

理學碩士學位論文

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Insulin-like growth factor binding protein-3
-5**

2002年 2月

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食品生命科學科

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指導教授 南 澤 正

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主 審

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Effects of glucose on nutrients metabolism and Insulin-like growth factor binding protein-3 and -5 expression in human fibroblasts

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Pukyong National University*

Abstract

Insulin-like growth factor-I(IGF-I) and IGF- are peptides structurally related to insulin. They exert a wide variety of metabolic and mitogenic effects, acting in autocrine, paracrine, and endocrine manners through specific IGF receptors as well as insulin receptors. In contrast to insulin, however, the bioavailability of IGFs to target cells is modulated by the IGF-binding protein(IGFBPs). Six human IGFBPs with different molecular masses, biological characteristics, and immunological properties have been characterized. IGFBPs affect the actions of IGF in both

a positive and negative manner. The level of IGFBP expression is modulated by a variety of systemic and local factors.

In diabetes, most study of the study of IGF systems have been investigated in insulin-dependent diabetes mellitus (IDDM), non-insulin-dependent diabetes mellitus (NIDDM), and streptozotocin-induced diabetes animals *in vivo*. Recently, little research regarding the IGFs system has been proposed in a portion of cells *in vitro*.

Human fibroblasts secrete IGFBPs that can modify IGF-I action. Previous to our study using either Northern blotting, and immunoblotting have shown that fibroblasts express mRNA IGFBP-3, -4, -5 and synthesize these proteins. In addition, fibroblast cell lysates revealed that the IGFBP-3 and IGFBP-5 were most abundant. For these reasons, we undertook to gain further insight into the effects of high and low glucose on metabolism and IGFBP-3, IGFBP-5 expression.

IGFBP-3 and -5 level are more decreased in cell lysates cultured at a high glucose level. IGFBP-3 mRNA of fibroblast cultured in low glucose is increased, whereas IGFBP-5 mRNA remained constant. Although high glucose and low glucose cultures expressed similar levels of IGFBP-5 mRNA, *in vitro* protease assays demonstrated an increase in IGFBP-5 proteolysis in high glucose compared with low glucose cultures. In contrast, IGFBP-3 is not affected by proteolysis. In summary, high glucose acts on human fibroblast to induce a change in IGFBP-3 and -5

mRNA abundance or peptide stability. Our study may be used as a model to investigate the potential roles of IGF-I and IGFBPs in pathophysiologic processes such as diabetes mellitus.

1

가 (1 3). 가 가

(4).

가
(Insulin-Dependent Diabetes Mellitus, IDDM)
(Noninsulin-Dependent Diabetes Mellitus, NIDDM)

(5 7).

,
 , Insulin-like Growth
 Factors(IGFs) insulin 가
 가
 가 (8 12).

, . IGFs
 (IGF Binding proteins, IGFBPs)

.
 IGF-I IGF-
 (mitogenic)
 , prolinsulin
 (13,14). IGFs 가
 (autocrine) (paracrine)
 (15,16).

IGF-I
 . IGF-I
 ,
 . recombinant human IGF-I(rhIGF-I)
 가 IGF-I metabolic, anabolic
 . I
 IGF-I
 , IGF-I .

rhIGF-I glucose

.

IGF-I

, Yde H(1969) IGF-I

(17,18). Maes M

IGF-I

(19). Baxter(20)

IGF-I

, Rieu Binous(21) 가

IGFs IGFBPs ..

IGFs Insulin-like

growth factor binding proteins(IGFBPs)

IGFBPs(IGFBP-1 6) (22). IGFBPs

, IGFBP-1, 3 4 IGF-I

IGF- , IGFBP-2, 5 6 IGF-I

IGF- (23). IGFBPs 가

IGFs IGFs

. IGFBPs

, IGFs (24 28).

IGFBPs IGFBP-1 free IGF-I

IGFBP-1 IGF (29,30).

IGFBP-1 IGFs hypoglycemic activity glucose

(29 32),

, insulin (33), glucose(34) (35)

IGFBP-1 가 (20,27,36,37).

NIDDM IGFBP-1 가 (38 43),
 가 (43 46). IGFBP-2
 (47), 가 (48)
 , streptozotocin (STZ) IGFBP-2 mRNA
 6 8 가 (49), NIDDM 가
 , GH IGF-I (50). IGFBPs
 가 IGFBP-3 IGF-I
 150kDa IGF-I ,
 IGF (20,51).
 insulin catabolic IGF-I
 IGFBP-3 proteolytic activity 가 free IGF-I
 가 가 . IGF-I 가
 IGFBP-3 proteolytic activity가 IGFBP-3
 proteolytic activity 가 .
 IGFBP-3 (52,53),
 IGFBP-3 (54). NIDDM IDDM human
 skin fibroblast IGFBP-2, -3, -5 ,
 IGFBP-4 (55).
 IGFBP-1, -2 mRNA 가 , IGFBP-3 mRNA
 , IGFBP-4 가 (38).
 Heo YR (56) STZ IGFBP-3
 IGFBP-2 가 , IGFBP-4 가
 .

in vivo 가 ,

가 *in vitro*
 . glucose
 GM10 IGFBP-3가 가
 IGFBP-4 -5 , cell lysate
 IGFBP-3 -5가 ,
 IGFBP-4 (57).
 human fibroblast(GM10) glucose
 IGFBP-3 -5
 , GM10
in vitro model system 가 .

2

1.

1.1

Human fibroblasts GM10 . GM10
glucose DMEM(25 mM glucose, Gibco) glucose
DMEM(5 mM glucose, Gibco) sodium bicarbonate 3.7 mg/Mℓ,
penicillin 100 U/Mℓ, streptomycin 100 μg/Mℓ 10% fetal bovine
serum(FBS, Gibco) 가 37 , 5% CO₂
. 가 90% confluent DMEM
(24 h, 72 h, 120 h) 3000 rpm
15 .
IGF-I(100 ng/μℓ), insulin(100 ng/μℓ) 24
(3,000 rpm, 15 min)
- 70 .
PBS 2 , cell lysis buffer(100 mM NaF, 10
mM EDTA, 1 mM benzamidin, 1 mM PMSF, 50 mM Tris-HCl,
pH 7.5) (12,000 rpm, 4 , 3 min)
. cell lysates -70
triglycerides(TG) , total protein(TP) , 1
× sample buffer cell lysate
IGFBP-3 IGFBP-5 immunoblotting .

2.

2.1

DMEM(Dulbeco's Modified Eagle Medium), FBS(fetal bovine serum) Gibco, sodium bicarbonate, penicillin 100 U/Ml streptomycin 100 µg/Ml antibiotics solution, BSA(bovine serum albumin), 1×Trpsin-EDTA, insulin, TRI-reagent, cell extraction buffer Sigma. Rainbow high molecular marker Amersham(RPN756), BCIP, NBT Promega(S3771), Sigma. Mighty Small gel device(Hoefer Scientific Instruments, CA, U.S.A.) Immobilon-PS^Q transfer membrane(Millipore, pore size; 0.1 µm, U.S.A) Hoefer semidary electroblotter(Hoefer Scientific Instruments, CA, U.S.A.)

2.2

Human fibroblast(GM10) glucose(25 mM) DMEM glucose(5 mM) DMEM, 90% confluent monolayer PBS-EDTA 2 trypsin Brightline hemacytometer(Hausser Scientific. U.S.A.)

2.3 Glucose

1.1 GM10 human fibroblast
glucose $10\ \mu\ell$
GLZYME kit glucose .
Kit manual , kit
standard 37 15 가 500 nm
glucose UV-visible spectrophotometer(Varian 1C cary 300.
U.S.A.) .

2.4 Triglycerides (TG)

Human fibroblast(GM10) glucose
monolayer PBS-EDTA 2 trypsin
PBS cell (3,000 rpm, 4 , 3 min)
methanol : chloroform : H₂O = 2 : 1 :
8 가 TG . TG
kit UV-visible spectrophotometer
(Varian 1C cary 300. U.S.A.) 505 nm .

2.5

Human fibroblast glucose cell
lysate . $25\ \mu\ell$ cell lysate
Biuret .
UV-visible spectrophotometer(Varian 1C cary 300. U.S.A.)
450 nm .

2.6

GM 10 glucose(25 mM) glucose(5 mM)
 DMEM 가 90% confluent DMEM
 5 (120 h) lithum
 loading buffer(pH 2.2) (10 ml)
 (Sykam amino acid analyzer. S433) .

2.7 IGFBP-3 mRNA

2.7.1 Total RNA

human fibroblast
 , TRI reagent(Sigma) 1 ml 가 cell lysate
 . Chloroform : isoamyl alcohol (24:1) 200 μ l 가
 30 가 (4 , 12,000 rpm, 15
) RNA가
 isopropanol 가 -70 overnight
 , total RNA 70% ice-cold
 ethanol 2 . total RNA 0.1%
 DEPC 260 nm/280 nm
 , 3 μ g 1% formaldehyde agarose gel
 total RNA band .

2.7.2 Total RNA

Total RNA 10 μ g RNA loading buffer(Sigma, USA) 가
 65 10 1% formaldehyde
 agarose gel . , transilluminater
 (VILBER LOURMAT, FRANCE) UV RNA band
 nylon membrane(ICN BIOTRANSTM) 18 transfer
 . Transfer가 membrane 75 2 baking

hybridization .

2.7.3 Probe labeling

IGFBP-3 DNA North Carolina
David R. Clemmons
, probe PCR DIG Probe synthesis Kit(Boehringer
Mannheim) . MgCl_2 1× PCR reaction
buffer, 200 μM dNTP mixture, PCR DIG-dUTP mixture, 2.6 U
enzyme mixture(ExpandTM High Fidelity), IGFBP-3 sense,
antisense primer 40 pmol, 0.1 U AMV-optimized Tag,
IGFBP-3 DNA PCR PCR Thermal
Cycler 480(TAKARA) PCR . PCR
95 45 , 55 45 , 72 2 10 cycles, 95
45 , 55 45 , 72 2 1 cycle
20 20 cycles . PCR , EtBr(10
mg/ml) 2% agarose gel PCR
band gel . QIAEXII Gel Extraction
Kit(QIAGEN) gel DIG- dUTP label
probe DNA .

2.7.4 Hybridization

Baking membrane 68 1 pre-hybridization(30%
formamide, 0.15 M NaCl, 0.12 M Na_2HPO_4 , 7% SDS, 1 mM
EDTA) pre-hybridization DIG- dUTP label DNA
20 μl 가 hybridization 68 18
hybridization .

2.7.5 Washing blocking

Hybridization membrane 0.1% SDS 2 × SSC
 15 2 , 68 0.5 × SSC 15
 2 washing buffer(0.1 M Maleic acid, 0.15 M NaCl; pH
 7.5; 0.3% (v/v) Tween 20) 5 .
 hybridization , blocking buffer(0.1 M Maleic acid, 0.15 M NaCl;
 pH 7.5; 1% (w/v) blocking reagent) 30
 membrane blocking .

2.7.6 Detection

Anti-Digoxigenin-AP 75 mU/M₀(1:10000) blocking
 buffer(0.1 M Maleic acid, 0.15 M NaCl; pH 7.5; 1% (w/v) reagent)
 30 shaking incubation , washing
 buffer(0.1 M Maleic acid, 0.15 M NaCl; pH 7.5; 0.3% (v/v) Tween
 20) 15 2 . membrane
 detection buffer(0.1 M Tris HCl, 0.1 M NaCl, pH 9.5) 5
 , detection buffer 1:100
 chemiluminescence substrate CSPD solution 5 .
 luminescent , membrane 37 30 1
 가 X-ray .

2.8 IGFBPs

GM10 human fibroblast glucose
 cell lysate . cell lysate 4 × Laemmli
 sample buffer(13.3% SDS, 0.4 M Tris, 0.013% bromophenol blue,
 40% glycerol pH 6.5) 1 × 가 1M DDT (
 0.1 M) 가 Mighty Small II Apparatus(Hofer Science

Instrument) 12.5% nonreducing SDS-polyacrylamide gel
 . maker Rainbow high
 molecular marker(Amersham) . gel
 Hoefer Semi-dry Transfer Unit
 Immobilon-PS^Q transfer membrane(Millipore, pore size; 0.1 μ m,
 U.S.A.) transfer .

2.8.1 IGFBP-3 western immunoblot

Transfer membrane 5% non fat dry milk TBS-T buffer (20
 mM Tris-base, 137 mM NaCl, 1 M HCl, 0.1% Tween 20, pH 7.6)
 1 blocking TBS-T 10 3
 . membrane TBS-T 1:1,500 IGFBP-3
 anti-rabbit 1 1 . 1
 , TBS-T 10 3 TBS-T 1:15,000
 anti-rabbit IgG-conjugated horseradish peroxidase 2
 1 . TBS-T 10 3
 enhanced chemiluminescence sustrate (ECL Western blotting
 detection reagent, Amersham, RPN 2209) IGFBP-3
 .

2.8.2 IGFBP-5 western immunoblot

Transfer menbrane 1 \times TBS 10 ,
 3% BSA가 1 \times TBS IGFBP-5 (1:1,000) 1
 . 1 \times TBS + 0.1% IGEPAL CA-630 + 0.03%
 TritonX-100 1 \times TBS 2 (1:1,000,
 anti- guina pig alkaline phosphase conjugate) 3
 . 1 \times TBS + 0.1% IGEPAL CA-630 + 0.03% TritonX-100,

AP buffer (100 mM Tris, 100 mM NaCl, 5 mM MgCl₂, pH 9.5)
 color substrate solution(NBT/BCIP, Promega)
 , stop buffer (20 mM Tris, 5 mM EDTA, pH 8.0)

2.9 IGFBPs proteolysis

2.9.1 IGFBP-3 proteolysis

GM10 human fibroblasts glucose(5 mM)
 glucose(25 mM) 90% confluent DMEM
 glucose DMEM IGF-I(100 ng/ml),
 insulin(100 ng/ml) 24 50 μ l
 0.5 M tris 10 μ l 100 mM CaCl₂ 2 μ l 7† intact
 IGFBP-3 100 ng 37 water bath
 . 4× sample buffer .
 12.5% nonreducing SDS-polyacrylamide gel .
 transfer membrane 1×TBS 5 3%
 BSA 1×TBS 30 blocking . 1×BSA7†
 1×TBS IGFBP-3(1:1500) 1 . 1×
 TBS + 0.1% IGEPAL CA-630 + 0.03% TritonX-100
 1×TBS 2 (1:2000, anti-rabbit IgG alkaline
 phosphatase conjugate) 3 . 1×TBS + 0.1%
 IGEPAL CA-630 + 0.03% TritonX-100, 1×TBS, AP buffer(100
 mM Tris, 100 mM NaCl, 5 mM MgCl₂, pH 9.5)
 color substrate solution(NBT/BCIP, Promega) .

band stop buffer(20 mM Tris, 5 mM EDTA, pH 8.0) . membrane .

2.9.2 IGFBP-5 proteolysis

glucose(25 mM) 90% confluent
 IGF-I(100 ng/ml), insulin(100 ng/ml)
 50 μ l glucose(5 mM) glucose(25 mM)
 50 μ l 0.5 M tris 10 μ l, 100 mM CaCl₂ 2 μ l
 가 (50
 mM EDTA, 2 mM PMSF, 1 mM 1,10-phenathroline, 5 mM
 aprotinin, 0.2 mM E64, 3 mM Benzamide, 1 μ g/Ml heparin)
 30 intact IGFBP-5 100 ng 가
 37 water bath . 4 \times sample buffer
 12.5% nonreducing SDS-polyacrylamide gel
 transfer membrane western immunoblot
 IGFBP-5 , inhibitor
 IGFBP-5 .

2.10 IGFBP-5 mRNA

2.10.1 Total RNA

GM10 human fibroblast
 , TRI reagent(Sigma) 1 Ml 가 scrapping cell
 lysate . Chloroform : isoamyl alcohol (24:1) 200 μ l
 가 30 (4 , 12,000 rpm, 15
) RNA가

isopropanol 가 -70 overnight . overnight
 , total RNA 70% ice-cold
 ethanol 2 . total RNA 0.1%
 DEPC 260 nm/280 nm
 , 3 μ g 1% formaldehyde agarose gel
 total RNA band -70 RT-PCR

2.10.2 Reverse transcription-polymerase chain reaction(RT-PCR)

Reverse transcription-polymerase chain reaction(RT-PCR)
 TAKARA one step RNA PCR kit . Total RNA(10
 μ g) 1 \times one step RNA PCR reaction buffer, 5 mM MgCl₂, 1 mM
 dNTP mixture, 0.8 U RNase inhibitor, IGFBP-5 sense, antisense
 primer 40 pmol, 0.1 U AMV reverse transcriptase, 0.1 U
 AMV-optimized Tag . PCR Thermal
 Cycler 480(TAKARA) 50 30 , 94 2
 reverse transcription 94 30 , 65 30 , 72
 30 30 cycles, 72 5 .
 RT-PCR IGFBP-5 sense, antisense primer

5'- GAA GGC TGA AGC AGT GAA GAA GGA CCG -3'
 (Sense)

5'-ACT CAA CGT TGC TGC TGT CGA AGG TGT -3'
 (Antisense)

RT-PCR , EtBr(10 mg/ml) 2% agarose gel
 RT-PCR , 100 bp molecular

weight marker(Bio Labs, 100 bp DNA Ladder)

2.11 Gelatin zymography

GM10 human fibroblast (25 mM) (5 mM)
 glucose 7.5 mM DMEM 90% confluent
 DMEM glucose DMEM IGF-I(100 ng/ml),
 insulin(100 ng/ml) 7.5 mM .
 glucose 4 × sample buffer
 60 10 10% gelatin 10%
 SDS-polyacrylamide gel . gel 2.5%
 Triton X-100 4 10 4 2
 5-6 , 4 mM CaCl₂ 50 mM Tris buffer(pH
 7.4) 37 fix solution(7% acetic
 acid, 40% methanol) staining solution(0.1%
 coomassie brilliant blue G-250, 10% acetic acid)
 destaining solution(7% acetic acid, 20% methanol) .

1. Glucose 가

Human fibroblast glucose(5 mM) glucose(25 mM) ,
 glucose $47.9 \pm 5 \times 10^4$, $45.1 \pm 4.3 \times 10^4$ cells/
 cm^2 glucose cell growth 가
 (Table 1).

Stefano Giannini (55) ,
 (Noninsulin-Dependent Diabetes Mellitus, NIDDM),
 (Insulin-Dependent Diabetes Mellitus, IDDM),
 fibroblasts $3.8 \pm 0.7 \times 10^4$, $4.5 \pm 0.7 \times 10^4$,
 $4.1 \pm 0.6 \times 10^4$, $3.8 \pm 0.7 \times 10^4$ cells/ cm^2 cell growth
 , Busiguina S (58)
 $19.4 \pm 0.5 \times 10^4$, $18.7 \pm 1.0 \times 10^4$ cells/ cm^2
 ,
 . glucose

glucose

Table 1. Effect of glucose levels on cell numbers

Glucose concentration	Cell number (cells/cm ²)
5 mM	$45.1 \pm 4.3 \times 10^4$
25 mM	$47.9 \pm 5.0 \times 10^4$

- Values are the mean \pm SE (n=3)
- Cell : Human fibroblast (GM10)

2. Glucose 가

2.1

Figure 1 human fibroblast(GM 10) glucose(25 mM)
 glucose(5 mM) , glucose
 (Figure 1). glucose 72
 glucose 24 10%
 , 120 25% .
 glucose 72 35%
 , 120 84% .
 glucose ,
 glucose glucose
 . Henry RP (59)
 glycogen ,
 ,
 Kahn CR(60) streptozotocin(STZ)
 , STZ Langerhan's -
 가 , 가
 가 ,
 (61) STZ-diabetic rats ,
 . Luciano
 R (62) IGF-I glucose uptake
 가 가 IGF-I
 glucose .

in vivo

.

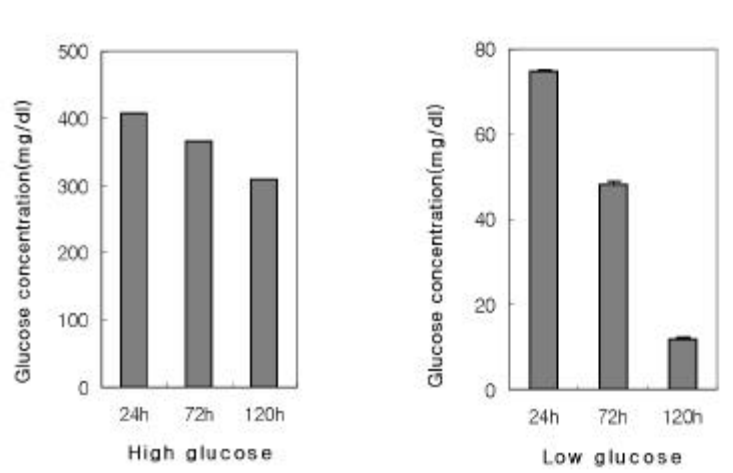


Figure 1. Time course of glucose consumption in conditioned medium of human fibroblast.

Cells were cultured for 24, 48, and 120 h in a medium containing either high glucose(25 mM) and low glucose(5 mM). conditioned medium were collected after the indicated hours of cultures and analysed by GLZYME kit as described in materials and methods.

Figure 2

Figure 1

가

glucose

triglyceride

glucose

glucose

glucose

triglyceride(TG)

가

glucose

Mardar Z (65)

, Niall (66)

(IDDM)

lipoprotein lipase

가

가

chylomicron

(67),

90%

가

(62,63),

HDL-

HDL-

(64).

,
in vivo

glucose

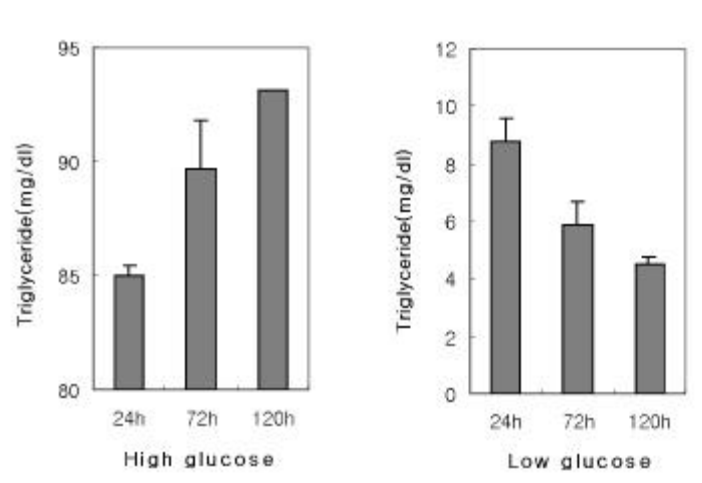


Figure 2. Time course of triglyceride levels in human fibroblast.

Cells were cultured for 24, 48, and 120 h in a medium containing either high glucose(25 mM) and low glucose(5 mM). cell lysate were collected after the indicated hours of cultures and analysed TRIGLYZYME-V kit as described in Materials and Methods.

2.3

Figure 3 GM10 glucose
total protein , glucose
total protein 가 ,
glucose
. Insulin
가 가 .
가 가 가
glucogenesis 가
(68). Green M (69) STZ
insulin
(70 72). ,
glucose 가 가
Table 2 glucose 5 (120 h)
,
, glucose
glucose .
glucose glucose
glucose
glucose

glucose

. cystine, glutamine, serine

glucose 58%, 80%, 49% ,

glucose 60%, 81%, 61%

glucose . cystine, glutamine,

serine

. , control glycine, phenylalanine, threonine

glucose 가 .

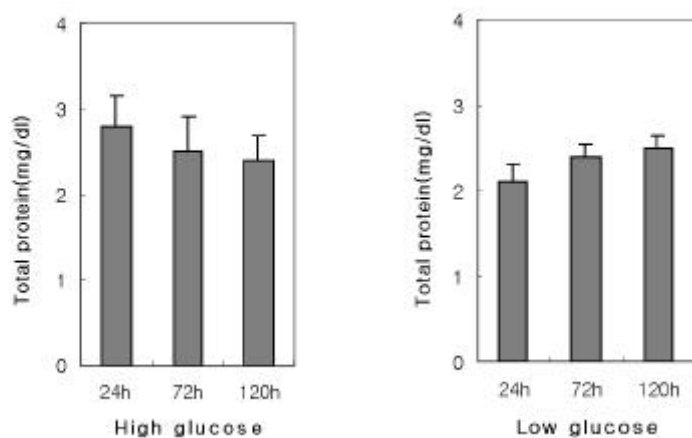


Figure 3. Time course of total protein concentration in cell lysate of human fibroblast.

Cells were cultured for 24, 48, and 120 h in a medium containing either high glucose(25 mM) and low glucose(5 mM). Cell lysate were collected after the indicated hours of cultures and analysed by TP-M kit as described in materials and methods.

Table 2. Free amino acid composition of human fibroblast high and low glucose conditioned medium.

Amino acid	Sample		
	Control	High (120h)	Low (120h)
Arginine	84	68	61
Cystine	63	27	25
Glutamine	584	114	111
Glycine	30	55	54
Histidine	42	28	26
Leucine	105	97	84
Lysine	146	125	117
Methionine	30	29	27
Phenylalanine	66	73	68
Serine	42	21	16
Threonine	95	107	102
Tryptophan	16	-	-
Tyrosine	104	73	67
Valine	94	95	88
Total	1,501	912	846

3. Glucose 가 IGFBP-3

Figure 4 GM10 glucose
IGFBP-3 ,
glucose . Baxter Martin (20)
IGFBP-3
, Rine Binoux (76)
IGFBP-3가 ,
, IGFBP-3 (77) (20)
(78 82).

STZ
IGFBP-3 (54,56),
IGFBP-3 protease activity 가 intact IGFBP-3가
(83), GH (84) (54)
(85) IGFBP-3가 . *in vivo*
glucose IGFBP-3

Figure 5 glucose IGFBP-3
mRNA , glucose IGFBP-3
mRNA .

IGFBP-3 mRNA
(38) Spoerri PE (86) IGFBP-3 mRNA
가 .

Figure 6 glucose
glucose IGF-I insulin

IGFBP-3 proteolysis human fibroblast
 (GM10) IGFBP-3 protease
 . Clemmons(81) human dermal
 fibroblasts porcine aortic smooth muscle cells
 IGFBP-3 , IGFBP-3 proteolytic
 . IGFBPs 가
 IGFBP-3 , glucose가
 IGFBP-3 protease
 , glucose
 .

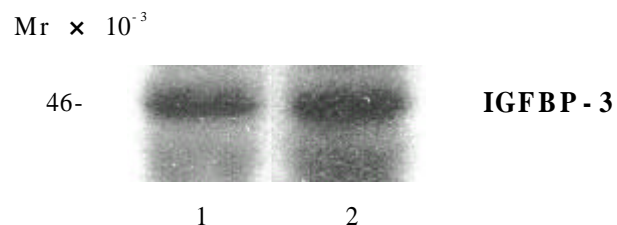


Figure 4. IGFBP-3 ECL western immunoblotting of human fibroblast cell lysates.

Collected cell lysates from medium containing either high glucose or low glucose, after 48 h culture, were subjected to SDS-polyacrylamide gel electrophoresis, blotted onto a nitrocellulose membrane, and analyzed ECL immunoblotting, as described in *Materials and Methods*. Molecular marker are indicated on the *left*.

lane 1 : High glucose(25 mM)

lane 2 : Low glucose(5 mM)

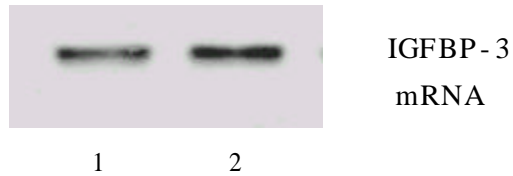


Figure 5. Effect of glucose on IGFBP-3 gene expression.

Cells were cultured for 24 h in a medium containing high glucose and low glucose. Total RNA was extracted from cultures of human fibroblast(GM10) and was analyzed by northern blotting using a specific IGFBP-3 cDNA as described in *Materials and Methods*.

lane 1 : high glucose(25 mM)

lane 2 : low glucose(5 mM)

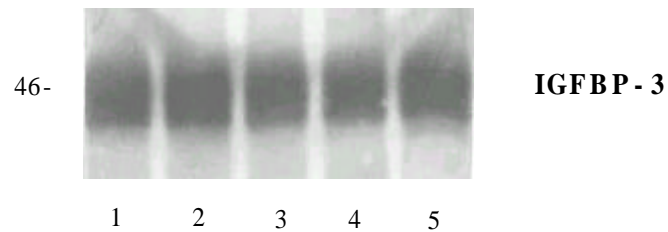


Figure 6. Effects of high glucose, IGF-I, insulin low glucose on IGFBP-3 protease activity.

Cells were cultured for 24 h in a medium containing high glucose, high glucose treated with IGF-I, high glucose treated with insulin and low glucose. Conditioned medium were incubated with intact IGFBP-3(100 ng) for 14 h at 37 °C, and the products were analyzed by immunoblotting.

- lane 1 : Conditioned medium plus intact IGFBP-3 and no incubation
- lane 2 : Conditioned medium containing high glucose(25 mM) plus intact IGFBP-3
- lane 3 : Conditioned medium containing high glucose(25 mM) treated with IGF-I(100 ng/mL) plus intact IGFBP-3
- lane 4 : Conditioned medium containing high glucose(25 mM) treated with insulin(100 ng/mL) plus intact IGFBP-3
- lane 5 : Conditioned medium containing low glucose(5 mM) plus intact IGFBP-3

4. Glucose 가 IGFBP-5

lysate glucose human fibroblast cell
 IGFBP-5
 glucose IGFBP-5 (Figure 7).
 human fibroblast IGF-I IGF- 가 IGFBP-5
 mRNA protease IGFBP-5 가
 (73), IGFBP-5 mRNA 가 (74,75),
 smooth muscle cell(88), chondrocytes osteoblast(89)
 IGF-I 가 IGFBP-5 mRNA 가
 . human fibroblast IGF-I 가
 IGFBP-5
 가 glucose IGF-I 가
 IGFBP-5 .
 (IDDM), (NIDDM) human
 skin fibroblast IGFBP-5 (55).
 Figure 8 IGFBP-5 ,
 glucose, glucose IGF-I insulin
 가 GM10 IGFBP-5 mRNA
 IGFBP-3 .
 IGFBP-5 mRNA 가
 (14), IGFBP-5 mRNA
 (90). Figure 7 IGFBP-5가
 , Figure 8 IGFBP-5 mRNA
 glucose 가 . Nam (91)

human fibroblast IGFBP-5 protease가
glucose
IGFBP-5 가 protease IGFBP-5
protease activity glucose (Figure 9).
, glucose 30 kDa intact IGFBP-5가
, 22 kDa IGFBP-5 가
, glucose IGFBP-5
가 glucose
IGFBP-5 IGFBP-3 protease
.

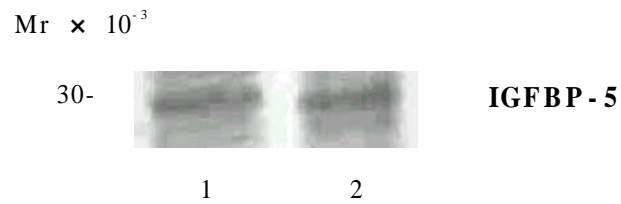


Figure 7. IGFBP-5 western immunoblot of human fibroblast cell's lysates .

After SDS-polyacrylamide gel electrophoresis (12.5% SDS-PAGE) of cell lysates were electroblotted onto immobilon-PS^Q membrane and incubated with IGFBP-3 antibody. Molecular markers are indicated on the *left*.

lane 1 : high glucose(25 mM)

lane 2 : low glucose(5 mM)

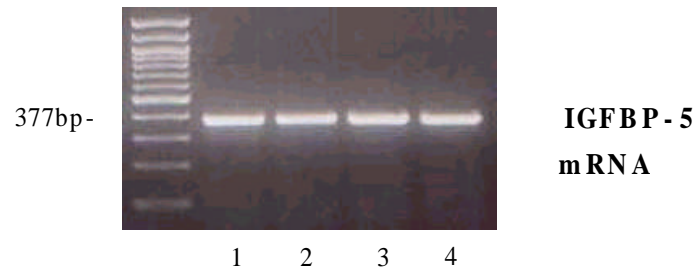


Figure 8. Effects of high glucose, IGF-I, insulin low glucose on of IGFBP - 5 gene expression.

Cells were cultured for 24 h in a medium containing high glucose, high glucose treated with IGF-I, high glucose treated with insulin and low glucose. Total RNA was extracted from cultures of human fibroblast(GM10) and used for RT-PCR as described in *Materials and Methods*.

lane 1 : high glucose(25 mM)

lane 2 : high glucose treated with IGF-I(100 ng/M \emptyset)

lane 3 : high glucose treated with insulin(100 ng/M \emptyset)

lane 4 : low glucose(5 mM)

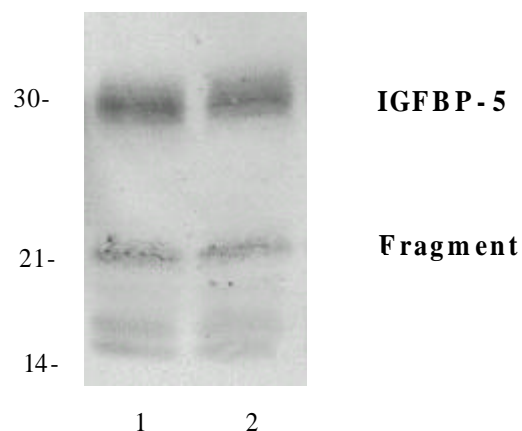


Figure 9. Effect of glucose on IGFBP-5 protease activity.

Cells were cultured for 24 h in a medium containing high glucose and low glucose. Conditioned medium were incubated with intact IGFBP-5(100 ng) for 14 h at 37 °C, and the products were analyzed by immunoblotting. The relative migration position of fragments are also noted by arrows.

lane 1 : Conditioned medium containing high glucose(25 mM)
plus intact IGFBP-5

lane 2 : Conditioned medium containing low glucose(5 mM)
plus intact IGFBP-5

5. Protease inhibitor

IGFBP-5 protease

glucose IGFBP-5
 protease activity 가 glucose IGF-I, insulin
 가 IGFBP-5
 inhibitor proteolysis
 .
 Figure 10 13 glucose IGF-I
 insulin 가 EDTA, PMSF,
 1,10-phenathroline, aprotinin, E64, benzamide, heparin
 protease western immunoblot . ,
 metalloprotease inhibitor EDTA ,
 1,10-phenathroline . Serine
 protease inhibitor PMSF IGF-I
 protease , aprotinin
 glucose protease
 . serine protease metalloprotease
 inhibitor heparin protease
 . E64, benzamide
 .
 glucose IGF-I insulin IGFBP-5 protease
 , glucose IGF-I
 22 kDa 20 kDa fragment ,
 heparin 20 kDa fragment가 .

Insulin 22 kDa 17, 16 kDa fragment .
, IGF-I insulin IGFBP-5 protease activity
, human fibroblast IGFBP-5
serine protease metalloprotease
. Human fibroblast IGFBP-5
serine protease metalloprotease가 , serine
protease가 22, 17, 16 kDa fragment
, metalloprotease가 20 kDa fragment
. serine protease inhibitor aprotinin,
3,4-DC , phenylmethyl-sulfonyl fluoride(PMSF) metalloprotease
inhibitor ethylenediaminetetraacetic acid(EDTA), 10-phenanthroline
IGFBP-5 proteolysis , E64,
Benzamide, NEM, iodoacetic acid protease
(91,92)
glucose IGFBP-5 가
, glucose
inhibitor IGFBP-5
, glucose IGFBP-5
.
in vivo ,
가 glucose , IGF-I insulin 가
IGFBP-5 protease activity
IGFBP-5 protease가 glucose
가 가 .

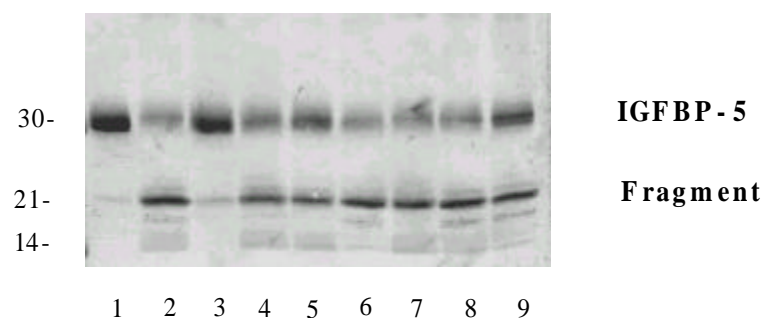


Figure 10. Analysis of effects of protease inhibitors on IGFBP-5 protease activity in high glucose conditioned medium.

Cells were cultured for 24 h in a medium containing high glucose. Conditioned medium were incubated with intact IGFBP-5(100 ng) and various protease inhibitor for 14 h at 37 °C, and the products were analyzed by immunoblotting. The relative migration positions of fragments are also noted by arrows.

- lane 1 : Conditioned medium plus intact IGFBP-5 and no incubation
- lane 2 : Conditioned medium containing high glucose(25 mM) plus intact IGFBP-5
- lane 3 : EDTA(50 mM)
- lane 4 : PMSF(2 mM)
- lane 5 : 1,10-phenanthroline(1 mM)
- lane 6 : aprotinin(5 mM)
- lane 7 : E64(0.2 mM)
- lane 8 : benzamide(3 mM)
- lane 9 : heparin(1 μ g/ml)

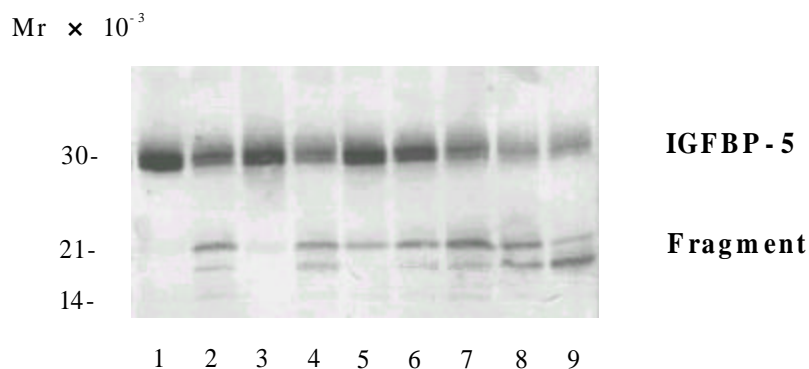


Figure 11. Analysis of effects of protease inhibitors on IGFBP-5 protease activity in human fibroblast high glucose conditioned medium treated with IGF-I.

Cells were cultured for 24 h in a medium containing high glucose. Conditioned medium were incubated with intact IGFBP-5(100 ng) and various protease inhibitor for 14 h at 37 °C, and the products were analyzed by immunoblotting. The relative migration positions of fragments are also noted by arrows.

- lane 1 : Conditioned medium plus intact IGFBP-5 and no incubation
- lane 2 : Conditioned medium containing high glucose(25 mM) plus intact IGFBP-5
- lane 3 : EDTA(50 mM)
- lane 4 : PMSF(2 mM)
- lane 5 : 1,10-phenanthroline(1 mM),
- lane 6 : aprotinin(5 mM)
- lane 7 : E64(0.2 mM)
- lane 8 : benzamide(3 mM)
- lane 9 : heparin(1 $\mu\text{g}/\text{Ml}$)

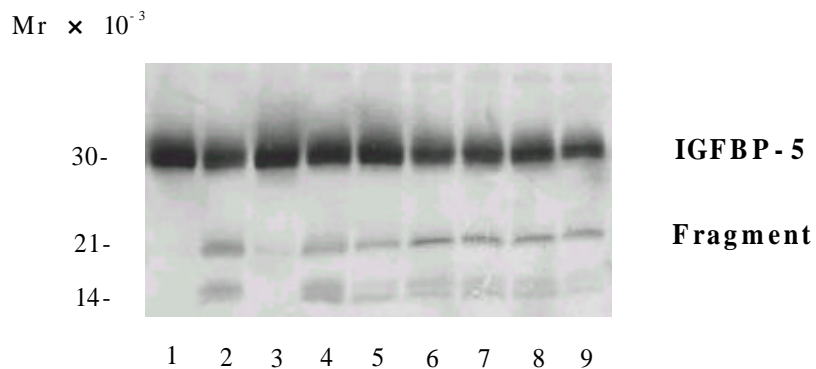


Figure 12. Analysis of effects of protease inhibitors on IGFBP-5 protease activity in human fibroblast high glucose conditioned medium treated with insulin.

Cells were cultured for 24 h in a medium containing high glucose. Conditioned medium were incubated with intact IGFBP-5(100 ng) and various protease inhibitor for 14 h at 37 °C, and the products were analyzed by immunoblotting. The relative migration positions of fragments are also noted by arrows.

- lane 1 : Conditioned medium plus intact IGFBP-5 and no incubation
- lane 2 : Conditioned medium containing high glucose(25 mM) plus intact IGFBP-5
- lane 3 : EDTA(50 mM)
- lane 4 : PMSF(2 mM)
- lane 5 : 1,10-phenanthroline(1 mM)
- lane 6 : aprotinin(5 mM)
- lane 7 : E64(0.2 mM)
- lane 8 : benzamide(3 mM)
- lane 9 : heparin(1 $\mu\text{g}/\text{Ml}$)

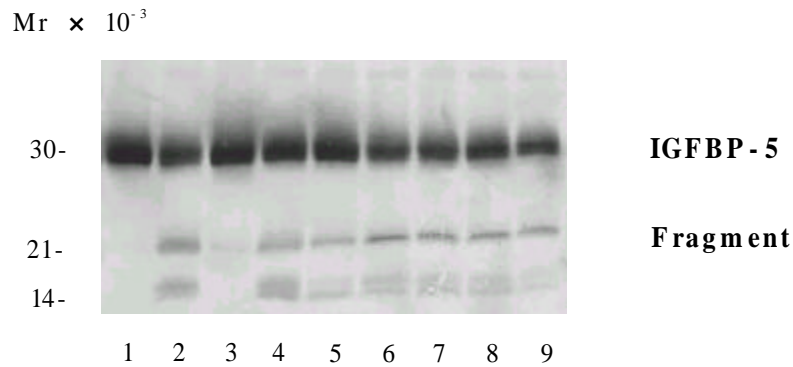


Figure 13. Analysis of effects of protease inhibitors on IGFBP-5 protease activity in human fibroblast low glucose conditioned medium.

Cells were cultured for 24 h in a medium containing high glucose. Conditioned medium were incubated with intact IGFBP-5(100 ng) and various protease inhibitor for 14 h at 37 °C, and the products were analyzed by immunoblotting. The relative migration positions of fragments are also noted by arrows.

- lane 1 : Conditioned medium plus intact IGFBP-5 and no incubation
- lane 2 : Conditioned medium containing high glucose(25 mM) plus intact IGFBP-5
- lane 3 : EDTA(50 mM)
- lane 4 : PMSF(2 mM)
- lane 5 : 1,10-phenanthroline(1 mM),
- lane 6 : aprotinin(5 mM)
- lane 7 : E64(0.2 mM)
- lane 8 : benzamide(3 mM)
- lane 9 : heparin(1 $\mu\text{g}/\text{Ml}$)

6. IGFBP-5 Protease

IGFBP-5 metalloprotease가 gelatin zymography . glucose protease . Figure 14 glucose IGF-I insulin metalloprotease 72, 69, 55 kDa gelatinase 가 , 69 kDa gelatinase가 가 glucose band . Figure 15 glucos protease gelatin zymography , glucose protease 가 , Figure 14 가 glucose protease . metalloprotease (93) *in vivo* 가 glucose IGFBPs protease .

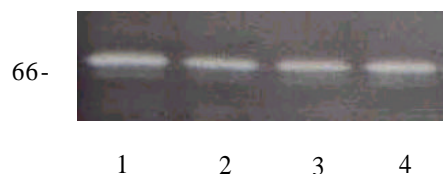


Figure 14. Characterization of gelatin-degrading protease activity in human fibroblast conditioned medium

Cells were cultured for 24 h in a medium containing high glucose, high glucose treated with IGF-I, high glucose treated with insulin and low glucose. Conditioned medium were analyzed by gelatin-substrate zymography. Sample were electrophoresed on 10% SDS-polyacrylamide gels that contained cross-linked gelatin. After electrophoresis, the gel incubated for 14 h at 37 °C. After fixing, staining, and destaining, take a photography of gel. Areas of lysis(*clear zone*) represent gelatin-degrading protease activity in the sample. Molecular markers are indicated on the *left*.

lane 1 : Conditioned medium containing high glucose(25 mM)

lane 2 : Conditioned medium containing high glucose(25 mM)
treated with IGF-I(100 ng/Ml)

lane 3 : Conditioned medium containing high glucose(25 mM)
treated with insulin(100 ng/Ml)

lane 4 : Conditioned medium containing low glucose(5 mM)

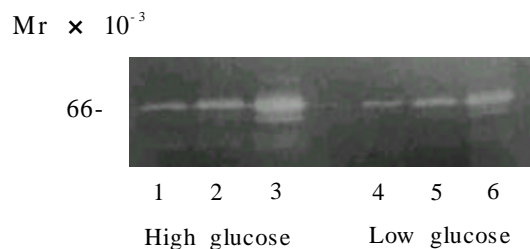


Figure 15. Time course of characterization of gelatin-degrading protease activity in human fibroblast conditioned medium.

Cells were cultured for 24 h in a medium containing high glucose, high glucose treated with IGF-I, high glucose treated with insulin and low glucose. Conditioned medium were analyzed by gelatin-substrate zymography. Sample were electrophoresed on 10% SDS-polyacrylamide gels that contained cross-linked gelatin. After electrophoresis, the gel incubated for 14 h at 37 °C. After fixing, staining, and destaining, take a photography of gel. Areas of lysis (*clear zone*) represent gelatin-degrading protease activity in the sample. Molecular markers are indicated on the *left*.

lane 1(24 h), 2(72 h), 3(120 h) : Conditioned medium containing high glucose(25 mM)
 lane 4(24 h), 5(72 h), 6(120 h) : Conditioned medium containing low glucose(5 mM)

4

Human fibroblasts glucose 가 IGFBP-3 -5

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1. GM10 human fibroblast glucose

.

2. glucose(25 mM) glucose(5 mM)
glucose , glucose
glucose .

3. triglyceride glucose 가
, glucose .

4. glucose ,
glucose 가 .

5. glucose 5
glucose 가
.

6. glucose , glucose IGFBP-3
, human
fibroblast IGFBP-3 protease
.

7. glucose IGFBP-5 IGFBP-5

mRNA 가 .

8. glucose IGFBP-5 가 ,
GM10 human fibroblast IGFBP-5 serine
protease metalloprotease .

9. Gelatin zymography protease glucose
, 가 .

in vivo , IGFs
가 가

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