

백서의 실험적 급성 심근경색후 순환 및 국소 Angiotensinogen mRNA 발현의 변화

김 효 수

Change of Angiotensinogen mRNA Expression in Myocardium and Liver after Myocardial Infarction in Rat

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ABSTRACT

Background : The renin-angiotensin system (RAS) plays a crucial role in pathophysiology of congestive heart failure and ventricular remodeling after myocardial infarction (MI). There are two components, systemic and local, in RAS. There has not been a study to analyze differentially the sequential changes of systemic and local RAS after MI. The aim of this study was to analyze the sequential change of the expression of angiotensinogen mRNA, the first component of the renin-angiotensin system, in liver and non-infarcted myocardium in rats after myocardial infarction. **Methods :** Female Sprague-Dawley rats (body weight 200–250 g) were subjected either to left coronary artery occlusion or to sham operation. And the rats were sacrificed at 1, 4, 18, 24 hours, 3 days, 2, 3 weeks. Hemodynamic measurement was performed and RNA was extracted from various tissues including liver and ventricle for the analysis of the expression of the angiotensinogen mRNA by northern blot analysis or RT-PCR. **Results :** Coronary artery ligation resulted in comparable infarct sizes among rats at 3 weeks after MI and was accompanied by significant increases of LVEDP (preMI 11 ± 2 vs postMI 21 ± 3 mmHg, $n = 4$). Systolic arterial pressure was reduced in animals with infarction (preMI 130 ± 15 vs postMI 90 ± 10 mmHg, $n = 4$). The liver angiotensinogen mRNA levels increased at 4, 18, 24 hours after myocardial infarction and decreased at 3rd day to control values (Angiotensinogen/GAPDH ; preMI 1.35 ± 0.20 vs postMI 5.97 ± 0.25 , max 4-fold, $n = 3$). After sham operation, the liver angiotensinogen mRNA levels increased also at 4, 18, 24 hours, but in a less degree (Angiotensinogen/GAPDH ; preop. 2.15 ± 1.17 vs postop. 3.41 ± 1.76 , max 1.5-fold, $n = 3$). In contrast to the liver, small amounts of angiotensinogen mRNA were detectable in normal left ventricle of rat with RT-PCR. The myocardial angiotensinogen mRNA levels decreased transiently in acute phase after MI, and recovered at 3-day after MI and increased further afterwards upto 3rd month after MI. **Conclusion :** The angiotensinogen in liver was activated early during acute phase after MI and decreased toward normal as the stable state was achieved. In contrast to the circulating RAS that was activated in acute phase after MI, the local RAS in heart was activated in chronic phase after MI. (Korean Circulation J 1999;29(3):322-334)

KEY WORDS : Myocardial infarction · Remodeling · Angiotensinogen · Gene expression.

323

(Female Sprague - Dawley rat, weight 200 ~ 250 g).

심근경색의 모델 및 수술적 처치

가

가 가 .

3

1) .

2) 30 atropine 0.05 mg/kg
pentobarbital 25 mg/kg

3) supine position
rodent ventilator (Harvard apparatus)

4

cDNA northern blotting probe

probe

肝 心筋

curved needle 6 0 silk

northern blot analysis RTPCR

4) (1 , 4 , 18 ,
24 , 3 , 7 , 2 , 3)

mRNA .

(- 70 liquid
nitrogen) - 70

대상 및 방법

연구 디자인

5) control(0) ,
sham control

Table 1

1 , 4 , 18 , , 3 , 2 , 3 , 3

肝 心臟

mRNA

조직으로부터 RNA 추출

1) - 70 RNAzol - B
(CINNA/BIOTECX Lab, Inc. : Guanidine thio-
cyanate, 2 - Mercaptoethanol, Phenol)

RNA . - 70

100 mg RNAzol 2 ml
(homogenizer) 30 3

실험동물

가

Table 1. 연구 디자인(각군의 각 시간대의 샘플 수는 4마리임.)

0 hour	1st	4th	18th	24th	3rd	2nd	3rd	3rd
	hour	hour	hour	hour	day	week	week	month
N = 4 AMI	N = 4	N = 4	N = 4	N = 4	N = 4	N = 4	N = 4	N = 4
N = 4 sham op.	N = 4	N = 4	N = 4	N = 4	N = 4	N = 4	N = 4	N = 4

sample 1/10 chloro - 15 g, LB medium 1l, ampicillin 100mg)
 form 15 colony LB media broth
 15000 rpm 15 4 (during overnight with 37 shaking incubator).
 isopropanol 가 - 20 45 3) E.coli Quigen plasmid preparation
 15000 rpm 15 kit (lysis) plasmid
 RNA 75% ethanol (restriction enzyme ; BamH1, Hind
 12000 rpm 8 RNA , Boehringer Mannheim) 37 12
 RNA 0.5% SDS(sodium dodecy - 4) probe re -
 Isulfate, pH 7.2) suspension combinant plasmid Genomed
 - 70 - 20 Jetsorb gel extraction kit gel
 2) RNA 260 nm spe - TE buffer(10 mM
 ctrophotometer , Tris/HCl, 1 mM EDTA, pH 8.0) suspension
 0.8% agarose gel(ethidium bromide - 20
 stained) UV transilluminator(UVP)
 RNA (degradation)
 18S 28S band

안지오텐신노젠 유전자 probe의 생산

Angiotensinogen probe 생산

1) probe ,
 cDNA ,²⁷⁾
 BamH1 Hind
 가 primer
 : sense promoter
 5/- ACTGCCGGATCCCCCGGGCTG - 3/(+501 -
 +522nt), +1 =
 : antisense primer
 5/- CCCGCTTCGAAGATTC - 3/(+1038 - +1052)
 poly(A) - RNA cDNA
 primer PCR ,
 DNA pBluescript SKII(-) BamH1 -
 Hind subcloning
 model 373A DNA sequencing system(Applied Bio -
 system)
 2)
 recombinant plasmid(pBluescript SK, Gen -
 bank) E.coli , LB agar plate(Bacto agar

GAPDH probe
 Internal control GAPDH(GlycerAl -
 dehyde - 3 - Phosphate DeHydrogenase) probe
 , GAPDH cDNA
 ,²⁸⁾ EcoRV, EcoRI 酵素
 가 primer
 : sense primer
 5' - GCCAAGGATATCATGACAACT - 3'
 (+474 - +495 nt), +1 =
 : antisense primer
 5' - CATCCACAGAATTCTGGGTGGCAGTGAT - 3'
 (+534 - +561 nt).

肝 poly(A) - RNA 逆轉寫
 cDNA primer PCR
 87 pBluescript SK(-)(Stra -
 tagene Inc, La Jolla, USA) EcoR1 - EcoRV
 subcloning

DNA thermal cycler 480
 (Perkin - Elmer Corp, Norwalk, Cetus Corp, Calif,
 USA) Taq dye terminator cycle sequencing kit
 (Applied Biosystems Inc, Foster city, Calif, USA)

model 373A DNA sequencing systems
 (Applied Biosystems)

Northern blot analysis

capillary transfer to membrane

1) 0.5% SDS(sodium dodecyl sulfate, pH 7.2) suspension RNA 1/10 2M Na acetate(pH 5.6) 2.5 100% ethanol - 70 20 15,000rpm 20 70% ethanol 15,000 rpm 15 RNA RNA 3 ul DEPC(diethyl pyrocarbonate) - treated water 20 ul RNA loading buffer (deionized formamide, 37% deionized formaldehyde, 10× MOPS, 50% glycerol) 37% formaldehyde 1% agarose gel(ehidium bromide stained) 18 24

2) agarose gel UV transilluminator RNA loading DEPC - treated water shaking incubator 20 20× SSPE(NaCl, NaH₂PO₄ · H₂O, EDTA, pH 7.4) 20 anarose gel paper tower 24 transfer membrane (Hybond - N+, Amersham) capillary transfer . Transfer buffer 20× SSPE

3) Transfer membrane UV transilluminator transfer UV cross - linker (UVP, model CL - 1000) 12× 104J cross - link . membrane hybrid 가 - 20

Hybridization

1) Membrane hybridization prehybrid buffer(deionized formamide, 20× SSPE, 100× Denhardt's solution, 0.5% SDS, denatured Salmom sperm DNA) 42 rolling incubator rolling 2 overnight prehybridization

2) Amersham Megaprime DNA labelling kit (RPN 1606) 32

P - labelled angiotensinogen DNA probe . Angiotensinogen DNA template 25 ng(2 ul) primer 5 ul, dH₂O 26 ul 95 100 5 denature , denatured DNA dATP, GTP, TTP가 reaction buffer 10 ul radiolabelled dCTP 5 ul DNA polymerase (Klenow enzyme) 2 ul 37 40 incubation . DNA polymerase - 20 - 70 dry ice - 20 microcon(Amicon, 30,000 dalton) 32P labelled angiotensinogen DNA probe 100 5 denature ice snap cooling isotope labelled angiotensinogen DNA probe .

3) Isotope labelled angiotensinogen DNA probe hybrid buffer(prehybrid buffer) 42 rolling incubator 20 hybridization

4) Hybridization membrane 42 50 . 2× SSPE rinse 2× SSPE, 0.1% SDS shaking 10 . 3 5 1 1.5 kcpm radioactivity film cassette - 70 film (X - OMAT, Kodak Co.) autoradiography

RT-PCR(Reverse Transcription and Polymerase Chain Reaction) and Southern blot analysis

Reverse transcription

Promega reverse transcription system . RNA 2 ug , 25 mM MgCl₂ 4 ul, 10× buffer 2 ul, 10 mM dNTP mixture 2 ul, rRNasin ribonuclease inhibitor 0.5 ul, AMV reverse transcriptase 15 units 20 ul가 RNase - free water . 42 30 , 99 5 , 0

5 reverse transcrip -
tase .

PCR of cDNA from post - infarction ventricle

mRNA primer
cDNA²⁷⁾ 2
(cDNA 718 738) 5
(1321 1301)
upstream primer
5' - CGCCATCTTCCCTCGCTCTCT - 3' (718
738 nt)
downstream primer
5' - CTGCTCCTCCTCGCCTGCTTG - 3' (1321
1301 nt)

genomic DNA

가
Fig. 1 610 가

Fig. 2 .

Reverse transcription product 20 ul 5 ul

PCR . PCR

, 30, 35, 40

denaturation at 94 , for 60 sec

annealing at 60 , for 60 sec

extension at 72 , for 120 sec

Internal control GAPDH(GlycerAl -
dehyde - 3 - Phosphate DeHydro - genase)

GAPDH

cDNA ,²⁸⁾ 가

primer .

upstream primer

5' - GCCAAGGATATCATGACAACT - 3'

(+474 - +495 nt), +1 =

downstream primer

5' - ATCCACAGAATTCTGGGTGGCAGTGAT - 3'

(+534 - +561 nt).

GAPDH 190 가

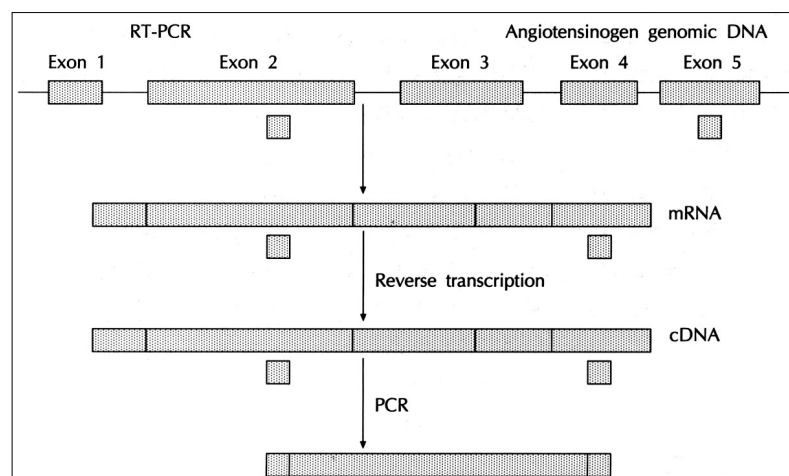


Fig. 1.

Sequences and loci of primers for PCR and PCR product on cDNA of angiotensinogen

718 738
cgccatcttccctcgctctctggacttaccactgaccagttcttctgctgccagaaatcaacaggtttgtgc
aggctgtgacaggggtggaagatgaacttgccactagaggggtcagcacggacagcacctattttcaa
cacctacgttcaactccaaggaagatgagaggcttctccagctgactgggtccatgagttctgggtgga
caacagcacctcagtgctctgctgcccagctctcgggcactggcaacttccagcactggagtgacgcccag
aacaacttctcgtgacacagcgtgccctgggtgagagtgacacctgctgctgatccagccccagtgctg
cctcagatctcgacaggggtggaggtcctcgtcttccagcacgacttctgactggataaagaacccgct
cctcgggccatcctgctgaccctgccgagctggaaattcggggattcctacaacctgcaggacctgctgg
ctcaggccaagctgtctacccctttgggtgctgaggcaaatctgggcaagatgggtgacaccaaccccg
agtgggagaggttctcaacagcatcctcctgaactccaagcaggcaggaggagcag
1300 1321

Ohkubo Proc Natl Acad Sci USA 1983

Fig. 2.

cDNA 572 1107
(Fig. 4).

Southern blot analysis of RT - PCR products

mRNA 검색후 간에서의 안지오텐신노젠 mRNA 발현의 변화 양상

cDNA Southern blotting northern blot analysis mRNA

4, 18, 24 가 (Fig. 5),

cDNA 572 mRNA/GAPDH mRNA

1107 PCR product 4 (Angiotensinogen/GAPDH ;

Fig. 3 preMI 1.35 ± 0.20 vs postMI 5.97 ± 0.25 , max 4 - fold, n = 3).

Southern blotting Amersham , Sham - operation ,

Fluorescence Gene Image System mRNA/GAPDH mRNA 가 1.5

(Angiotensinogn/GAPDH ; preop. 2.15 ± 1.17 vs postop 3.41 ± 1.76 , max 1.5 - fold, n = 3)(Fig. 6).

결 과

mRNA

심근 경색증 군에서의 혈역학적 변화 ,

3 acute phase reactant

가 ,

(preMI 11 2 vs postMI 21 3 mmHg, n = 4), , sham - op

(preMI 130 15 vs postMI 90 mRNA 가가

10 mmHg, n = 4). mRNA

acute phase response

안지오텐신노젠 탐식자의 클로닝과 염기 배열 결정 ,

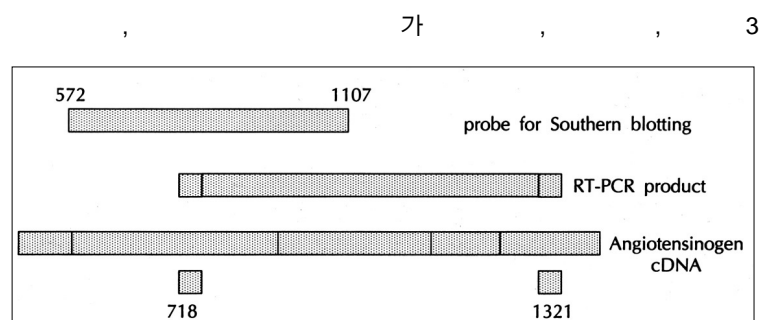


Fig. 3. Scheme of southern blotting of RT-PCR product.

```
CTGCAGGCTGTTACAGGGCTTGCTGGTCACCCAGGGTGGAAGCAGCAGTC
CAGACACCCCTGCTACAGTCCACCGTGGTGGGCCCTTCTCACTGCCCCAGGC
TTGCGCCTAAACAGCCATTGTTGAGAGCTTGGGTCCCTTACCCCCCGCC
ATCTTCCCTCGCTCTCTGGACTTATCCACTGACCCAGTTCCTTGCTGCCAG
AAAATCAACAGGTTTGTGACAGGCTGTGACAGGGTGGAAGATGAACCTTGCCA
CTAGAGGGGTCAGCACGGACAGCACCCCTATTTTCAACACCTACGTTTAC
TTCCAAGGAAGATGAGAGGCTTCTCCAGCTGACTGGGCTCCATGAGTTCT
GGGTGGACAACAGCACCTCAGTGTCTGTGCCATGCTCTCGGGCACTGGC
AACTTCCAGCACTGGAGTGAGCCCAGAACAACTTCTCCGTGACACGCGTG
CCCCTGGGTGAGAGTGTACCCCTGCTGCTGATCCAGCCCCAGTGCGCCT
CAGATCTCGACAGGGGTGGAGGTCTCTGCTCTTC
```

Fig. 4. 클로닝된 안지오텐신노젠 탐식자의 염기 배열 순서.

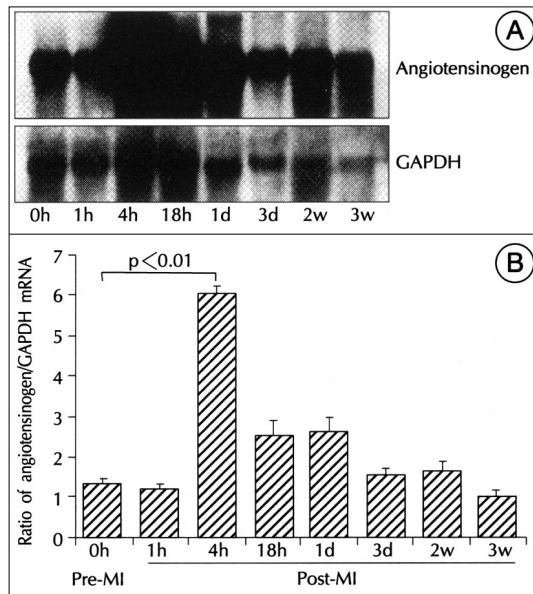


Fig. 5. Sequential change in angiotensinogen (Ao) mRNA expression (a) and the ratio of Ao/GAPDH mRNA (b) in liver after experimental MI. Ao expression remarkably increased from 4 to 24h after MI and returned to its baseline level in 3 days. The increase during the acute phase after MI was greater than that after the sham operation.

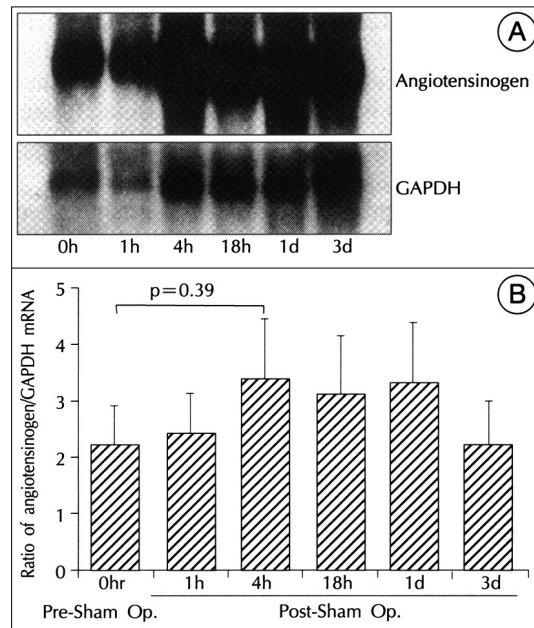


Fig. 6. Sequential change of expression of angiotensinogen (Ao) mRNA (a) and the ratio of Ao/GAPDH mRNA (b) in liver after sham operation. Ao expression increased slightly from 4 to 24h after sham operation, indicating that Ao behaves as an acute phase reactant. GAPDH, glyceral-dehyde-3-phosphate dehydrogenase : h, hours ; d, days ; w, weeks ; m, months

mRNA
가 (Fig. 7).
mRNA

심근경색후 비경색심근에서의 안지오텐신노젠 mRNA 발현의 변화 양상

mRNA Fig. 7
northern blot
analysis 가
semiquantitative RT - PCR

mRNA 가
24 가
3 3, 3 가
GAPDH mRNA

(Fig. 8).
Southern blotting
mRNA radioac -
tivity densitometry , Fig. 8
mRNA/GAPDH mRNA 比

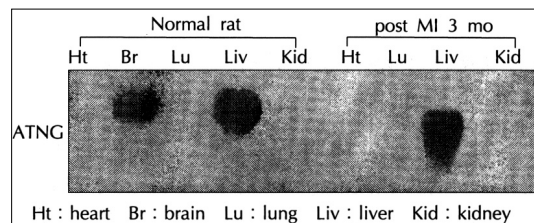


Fig. 7. Angiotensinogen mRNA expression in various tissues in normal or post M1 rats.

3 가 , 3
1.7 .
간과 심근에서의 안지오텐신노젠 mRNA 발현의 시간차

Fig. 9
mRNA
가 ,
3 ,
mRNA 가 ,
3 , 3 2

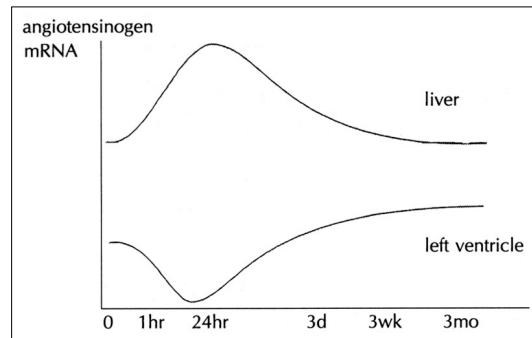
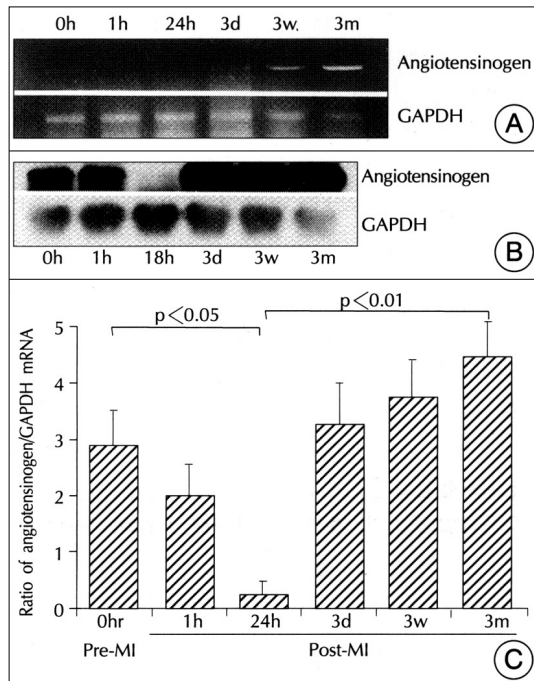


Fig. 9. Sequential change of angiotensinogen mRNA expression in liver and left ventricle after MI.

고 찰

가 mRNA 가

가 가

순환과 국소 안지오텐신노젠 mRNA 발현의 시간적 차이

mRNA

가

12

mRNA mRNA

mRNA

가 , 가 .²⁶⁾

-

가 , -

가 , -

가 .

- 가

.

, 29 - 31)

-

- 가

가 .

-

.³²⁾

경색후 간에서의 안지오텐신노젠 mRNA 발현 양상과 그 기전

, 4

mRNA ,

endotoxemia, surgery 22)23)

3 16 (SAVE study),¹⁵⁾ 3 10
(AIRE study)¹⁷⁾
가 , 24
가
(GISSI - 3, ISIS - 4)¹⁶⁾¹⁸⁾ ,
(CONSENSUS - 2).⁴⁷⁾
24
2 ,
remodeling ,⁴¹⁾ 36
1
가 ⁴²⁾ .
가
가
가 (ISIS - 4).¹⁶⁾
, ,
, 가
; Pfeffer ⁷⁾
captpril
21
2 - , -
2 , 3
가 . Gay
³⁷⁾ 3 , 1
2 captpril
3 , 2 2 . , 恒常性
4 , 3
3 4 .
가 ,
Shoemaker ³⁸⁾
1 3 captpril
, 3 5
, ()
2 14
¹²⁾¹³⁾³⁹⁾ ,

요 약

연구배경 :

mRNA
 방 법 :
 , 18 , 24 , 3 , 2 , 3 , 3 (1 , 4)
 RNA
 3
 mRNA Northern blot analysis
 RT - PCR mRNA
 결 과 :
 , 4, 8, 24
 mRNA 가 3
 ,
 가, 3
 mRNA , 3 ,
 3 가
 결 론 :
 가 ,
 mRNA 가 , 가
 , 가
 가
 가
 가
 중심 단어 : Angiotensinogen · MI · Myocardium · Liver

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