

Genetic engineering toolbox for industrial yeast

Vratislav Stovicek

Strain development for diacid production
Industrial yeast development



The Novo Nordisk Foundation Center for Biosustainability,
Technical University of Denmark, Kogle Alle 6, 2970 Hørsholm



- Metabolic engineering of strains for production of wide variety of chemicals needs several rounds of genetic interventions

Industrial x laboratory yeast strains

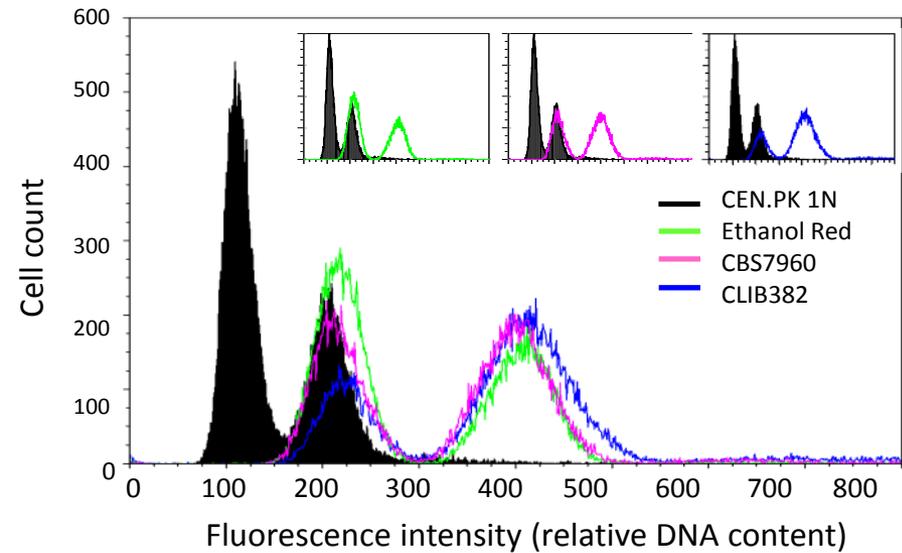
- stress resistance
- robustness
- high fermentation capacity
- multiple ploidy, aneuploidy
- prototrophic strains
- low transformation efficiency

⇒ tools for efficient genome editing

- ① gene delivery & insertion
- ② gene disruption

Industrial *S. cerevisiae* strains

CBS7960	ethanol producer from cane-sugar syrup
Ethanol Red	industrial ethanol producer
CLIB382	Irish brewer's strain
CEN.PK	laboratory strain

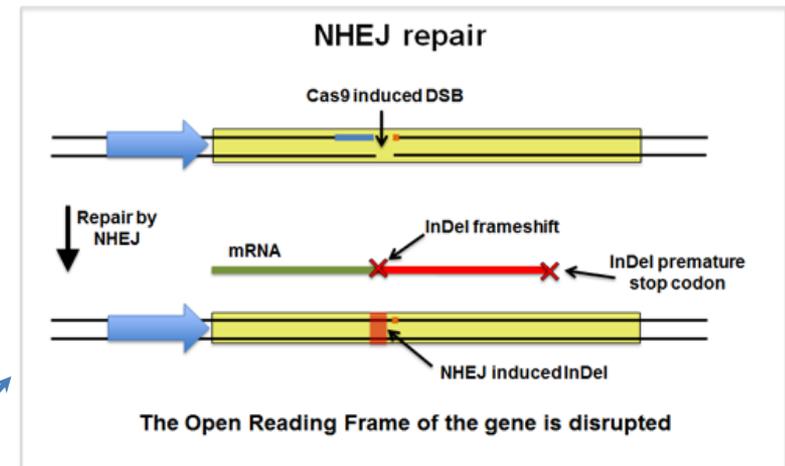
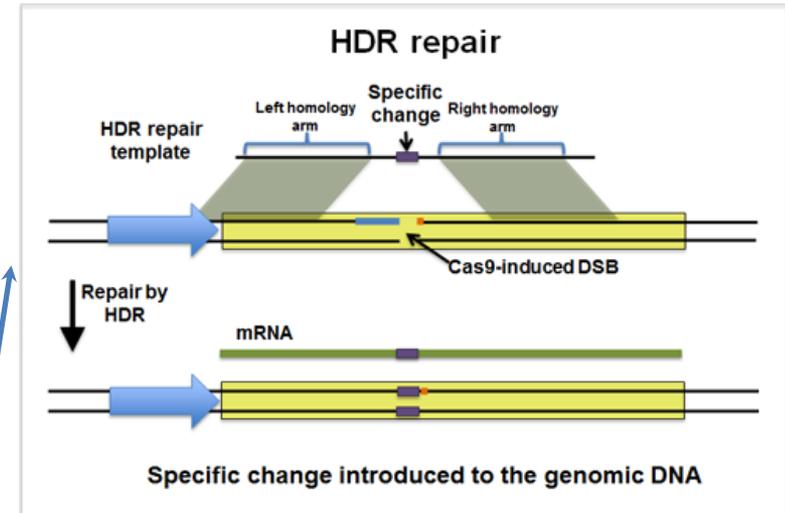
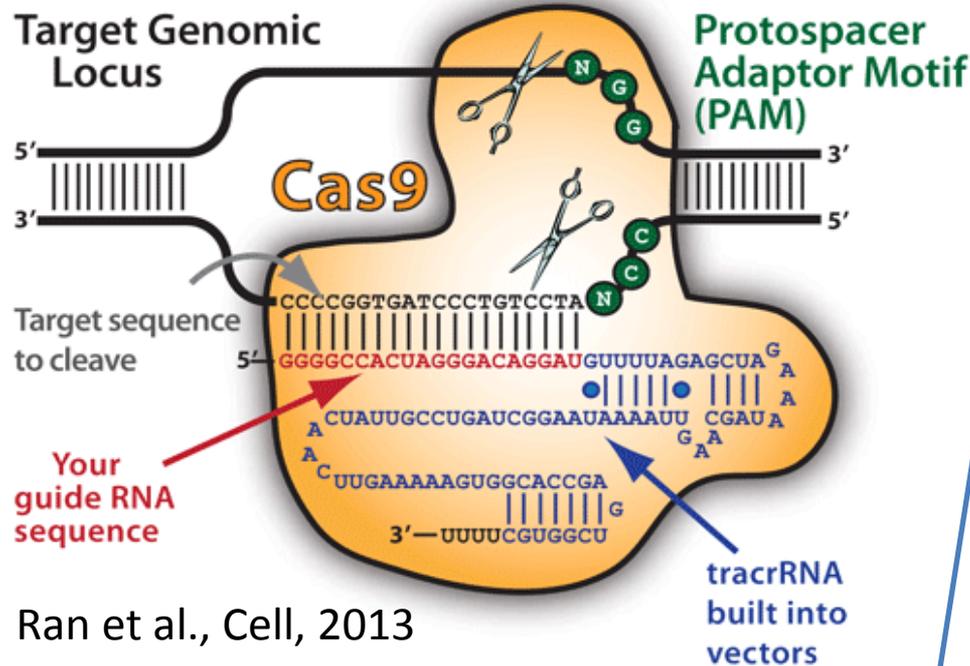


- 1) Development of genetic engineering toolbox for industrial *S. cerevisiae* strains
 - fast and efficient strategy for gene disruption in industrial strains
 - heterologous gene insertion strategy

CRISPR-Cas9 genome editing

- RNA guided specific endonuclease-mediated genome targeting

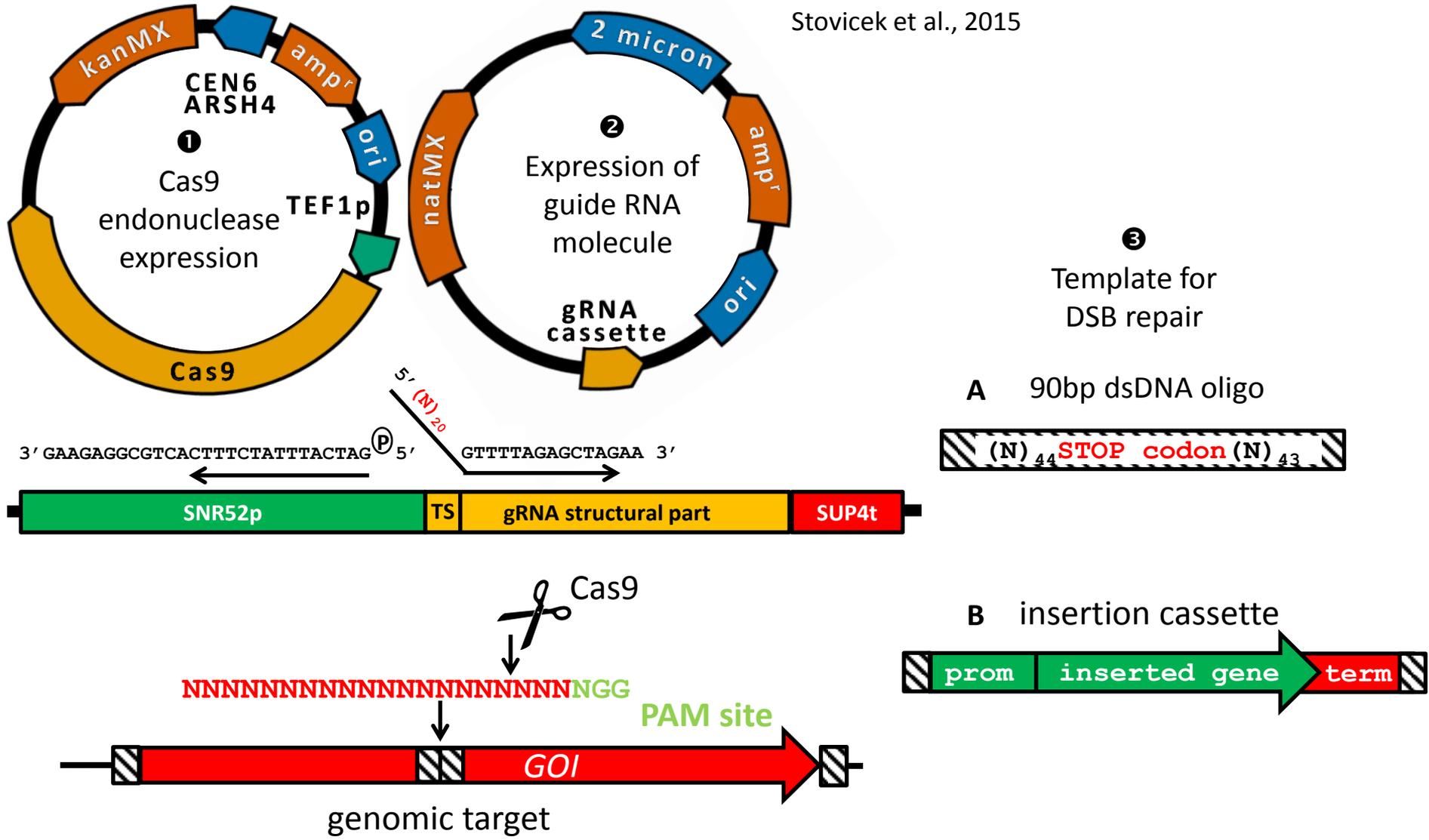
The CRISPR-Cas9 Nuclease Heterocomplex



- Cas9 introduces DSB at specific genomic location
- Cas9 can be recruited to any sequence of choice

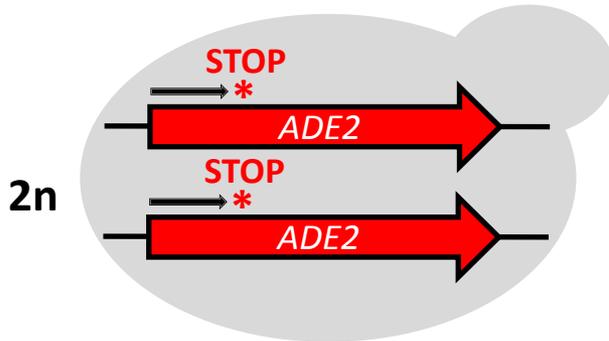
CRISPR-Cas9 system for industrial yeast gene targeting

Stovicek et al., 2015

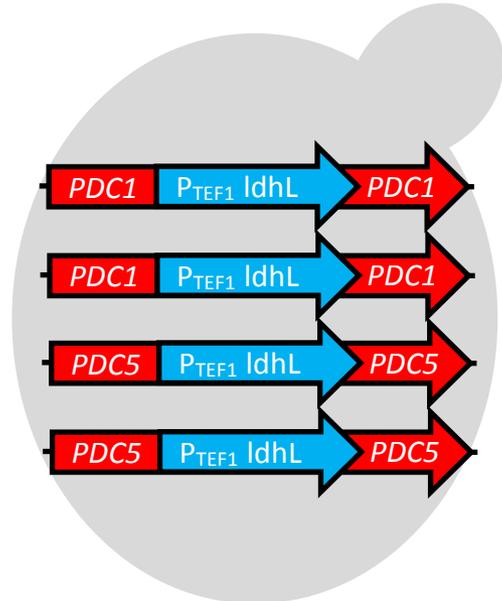


Proof-of-concept & case study

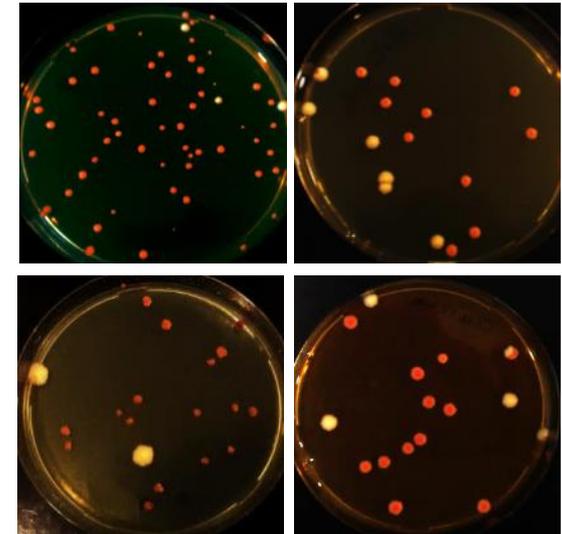
ADE2 disruption



Lactate producing strain

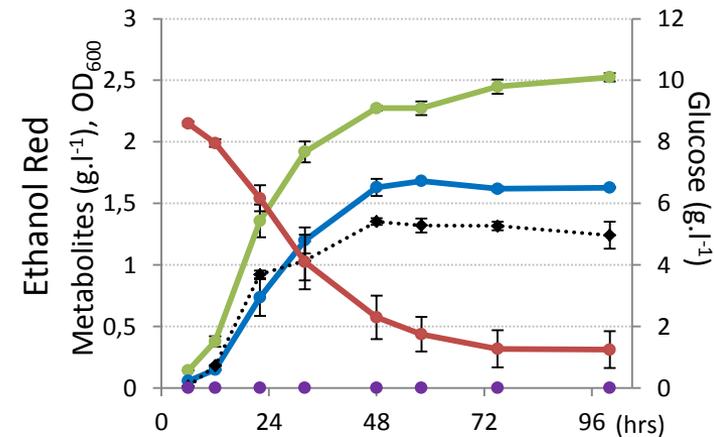
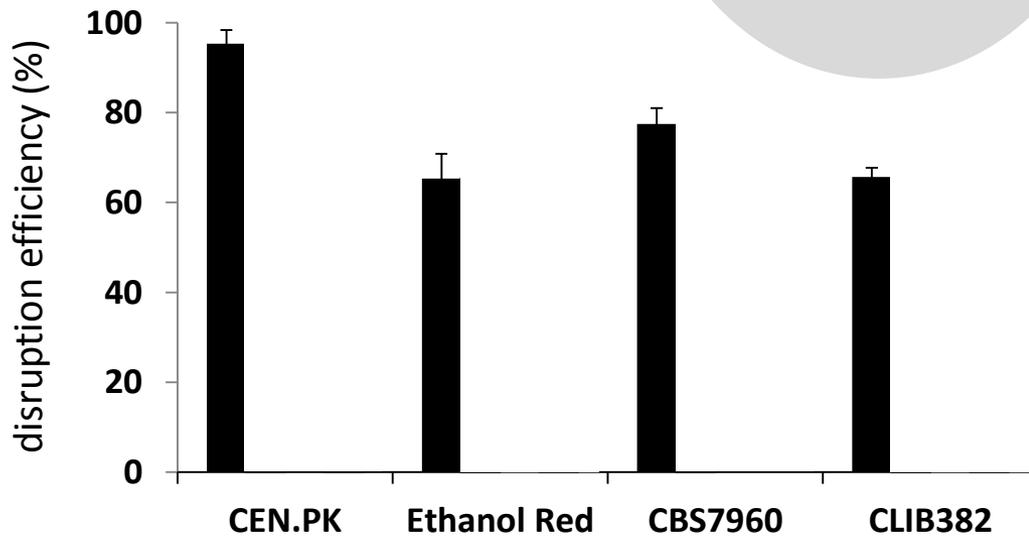


CEN.PK113-7D Ethanol Red



CBS7960

CLIB382



- **Highly efficient marker-free system for gene disruption in industrial yeast based on CRISPR-Cas9**
 - **demonstrated in metabolic engineering test case** ✓

Stovicek V., Borodina I., Förster J. (2015): CRISPR-Cas system enables fast and simple genome editing of industrial *Saccharomyces cerevisiae* strains. *Metabolic Engineering Communications* 2:13-22

- **New generation of integrative vectors suitable for gene integrations in industrial strains**
 - **demonstrated in metabolic engineering test case** ✓

Acknowledgements

Jochen Förster Irina Borodina Xiao Chen

Laura Dato

Tomas Strucko

Michael Krogh Jensen

Tadas Jakociunas

Mathew Malcolm Jessop Fabre

Gheorghe M. Borja

Mette Kristensen

Anders Holmgaard Hansen



EU FP7

grant no: 613771

novo nordisk fonden

Thank you for your attention !!!

vrast@biosustain.dtu.dk

