

Co-production of hydrogen and ethanol of *Escherichia coli* SS1 isolate

ABSTRACT

Background: The development of a potential single culture that can co-produce hydrogen and ethanol is beneficial for industrial application. Strain improvement via molecular approach was proposed on hydrogen and ethanol co-producing bacterium, *Escherichia coli* SS1. Thus, the effect of additional copy of native hydrogenase gene *hybC* on hydrogen and ethanol co-production by *E. coli* SS1 was investigated. Results: Both *E. coli* SS1 and the recombinant *hybC* were subjected to fermentation using 10 g/L of glycerol at initial pH 7.5. Recombinant *hybC* had about 2-fold higher cell growth, 5.2-fold higher glycerol consumption rate and 3-fold higher ethanol productivity in comparison to wild-type SS1. Nevertheless, wild-type SS1 reported hydrogen yield of 0.57 mol/mol glycerol and ethanol yield of 0.88 mol/mol glycerol, which were 4- and 1.4-fold higher in comparison to recombinant *hybC*. Glucose fermentation was also conducted for comparison study. The performance of wild-type SS1 and recombinant *hybC* showed relatively similar results during glucose fermentation. Additional copy of *hybC* gene could manipulate the glycerol metabolic pathway of *E. coli* SS1 under slightly alkaline condition. Conclusions: *HybC* could improve glycerol consumption rate and ethanol productivity of *E. coli* despite lower hydrogen and ethanol yields. Higher glycerol consumption rate of recombinant *hybC* could be an advantage for bioconversion of glycerol into biofuels. This study could serve as a useful guidance for dissecting the role of hydrogenase in glycerol metabolism and future development of effective strain for biofuels production.

Keyword: Hydrogen; Ethanol; Co-production; *Escherichia coli*; Glycerol