

Multiplex Genome Engineering of Polyploid Industrial Yeast Strains using an Optimized CRISPR/Cas9 System

Background/objective

- ❖ Compared with lab strains, industrial yeast strains have the advantages of higher productivity and robustness under harsh industrial conditions.
- ❖ Although CRISPR/Cas9 has been widely used for industrial yeast strain engineering, its application in industrial yeast strains is less successful (15-60% efficiency for single gene deletion in ATCC 4124).

Approach

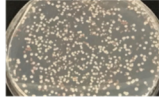





- ❖ Increasing gRNA abundance via super-high copy number plasmids.

Results

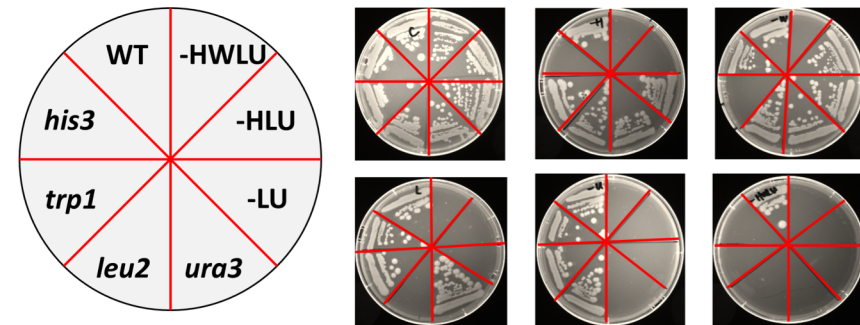
- ❖ Plasmids with higher copy numbers increased gRNA levels, resulting in higher genome editing efficiencies (up to 100% efficiency).
- ❖ Up to 12 alleles were disrupted in a single step with 100% knock-out efficiency.

Significance

- ❖ Facile engineering of industrial yeast strains for practical applications, i.e. biofuel production.

	w/ SgRNA & KO efficiency		w/ mCherry & PCN		Relative gRNA abundance
KanWT		~20%		~5/cell	1
Kan100		~60%		~20/cell	~4
Kan20		100%		~80/cell	~60

Plasmids with increased copy numbers for multiplex genome engineering of industrial yeast strains.



All the single, double, triple, and quadruple knock-out strains were constructed in a single step with 100% efficiency.

¹ Lian, et al. 2017. "Engineered CRISPR/Cas9 system for multiplex genome engineering of polyploid industrial yeast strains." *Biotechnology and Bioengineering*, DOI: 10.1002/bit.26569