of fatty acyl-CoAs, and the activity of 3-oxoacyl-ACP synthase (*FabH*), which is inhibited by unsaturated fatty acyl-ACPs, was enhanced to promote the initiation of fatty acid biosynthesis by deleting the *fadR* gene. A modified thioesterase was employed to convert short chain fatty acyl-ACPs to the corresponding FFAs, which were consequently converted to short chain alkanes by the sequential reactions of *E. coli* fatty acyl-CoA synthetase, *Clostridium acetobutylicum* fatty acyl-CoA reductase and *Arabidopsis thaliana* fatty aldehyde decarboxylase. The resulting strain produced up to 580.8 mg L<sup>-1</sup> of SCAs [This work was supported by the Advanced Biomass Research and Development Center of Korea (ABC-2010-0029799) through the Global Frontier Research Program of the Ministry of Science, ICT and Future Planning (MSIP) through the National Research Foundation (NRF). Systems metabolic engineering work was supported by the Technology Development Program to Solve Climate Changes on Systems Metabolic Engineering for Biorefineries (NRF-2012-C1AAA001-2012M1A2A2026556) by MSIP through NRF].