

## Mate & Plate™ Library - Universal Human (Normalized)

**Catalog No.**  
630480

**Lot Number**  
1712612A

### Description

This yeast two-hybrid library was constructed from human cDNA that had been previously normalized to preferentially remove abundant cDNAs derived from high-copy-number mRNAs. The normalization process incorporates a Duplex-Specific Nuclease (DSN) treatment and SMART™ technology, and increases the representation of low-copy-number transcripts in the library. This reduces the number of clones that must be screened to identify positive interactions, and facilitates the identification and characterization of novel protein-protein interactions.

A Universal human cDNA library transformed into yeast strain Y187. The library can be readily mated to a MATa GAL4 reporter strain, such as AH109 or Y2HGold (Clontech, 2008).

### Package Contents

- 5 x 1.0 ml Mate & Plate Library - Universal Human (Normalized)
- 1 x 1.0 ml Mate & Plate Library - Control (pGADT7-T in Y187)

### Storage Conditions

- Store all components at  $-70^{\circ}\text{C}$
- Do not refreeze

### Shelf Life

- 1 year from date of receipt under proper storage conditions.

### mRNA Source

- Mixture of Poly A+ RNAs from a collection of adult human tissues chosen to represent a broad range of expressed genes. Both male and female donors are represented. Modeled after our Human Universal Reference Total RNA (Cat. no. 636538).

### Cloning Vector

- pGADT7-RecAB

### Cloning Site

- Sfi I A/Sfi I B

### Priming Method

- Sfi I (dT)<sub>30</sub> primed

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## Yeast Genotype

- (Y187): MAT $\alpha$ , ura3-52, his3-200, ade2-101, trp1-901, leu2-3, 112, gal4 $\Delta$ , gal80 $\Delta$ , met $-$ , URA3 :: GAL1<sub>UAS</sub>-GAL1<sub>TATA</sub>-LacZ, MEL1

## Shipping Conditions

- Dry ice (-70°C)

## Product Documents

Documents for our products are available for download at [takarabio.com/manuals](http://takarabio.com/manuals)

The following documents apply to this product:

- Matchmaker® Gold Yeast Two-Hybrid User Manual (PT4084-1)
- pGADT7-RecAB Vector Information (PT3718-5)

## References

Franz, O., Bruchhaus, I. & Roeder, T. Verification of differential gene transcription using virtual northern blotting. *Nucleic Acids Res.* **27**, e3 (1999).

Pretransformed Mate & Plate Libraries. *Clontechiques* **24**, 26–27 (2008).

Shagin, D. A. *et al.* A novel method for SNP detection using a new duplex-specific nuclease from crab hepatopancreas. *Genome Res.* **12**, 1935–1942 (2002).

Zhulidov, P. A. *et al.* Simple cDNA normalization using kamchatka crab duplex-specific nuclease. *Nucleic Acids Res.* **32**, e37 (2004).

## Quality Control Data

### 1. Quality Control Data

| Test                         | Result                  |
|------------------------------|-------------------------|
| Titer (yeast colonies)       | $>5 \times 10^7$ cfu/ml |
| Number of independent clones | $4.0 \times 10^6$       |
| Average cDNA size            | 1.34 kb                 |
| cDNA size range*             | 0.5 - 2.0 kb            |

\*the cDNA was size-selected by excision from an agarose gel prior to cloning

### 2. Quality Control Data for the Pretransformed Library in Yeast

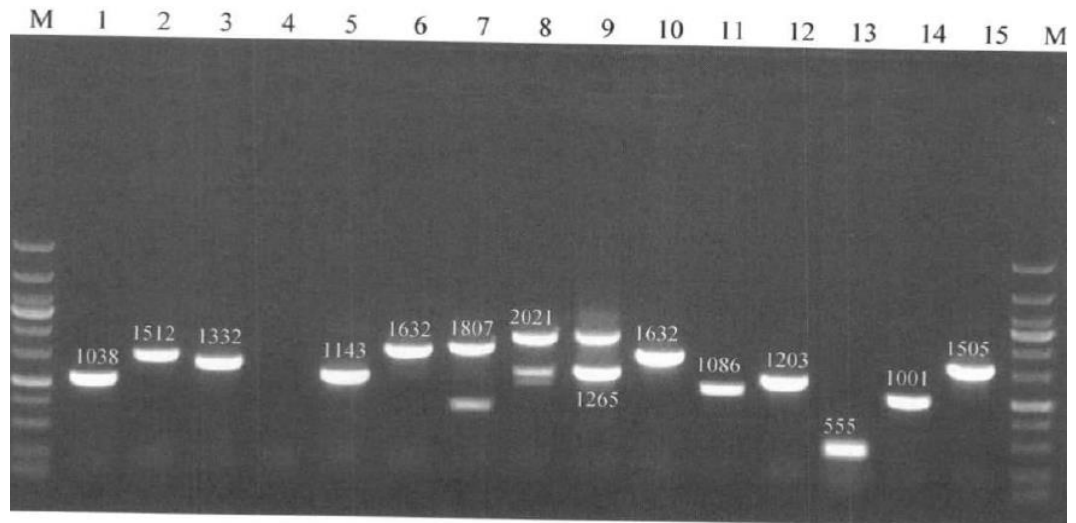
#### Library Insert Size Screening

15 yeast colonies were randomly picked and screened by PCR using the Matchmaker AD LD-Insert Screening Amplimer Set (Cat. No. 630433).

14 of 15 colonies contained inserts as determined by PCR.

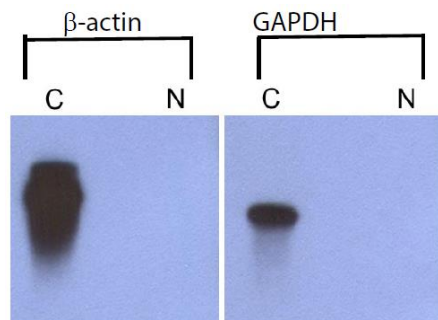
Lane M. 1 kb DNA ladder

- 1. 1.04 kb
- 2. 1.51 kb
- 3. 1.33 kb
- 4. ----
- 5. 1.14 kb
- 6. 1.63 kb
- 7. 1.81 kb
- 8. 2.02 kb
- 9. 1.27 kb
- 10. 1.63 kb
- 11. 1.09 kb
- 12. 1.20 kb
- 13. 0.56 kb
- 14. 1.00 kb
- 15. 1.51 kb



### 3. cDNA Normalization

cDNA generated using SMART™ technology was normalized using Duplex-Specific Nuclease (DSN) normalization (2, 3). Prior to cloning, the efficiency of normalization was assessed by virtual Northern blot analysis (4) comparing the abundance of GAPDH and β-actin in normalized and non-normalized human cDNA.



**Figure 1. DSN-Normalization reduces the amount of highly abundant transcripts.** Normalized (Lanes N) and non-normalized (Lanes C) Human Universal cDNA samples (PCR products) were electrophoresed on a 1.5% agarose gel and transferred to Hybond-N membrane. PCR-amplified probes of GAPDH and β-actin were labeled with <sup>32</sup>P-dATP and hybridized to the membrane. GenBank Accession numbers: GAPDH, NM\_002046 and β-actin, NM\_001101.

It is certified that this product meets the above specifications, as reviewed and approved by the Quality Department.

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