

## Saccharomyces cerevisiae DUALhybrid cDNA library P02108

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### Source data

Organism Saccharomyces cerevisiae  
RNA poly A+ RNA

### Construction data

Library vector pGAD-HA  
Cloned directional / Sfi I  
1st strand synthesis oligo dT / random hexamers  
5' adapter DSM4

### Quality control data

Complexity 1.0 x 10E7 independent clones  
Average insert size 1.9 kb  
Size range 0.6 - 10.0 kb  
% vectors with insert 85 %  
% inserts > 250 bp 100 %

DSM4 adapter 5' AAGCAGTGGTATCAACGCAGAGTGGCCATTACGCCGGG 3'  
oligo dT primer 5' ATTCTAGAGCCGAGGCGGCCGACATGTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTVN 3'  
random hexamer primer 5' ATTCTAGAGCCGAGGCGGCCGACATGNNNNNN 3'  
random pentadecamer primer 5' ATTCTAGAGCCGAGGCGGCCGACATGNNNNNNNNNNNNNNNN 3'

Vector sequences and maps can be found in the support section of <http://www.dualsystems.com>.

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