

1 가 .
 2 H - pYFRH - 1
 3 SDS - PAGE
 4 L - pYFRL - 1

ase chain reaction) , (cDNA library) (polymer

(iron)
 H - L - , 1 500kDa
 4,500 1,200
 (band)
 21kDa Heavy (H) - 19kDa Light(L) - 1 24
 24 H - L - (subunit) 가

H - Fe() Fe() , Fe() , (core)
 (ferroxidase) Fe() L - . L - H - 55% . H -

H - (Costanzo, F.,et al., EMBO J. 3:23 - 27, 1984)

(Levi, S.,et al., Gene 51:269 - 274. 1987)

PL , 15% H - 가 L -
 H - H - , H -

(Prozzi D.,et al., FEMS letter 234 : 61 - 4, 1988)

H - H - H -

L - (Santoro, C.,et al., Nucl. Acids. Res. 14:2863 - 2867. 1986)
(Levi, S.,et al., Biochemistry 28:5179 - 5184. 1989)

15%

L - 가 L -

(Levi S.,et al., J. Mol. Biol. 238:649 - 654, 1994)

H - L

- 가 ,
L - 가 ,
L - 가

H -
H -
H -

3 4 가

(*Saccharomyces cerevisiae*)
가

가 (pyrogen) 가

24 24 H - L - 가
(subunit)

(folding)

가 가

가

(*S. cerevisiae*)

GAPD

(*Saccharomyces cerevisiae*),
(*Pichia pastoris*) H - L -

(*Hansenula polym*

orpha)
가

가

H -

L -

H - L -

dNTP Mix 0.5mM Klenow 37 , 15min ,EcoRI 가 XhoI (blunt end) Klenow 1μl 70 20min
 GAL ,EcoRI 37 , 15min Xbal Klenow 1μl 70 20min GAPDH dNTP Mix 0.5mM Klenow
 ,EcoRI 가 (Ligase) H - E. coliXL1 - Blue 70 100
 20min 가 (2). DNA

2: H -
 1 pYFRH - 1 E. coliXL1 - Blue 5μg DNA
 30 YPD(YE 1%, Peptone 2%, Glucose 2%)
 Y2805WT(S. cerevisiaeY2805WT) 1/100 volume YPD broth 3 4
 4 2 , 4 1M (Sorbitol) 1
 1M 70μl volume DNA Electroporater (B
 io - rad Gene Pulser II) 2mm 가 (Cuvette) 200 , 250V, 1,5 μ F (sh
 ock) 1mL YPD (tube) 30
 0 , 가 (w/o Nitrogen base 6.7g/L, 20mg/L Histidine, Agar 1.5%) 200μl 3
 2 3 Y2805/pYFRH - 1

Y2805/ pYFRH - 1 2000 7 21

KCTC 0832BP

3: H -
 H - 가
 Y2805/pYFRH - 1(KCTC 0832BP) 30 URA (w/o Nitrogenbase 0.67
 %, Glucose 2%, Histidine 20μg/ml) O.D. 0.5(at 660nm) 50 YPD
 (Yeast extract 2%, Peptone2%, Glucose 2%) broth OD 660 6 7
 (20) SDS가
 1/2 (v/v) 5 (Marker) 15% SDS - PAGE (15% Tris -
 Glycine gel, Novex) (Silver) (3).

4: L - pYFRL - 1
 L - cDNA
 L - pBlueScript KS +
 가 L -

5' - ATTGAATTCACGATGAGCTCCAGATTCGTCAG - 3'

5' - AATTCTCGAGGTGGGCTCAAGAAGGCTCTTAGTC - 3'

dTTP 200 μ M. primer 100 μ l 1 μ l cDNA library (Clontech), dATP, dCTP, dGTP,
 200nM, Vent polymerase(New England Biolab) 1 Unit,
 10 μ l 50 μ l 95 5 , 95 1
 , 65 1 , 72 2 30 DNA , 72 10
 . 1.2% 가 (TBE) DNA (1)
 PCR purification kit (Promega) DNA EcoRI XhoI
 pBlueScirpt KS + E. coliXL1 - Blue IPTG(Isopropyl
 - D - thiogalactopyranoside) 1mM, X - gal(5 - Bromo - 4 - Chloro - 3 - Indolyl - D - galactopyranoside) 40 μ g/
 mL 100 μ g/mL 가 LB(Yeast Extract 0.5%, Tryptone 1%, NaCl 0.5%) 가 (Agar)
 37
 EcoRI XhoI 1.2% 가 DNA DNA가 L -
 가 XhoI Klenow 1 μ l
 dNTP Mix 0.5mM 37 , 15min (blunt end) 70 20m
 in Klenow ,EcoRI GAPDH
 GAL Xbal Klenow 1 μ l dNTP Mix 0.5mM
 37 , 15min 70 20min Klenow
 ,EcoRI L - 70
 20min 가 (Ligase) E. coliXL1 - Blue 100
 μ g/mL 1 DNA
 pYFRL - 1 (4).
 5: L -
 4 pYFRL - 1 E. coliXL1 - Blue 5 μ g DNA
 , 30 YPD(YE 1%, Peptone 2%, Glucose 2%)
 Y2805WT(S. cerevisiaeY2805WT) 1/100 volume YPD broth 3 4
 4 2 , 4 1M (Sorbitol) 1
 . 1M 70 μ l volume DNA Electroporater (B
 io - rad Gene Pulser II) 2mm 가 (Cuvette) 200 , 250V, 1,5 μ F (sh
 ock) 1ml YPD (tube) 30
 , 가 (w/o Nitrogen base 6.7g/L, 20mg/L Histidine, Agar 1.5%) 200 μ l 30
 2 3
 Y2805/pYFRL - 1
 Y2805/ pYFRL - 1 2000 7 21
 KCTC 0833BP
 6: L -
 L - 가
 Y2805/pYFRL - 1(KCTC 0833BP) 30 URA (w/o Nitrogenbase 0.67
 %, Glucose 2%, Histidine 20 μ g/Ml) O.D. 0.5(at 660nm) 50 YPD
 (Yeast extract 2%, Peptone2%, Glucose 2%) broth . OD 660 6 7
 (20) SDS가
 1/2 (v/v) 5 (Marker) 15% SDS - PAGE (15% Tris -
 Glycine gel, Novex) (Silver) (3).

가 H- L- 가 H- L

(57)

1.

H- DNA가

2.

1 H- DNA

3.

1

4.

3 (Hansenula polymorpha) (Saccharomyces cerevisiae), (Pichia pastoris)

5.

3 H- H-

6.

5 H-

7.

6 H-

8.

L- DNA가

9.

8 , L - DNA , , ,

10.

8 .

11.

10 , (Saccharomyces cerevisiae), (Hansen
ula polymorpha) (Pichia pastoris) .

12.

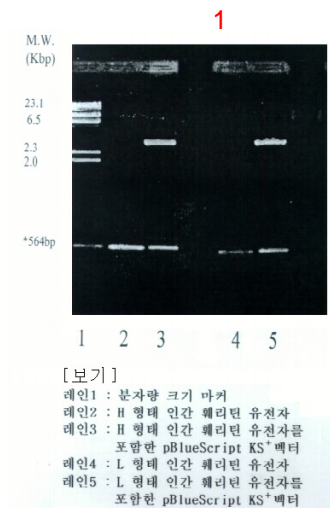
10 L - L -

13.

12 L - .

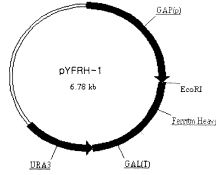
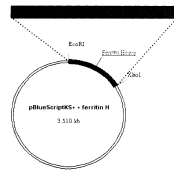
14.

13 L - .



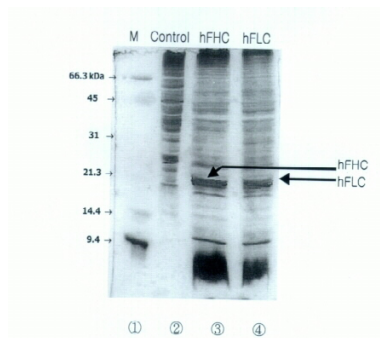
2

5'-ATTGAATTCACGATGACGACCGGTCACCTCG-3'
 3'-AATTCGAGCAGGAAGTACCCACGGCAATTG-5'



S. cerevisiae

3

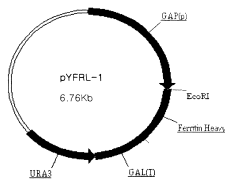
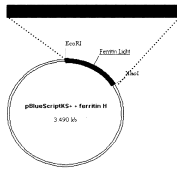


[보기]

- 라인1 : 분자량 크기 마커
- 라인2 : 웨리틴 발현백터로 형질전환되지 않은 효모 대조구
- 라인3 : H 형태 인간 웨리틴 발현백터(pYFRH-1)로 형질전환된 재조합 효모
- 라인4 : L 형태 인간 웨리틴 발현백터(pYFRL-1)로 형질전환된 재조합 효모

4

5'-ATTGAATTCACGATGAGCTCCAGATTCGTCAG-3'
 3'-AATTCGAGGTGGGCTCAAGAGGGCTCTTAGTC-5'



S. cerevisiae
 - 9 -