

# YKO parental strains

Cat. #YSC1048, YSC1049, YSC1050, YSC1061, YSC1062

A nearly complete set of yeast open reading frame (ORF) knock-outs has been produced by the Saccharomyces Genome Deletion Project (SGDP) using five Yeast Knockout (YKO) Parental Strains derived from Saccharomyces cerevisiae S288C¹. A PCR-based strategy was used to replace each ORF with a KanMX cassette containing unique tags, "barcodes", for each deletion. Four different mutant collections have been generated: haploids of mating types MATa and MATalpha, homozygous diploids for non-essential genes, and heterozygous diploids, which contain the essential and non essential ORFs.

## **Storage**

- 4 °C for up to one week
- -80 °C indefinitely

## **Product description**

Culture of *S. cerevisiae* in YPD broth with an inert growth indicator + 15% glycerol.

| BY4730 | MATa leu $2\Delta0$ met $15\Delta0$ ura $3\Delta0$   |
|--------|--|
| BY4739 | MATα leu2 $\Delta$ 0 lys2 $\Delta$ 0 ura3 $\Delta$ 0 |
| BY4741 | MATa his3Δ1 leu2Δ0 met15Δ0 ura3Δ0                    |
| BY4742 | MATα his3Δ1 leu2Δ0 lys2Δ0 ura3Δ0                     |
| BY4743 | 4741/4742  |

Homozygous diploids are in the BY4743 background unless 4730/4739 is indicated.

## **Useful websites and references**

Saccharomyces Genome Deletion Project, click here.

 Winzeler, E.A., et al., Functional characterization of the Saccharomyces cerevisiae Genome by gene deletion and parallel analysis, Science, 285, 901-906, 1999.

#### If you have any questions, contact

- t +44 (0) 1223 976 000 (UK) or +1 800 235 9880 (USA); +1 303 604 9499 (USA)
- **f** + 44 (0)1223 655 581

w horizondiscovery.com/contact-us or dharmacon.horizondiscovery.com/service-and-support

Horizon Discovery, 8100 Cambridge Research Park, Waterbeach, Cambridge, CB25 9TL, United Kingdom