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PRODUCT: Human Aorta Matchmaker™ cDNA Library

CATALOG No. 638813

LOT NUMBER: 8030090

STORAGE CONDITIONS

- Store library at -70°C.
- Avoid repeated freeze/thaw cycles.

SHELF LIFE

1 year from date of receipt under proper storage conditions

SHIPPING CONDITIONS

Dry ice (-70°C)

mRNA SOURCE

Normal, whole aortas pooled from 38 male/female Caucasians, ages 16–73; cause of death: trauma

NOTE: No further information on the mRNA source was made available to Clontech.

PACKAGE CONTENTS

- 2 x 1.0 ml library culture (in *E. coli* BNN132)
- Saccharomyces cerevisiae strain AH109
- Saccharomyces cerevisiae strain CG-1945
- Matchmaker[™] Two-Hybrid System 3 User Manual (PT3247-1)
- Clontech Yeast Protocols Handbook (PT3024-1)
- pACT2 Vector Information Packet (PT3022-5)
- *TITER: ≥10⁸ cfu (of BNN132)/ml in LB broth & 25% sterile glycerol

CLONING VECTOR: pACT2

CLONING SITE: Xho I/EcoR I

ADAPTOR 5' AATTCGCGGCCGCGTCGAC3'
SEQUENCE: 3' GCGCCGGCGCAGCTG5'

PRIMING METHOD: Xho I-(dT)₁₈ primed

FOR RESEARCH USE ONLY

QUALITY CONTROL DATA

1. Library Information

A. Estimated % of Colonies

with Inserts: 93%

B. Number of

Independent Clones: 3.5 x 10⁶
C. Average cDNA Size: 1.7 kb
D. cDNA Size Range: 0.4 – 3.6 kb

(The average cDNA size and size range were determined by examination of an autoradiogram of the cDNA prior to cloning.)

- E. Amplification: This library was amplified once in BNN132.
- This pACT2 library was released from a λACT2 library.
- * Please retiter the library before use. For library amplification, plate out the library to obtain isolated colonies on each plate (approximately 10⁴–10⁵ colonies per plate). Plating at a higher density may cause loss of library representation.

APPROVED BY:

(PA0X749)



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QUALITY CONTROL DATA continued

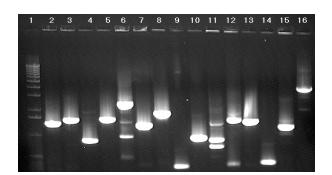
2. Insert Size Range Analysis

15 colonies were randomly picked and subjected to long-distance PCR using Clontech's Matchmaker $^{\text{TM}}$ AD LD-Insert Screening Amplimers (Cat. No. 630433) and Advantage cDNA PCR Kit (Cat. No. 639101) in a Perkin-Elmer DNAThermal Cycler.

14 of the 15 colonies contained inserts as determined by PCR.

Lane: 1. 1-kb DNA ladder

- 2. 1.5 kb
- 3. 1.6 kb
- 4. 0.9 kb
- 5. 1.6 kb
- 6. 2.6 kb
- 7. 1.4 kb
- 8. 2.0 kb
- 9. no insert
- 10. 1.0 kb
- 11. 0.9 kb
- 12. 1.6 kb
- 13. 1.5 kb
- 14. 0.4 kb
- 15. 1.3 kb
- 16. 3.5 kb



3. Coprecipitant Usage

This library was constructed using yeast tRNA as a coprecipitant.

☐ yes 🕅 no

The library was subjected to PCR screening using yeast-specific 5S rRNA primers. No amplified product was observed.

4. Analysis of Sequence Representation

Sequence representation is evaluted by colony or plaque hybridization using a gene-specific probe. Mammalian libraries are screened with a human β -actin probe which cross-reacts with all mammalian β -actin cDNA. All human cDNA libraries must show a minimum β -actin frequency of 0.10%, and all other mammalian cDNA Libraries must show minimum β -actin frequency of 0.05%. Nonmammalian cDNA libraries are screened with a ubiquitously expressed species-specific probe.

Note: The frequency of β -actin positive clones varies among libraries made with RNA from different tissues and species. A frequency of >0.10% in a human cDNA library suggests a reasonably high probability of finding a rare transcript [Hagen, F. S., et al. (1988) BioTechniques **6**:340–345.] For nonhuman mammalian cDNA libraries, a frequency of 0.05% suggests a reasonably high probability of finding a rare message in those libraries (Clontech observation, unpublished).

The percentage of colonies or plaques to which β -actin hybridized: 0.46%

Note: The β -actin probe may hybridize to other forms of actin.

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QUALITY CONTROL DATA continued

5. PCR Screening for Specific Sequences

1 μ I of the library lysate was screened by PCR using PCR primers for one or more of the following gene transcripts: β -actin, G3PDH, or transferrin receptor. β -actin and G3PDH represent high-abundance gene transcripts whereas transferrin receptor represents a low-abundance gene transcript. Furthermore, the β -actin primers are designed to amplify a 1.1-kb fragment near the 5' end and thus provide a stringent test for detecting the presence of full-length transcripts. (Note: Species-specific primer sets are used for human, mouse, and rat cDNA libraries; human primer sets are used for other mammalian cDNA libraries.)

The presence of the following 3 sequences were determined using PCR.

β-actin

G3PDH

transferrin receptor

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