

In silico improvement of organisms using flux sum of metabolites

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This study presents an *in silico* method for improving strain performance on the basis of flux sum (Φ) of metabolites. The flux sum method first screens key metabolites that eventually increase production yield of useful substance by defining the metabolite utilization of an organism as flux sum with respect to the production of the target substance and perturbing the flux sum. Then, the method deletes and/or amplifies genes associated with aforementioned screened key metabolites.

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