Precise precursor rebalancing for hexanoic acid production by fine-tuning of *atoB* expression in recombinant *Escherichia coli*

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In recombinant *E. coli* strains, hexanoic acid is generally synthesized from artificial pathways using acetyl–CoA and butyryl–CoA as precursors. Here, we demonstrate that rebalancing of metabolic fluxes at acetyl–CoA can substantially improve the titer and productivity of hexanoic acid. First, the expression of *mhact* coding ACT increased hexanoic acid production by 47% compared to that with an *E. coli* thioesterase, TesB. Next, metabolic flux was optimized at the acetyl–CoA branch point by fine–tuning the expression level of the gene for acetyl–CoA acetyltransferase (AtoB). Four different synthetic 5′–untranslated regions (5′–UTRs) were designed for *atoB* using UTR Designer to modulate the expression level of the gene. The flux–optimized variant produced 528 mg/L hexanoic acid in 36 h, and this titer represents an 8.7–fold increase compared to that of the non–optimized parental strain. Notably, the productivity of the optimized strain was 14.7 mg/L/h and this value was the highest among the recombinant *E. coli* strains when using a similar inoculum size for fermentation. Our results show that fine–tuning the expression level of atoB is critical for production of hexanoic acid.