

(19) (KR)  
(12) (A)

(51) . Int. Cl. 7 (11) 2001 - 0071530  
A61K 39/395 (43) 2001 07 28

(21) 10 - 2000 - 7014459  
(22) 2000 12 19  
2000 12 19  
(86) PCT/NL2000/00253 (87) WO 2000/63251  
(86) 2000 04 19 (87) 2000 10 26

(71) *...the first time I had seen him, he was wearing a dark suit and a white shirt with a tie.*

(72)

(74)

:

(54)

T

C

T

C

T

ICAM

HIV

C

가

,

,

,

C

가

,

T

,

,

C

가 T

ICAM

,

T

1 T

가

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가

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HIV

가

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C

가

T

,

가

가 /

T C T C  
" ICAM - 2 ICAM - 3 ICAM ICAM - 3  
. " ICAM

가 C / C  
C T ,

WO 96/23882 C  
, 10 ( " DEC - 205" ) 7.5 205 kDa  
C (MMR)

WO 96/23882 DEC - 205 DEC - 205 ( 가 )  
DEC - 205 가 T DEC - 205 가 ) ( ,  
(GVH) DEC - 205 ) T  
T ( , ) DEC - 205

Curtis , Proc.Natl.Acad.SciUSA, 89(1992) p.8356 - 8360 WO 93/01820  
CD4 gp120 gp120 , ,  
CD4 가 , , T , B ( , ). WO 93/01820  
0 1 2 C gp120  
C 가 (> 98 %).

Curtis WO 93/01820 C 가 HIV / gp120 HIV gp120  
가 (internalization) C gp120 / HIV , WO 93/01820  
HIV A , A N -  
- D - , HIV가

WO 93/01820 COS - 7 HIV gp120 gp120 ( " 110.1" )  
, gp120 C

, Curtis

WO 93/01820

C

T

WO 95/32734

)

B7 - 1/2

Fc R (CD 32)  
(APCs) Fc R (CD 32)

( ,

),  
ICAM - 3  
; Fc R가  
T  
( ,  
2가  
가IgG  
IgG Fc

Fc

WO 95/32734 T

( , ),

T  
T ICAM - 3

( , TcR/CD3

1  
WO 95/32734

( ) C

WO 98/02456

O 98/49306  
1633

(" PAP" )

C

C

. WO 98/4

WO 98/02456, WO 98/49306 WO 98/41633

C

C , C ,

C

;

(DC)

T

1 T

, DC HIV - 1  
HIV DCCD4<sup>+</sup> T

1

[Spira , 1996 ; Joag , 1997]. HIV - 1 DC  
[Granelli - Piperno , J.Virol 72(4), 2733 - 7, 1998][Blauvelt  
, Nat Med 3(12), 1369 - 75, 1997], HIV - 1 가 DC T

[Cameron , Scienc

e 257(5068), 383 - 7, 1992]. HIV - 1

DC T HIV - 1 AIDS CD4<sup>+</sup> T HIV

, WO 95/32734 WO 96/2

3882

DC T  
 DC - T  
 ICAM - 3  
 LFA - 1, D  
 ICAM - 3  
 DC C  
 DC - T  
 DC  
 , DC - SIGN DC  
 HIV - 1 T  
 HIV - 1 T  
 DC H  
 IV - 1

1 , , C

, T , , C

, T , , C

, C T ICAM ,  
 C

가 T  
 . C Curtis  
 6 - 8360 WO 93/01820 C gp120 , Proc.Natl.Acad.Sci.USA, 89(1992), p.835  
 가 (< 98 %). C , WO 93/01820 1  
 ; 44 kDa ; 404 1  
 2 (Ile 77 Val 249) (Met 1 C  
 3 (Cys 253 Ala 404) 가  
 , C T ICAM

, T T ( , / ),  
 ( , / )

가 T 가 , T T , , T  
 , T T , , T

가 , T 1 / ( , / T NHC -  
 ) T , , ,

; , T , , , WO 95/32  
 734 WO 96/23882

,  
/ ; (1 ) T .

T 가 T ICAM T C  
T , T T T 가

, , , /  
HIV T HIV가  
. , , , T HIV

가, HIV . T HIV " " , HIV가 , 가

T , HIV AIDS  
T , /

가 , HIV가 , HIV ( , gp120)  
HIV 가 ,

C 가 T HIV 가 C ,

WO 93/01820 C

, C , 1/ 9 C (  
가 ) , .  
D - ; L - ; ; A ; ( ;  
A ; N - - D - ;  
) ;  
. , gp120 C  
ICAM C .  
가

가 / HIV A

, WO 93/01820  
· , gp120  
(WO 93/01820 ).

C WO 95/327  
34, WO 96/23882, WO 98/02456, WO 98/41633 / WO 98/49306

, , , , , ( , le limpet hemocyanin) ) / Freund's, , ISCOM's, ( 1 7 ), , , WO 96/23882 ( , ) . C . C (keyho

C 가 , 1 2 9  
가 . C , ( ) 가 ;  
 , AZN - D1 AZN - D2 2 .

가 C , 1 2  
 9 , ( ) ,  
 AZN - D1  
 AZN - D2 ECACC 990400818 990400819  
 (European Collection of Cell Cultures) 1999 4 8

C ) 가 ( ) , 가 ( , ) / ) . - -

, C C ,

; , , , ( / ), , /

, / ( , US - A - 5,6 )

43,786

가

,

가 ( , ) .

( / C ) /  
/ , ELISA

FACS , / 가 ( )

D.

P. Sites, A.I.Terr, T.G. Parslow : " Basic and clinical immunology" , 8th Ed., Prentice - Hall(1994) ; I. Rott, J.Brostoff, D.Male:" Immunology" , 2nd Ed., Churchill Livingstone(1994) ;

Janeway - Travers : " Immunobiology, the immune system in health and disease" , 3rd Edition

2.7 2.17

c 2 , , / , A , ,

, LF3A

( , ), , , 가 (TNF) HIV ( , WO 95/3234 WO 9  
 6/23882 ), (TNF) HIV , Leu - 3A CD4  
 CD4

CD4

/ , , , ,  
( , ) , , / WO 93/01820, WO 95/327

34, WO 96/23882, WO 98/02456, WO 98/41633 / WO 98/49306  
WO 93/01820 .

WO 93/01820, WO 95/327

WO 95/32734, WO 96/23882, WO 98/02456, WO 98/41633

/ WO 98/49306

가 , , , ,

가 , c

T

T

1

,  
A A ; N -  
- D - ( ) ; ,

, gp120 C ICAM - , C , AZN - D1 AZN - D2 ( , ) .

gp100, g250, p53, MAGE, BAGE, GAGE, MART1,

, , , , , ( ) , (Haemophilus influenzae), (Neisseria), (Bordetella pertussis), (Pneumococcus) (Polyomyletus)

, , / HIV

가

1) C ; , 가 / , 2)

, /  
WO 93/01820, WO 95/32734 / WO 96/23882

, WO 93/01820, WO 95/32734,  
WO 96/23882, WO 98/02456, WO 98/41633 / WO 98/49306

, 44 kDa C  
, 1 9 , C  
, / - T .  
(80% 90% ) / C ;  
; /T  
T , WO 96/23882  
C

2a 2c ICAM - 3 DC AZN - D1 AZN - D2 가 DC

3a 3b SDS - PAGE( 3a) DC DC - SIGN HIV g  
p120 C DC - SIGN .

4a 4b COS7 DC - SIGN DC - SIGN AZN - D1 ICAM - 3

5 (A B) (C D) DC - SIGN C - SIGN (OM  $\times$  100). D

6a 6d ICAM - 3 DC DC - SIGN ( 6a ) ; K562 - ICAM - 3 DC  
 ( 6b ) ; PBL DC ; DC T DC -  
 SIGN - ICAM - 3 ( 6d ), ICAM - 3 DC DC - SIGN  
 DC - T

7 DC - SIGN DC HIV - 1

8 DC - SIGN ICAM - 3 ICAM - 3 K562

## 9 DC - SIGN

가 T , (DC) 2 . H  
 IV - 1 CD4<sup>+</sup> T DC DC C , HIV - 1  
 DC - SIGN , ICAM - 2 / ICAM - 3 DC - T DC T  
 , DC - SIGN , DC HIV - 1 CD4 HIV  
 , DC HIV - 1 DC -  
 가 DC , 1  
 SIGN

1 : ICAM - 3 DC

1 DC - T ICAM - 3 ICAM - 3 - Fc  
 (Geijtenbeek ), DC , (PBL) IL - 4 GM - CSF  
 7 DC 2 Ca<sup>2+</sup> ICAM - 3  
 (72%, 1a). 1 DC ICAM - 3  
 . 1a, 1b 1c 3 가 (SD < 5%).

1a : ICAM - 1 ICAM - 3

PBL, DC 30 37 , ICAM - 1Fc ICAM - 3Fc  
 , %

1b : 2 ICAM - 3

980 nM PMA 2 KIM185 (10 µg/Mℓ) , 37 30 ICAM -  
 3Fc . PMA LFA - 1  
 LFA - 1 NKL - L15 (20 µg/Mℓ)

1c : 2 (NKI - L19), 1 (AIIB2), ICAM - 3(CBR - IC3/1, CBR - IC3/2)  
 20 µg/Mℓ EDTA(5mM) EGTA(5mM) ICAM - 3 DC  
 1a

ICAM - 3 DC , PBL(resting PBL) ICAM - 3  
 ( 1a). ICAM - 3 DC 가 L 2  
 ( 1a). , ICAM - 3 가 L  
 - 1 2 ( D 2 , M 2 , X 2 ; DC ICAM - 3 )  
 . ( 1, 7 , 1b) DC  
 , ICAM - 3 DC 가

DC ICAM - 3 - Fc ICAM - 3 , ICAM - 3 , CBR3/1, C  
 BR3/2 가 ( 1b). , DC  
 ICAM - 3 EDTA EGTA ( 1b).

DC 가 1 2 Ca<sup>2+</sup> DC가 ICAM -  
 3 . . . :DCDC - SSpecificIC  
 AM - 3GGrabbingNNon - integrin (DC - SIGN).

2 : DC - SIGN DC - ICAM - 3

DC - SIGN , ICAM - 3 가 . DC  
 SP2/0 . . DC ICAM - 3  
 . 2 ,  
 AZN - D1 AZN - D2 DC 가 ICAM - 3 ( 2a).  
 , ICAM - 3 LFA - 1  
 2 AZN - D1 AZN - D1가 ICAM - 3 DC DC

2a : AZN - D1 AZN - D2 (20 µg/Mℓ) DC . 3 가 , ICAM - 3Fc (SD < 5%).

2b : DC , DC - SIGN 가 . DC GM - CSF IL - 4  
 . CD14, 2 LFA - 1 DC - SIGN  
 . .  
 . 3 가

2c : GM - CSF IL - 4 DC가 ICAM - 3 DC - SIGN  
 . 가  
 2 AZN - L19 AZN - D1 (20 µg/Mℓ) . AZN - D2 AZN - D1 ICAM - 3Fc  
 1 . . . . ICAM - 3 (SD < 5%).

2d : DC ICAM - 3 2 DC - SIGN  
 2c AZN - D1 AZN - L19

AZN - D1 DC - SIGN 1 DC - SIGN  
 ( 2b). DC - SIGN . DC - SIGN  
 . 가 ( 2b). 7 ( 2  
 b). 가 . MHC , , 2  
 p150, p95 CD86 ( ) DC ,  
 . DC  
 LFA - 1

, . 가 DC , ICAM - 3 DC - SIGN ( 2c). (0 ) DC ICAM - 3 2 L19  
 . .  
 ( 2c), 2 (LFA - 1) . DC - SIGN  
 1 ( 2b), ICAM - 3 2 (LFA - 1) DC - SIGN ( 2c)

. ICAM - 3 가 2 7 , DC - SIGN  
     , 2 ( 2c) , 2 , DC - SIGN  
 . DC - SIGN 7 ( 2c). DC - SIGN

, DC - SIGN 가 ICAM - 3 가  
 ( 2a 2b). AZN - D1 AZN - D2 DC - SIGN DC  
 ICAM - 3 .

3 : DC - SIGN 44kDa

DC - SIGN , DC - SIGN <sup>125</sup>I - DC  
 SDS - PAGE 44kDa ( 3a, 1 2). 3  
 DC - SIGN HIV gp120 C  
 3a : DC - SIGN 44kDa DC I <sup>125</sup> , DC - SIGN DC - SIGN  
 AZN - D1( 1), AZN - D2( 2) AZN - L19( 2 ; 3) . SDS - P  
 AGE (5 15% )  
 2 LFA - 1( L, 180 kDa), MAC - 1( M, 165 kDa) p150, p95( X, 150 kDa),  
 (95 kDa) DC - SIGN (44 kDa) 3

3b : DC DC - SIGN 2 ( gp120 C DC -  
 SIGN : 296 306 187 197 ) Edman  
 gp120 C cDNA DC - SIGN DC  
 7 8 (R1 R8) ,  
 .

SDS - PAGE , DC - SIGN DC 44 kDa  
 ( ). , ICAM - 3 - Fc ICAM - 3 - Fc  
 , DC - SIGN DC - SIGN 44 kDa  
 DC ( ). , ICAM - 3 Fc DC 44 kDa  
 DC - SIGN ICAM - 3 가 , DC  
 FA - 1 DB 가 ( 3a), ICAM - 3 L  
 (Vandervieren , Immunity.3, 683 - 690, 1995). DC - SIGN LFA - 1 가  
 ( 3a, 1 3), LFA - 1 DC - SIGN DC  
 . , DC - SIGN 44 kDa DC ICAM  
 - 3 1 .

4 : DC - SIGN HIV gp120 C

DC - SIGN , DC - SIGN AZN - D1 DC SDS - PA  
 GE 44 kDa HPLC . Edman 2 (0.5 1 pmol)

I) 11 ( 3b)  
 EMBL HIV  
 gp120 C 100% (HIV) gp120 (Curtis CD4 , 1992).

gp120 C DC RNA RT - PCR  
 , 1237 nt PCR DC PCR  
 gp120 100% ( 3b). gp120 C  
 cDNA COS7 , gp120 C  
 DC - SIGN ( 4a). 4 COS7 DC - SI  
 GN DC - SIGN AZN - D1 ICAM - 3 .  
 .

4a : AZN - D1 DC - SIGN cDNA COS7 ( ), ( ).  
 COS7 ( ) . AZN - D2 ( ).

4b : ICAM - 3 COS7 - DC - SIGN . COS7 , 1a

EGTA(5 mM) DC - SIGN(AZN - D1), ICAM - 3 (CBR - IC3/1, CBR - IC3/2) 2 (AZN - L19)  
 ICAM - 3 COS7 - DC - SIGN . 3 가  
 (SD < 5%).

COS7 30% 가 DC - SIGN DC - SIGN  
 , COS7 - DC - SIGN ICAM - 3 COS7  
 ICAM - 3 ( 4b). COS7 DC - SIGN ICAM - 3 DC - SIG  
 N , EGTA 가 Ca<sup>2+</sup>  
 ( 4b).

DC(DC - SIGN) ICAM - 3 HIV gp120 C (Curtis , 1992),  
 3 404 a.a 7 1  
 C (Cys253 - Ala404) Ca<sup>2+</sup> (C )  
 ( 3b).

5 : DC - SIGN DC

AZN - D1/D2 ( 1). DC DC - S  
 IGN , RT - PCR DC - SIGN mRNA DC - SIGN  
 DC

DC - SIGN , 2 DC - SIGN

DC - SIGN , T DC - SI  
 ( 5). 5 DC - SIGN :

GN (OM × 100). . AZ  
 N - D1 DC - SIGN . (GC), T (T) B (B)

, DC - SIGN  
 ( 5). CD3 CD14 DC - SIGN  
 RT - PCR , T ( )  
 ( 1).

6 : DC - SIGN/ICAM - 3 DC - T

DC - SIGN ICAM - 3 DC 가 , ICAM - 3  
 cDNA K562가 DC 6a ,  
 DC - SIGN , DC DC - SIGN , K562 - ICAM - 3  
 DC K562 , , ICAM - 3 가 DC - SI  
 GN 6 DC ICAM - 3 DC - SIGN DC - T

6a : DC ICAM - 3 DC - SIGN D  
 DC - SIGN AZN - D1 (-) (-) DC 가 ICAM - 3  
 , 2 가 (SD < 5%).

6b : K562 - ICAM - 3 DC ICAM - 3 cDNA  
 K562 K562 (K562 - ICAM - 3) HE( ) DC  
 SFDA 37 DC - SIGN , K562 K562 - I  
 CAM - 3 DC( $50 \times 10^3$  cells/cell type)  
 (0.5%) 2 가

6c : PbL DC DC - SIGN DC(50 × 10<sup>3</sup> cells)  
 10 DC - SIGN AZN - D1 AZN - D2( $10 \mu\text{g}/\text{M}\ell$ )  
 Calcein - A PBL ( $1 \times 10^6$  cells)( $25 \mu\text{g}/10^7$  cells/ $\text{M}\ell$ , 37 30 )  
 가 37  
 2 가

6d : DC - SIGN - ICAM - 3 DC T T (10  
 $0 \times 10^3$  LFA - 3(TS2/9) DC - SIGN (AZN - D1, AZN - D2) (20  $\mu\text{g}/\text{M}\ell$ ) DC -  
 $(1.5 \times 10^3)$  가 4 4 6  
 $[^3\text{H}]$  - 가 3 CPM %

DC - SIGN , DC - SIGN - ICAM - 3 DC 가  
 , 60 TCR DC

DC T 가 DC - SIGN  
 DC T (DC:T , 1:20)

DC - T . . 6b , DC T  
 20 ( 50%), (32%) DC - T DC - SIGN  
 /ICAM - 3 , DC - T , DC - SIGN  
 A K562 , ICAM - 3 , ICAM - 3 cDNA , DC - SIGN cDN  
 , DC - SIGN K562

7 : DC T DC - SIGN

ICAM - 3 DC - SIGN 가 DC T , DC T  
 DC - SIGN DC 6c , DC - SIGN DC - SIGN  
 T 75% , T CD2 AZN - D1 AZN - D2  
 LFA - 3 T T LFA - 3 DC - SIG  
 N T ( 6c).

8 : DC - SIGN DC HIV - 1

DC - SIGN HIV gp120 , HIV - 1 DC 가 (Blauvelt  
 DC, DC - SIGN , , , ,  
 DC - T , J. Exp. Med. 184(6), 2433 - 8, 1996) DC - SIGN 7  
 , 1997 ; Granelli - Piperno ( 7), IL - 4 GM - CSF DC , ,  
 , DC 2 HIV - 1 가 PBMC 3 HIV - 1  
 . DC - T p24 Gag ( 7) , [Blauvelt , 1997 ; Gr  
 , p24 , , , ,  
 , anelli - Piperno , 1998 ; Granelli - Piperno , Curr Biol.9(1), 21 - 29, 1999]. , HIV - 1  
 가 DC DC - SIGN PBMC  
 , HIV - 1 75 % , T 3 5  
 ( 7). HIV - 1 가 DC - SIGN DC , DC - T HI  
 V - 1 ( 7). DC - SIGN DC - SIGN HIV - 1 DC T  
 , DC HIV - 1 , , , , DC - SI  
 GN DC HIV - 1 , ICAM - 3 HIV - 1 DC

, DC T  
 DC TCR T , DC  
 , DC - SIGN ICAM - 3 DC - T , DC  
 . DC - SIGN DC , DC - SIGN DC HIV - 1  
 . HIV - 1 DC , DC - SIGN DC HIV - 1  
 IV - 1 T [Cameron CD4+ T H  
 (1), 61 - 71, 1994]. DC HIV - 1 , 1992 ; Cameron: AIDS Res Hum Retroviruses 10  
 [Granelli - Piperno , 1998]. IL - 4 GM - CSF , DC

DC M HIV - 1 [Granelli - Piperno, 1996; Granelli -  
 Piperno, 1998] [Blaauvelt, 1997], DC M T HIV - 1  
 [Granelli, 1998, 1999].  
 T HIV - 1  
 HIV - 1  
 gp120 CD4 , CD4  
 , CCR5 CXCR4  
 [Littman, 1998] [Dragic, 1996 ; Feng, 1996], Sci  
 ence 272(5263), 872 - 7, 1996]. DC CD4  
 DC HIV - 1  
 DC . DC CD4 , HIV - 1 DC 가 , HIV - 1 DC 가  
 HIV - 1 T DC CD4 DC DC SIGN DC SIGN  
 DC DC SIGN DC DC SIGN  
 DC SIGN DC DC CD4  
 DC T DC CD4 CD4 가  
 DC T HIV - 1  
 [Tsunetsugu - Yokota, 1997]. DC - T DC - SIGN DC SIGN  
 HIV - 1 가 , DC - SIGN HIV - 1 T DC SIGN  
 HIV - 1 DC DC SIGN DC SIGN  
 C가 DC - SIGN CD4 ( 7) HIV - 1 DC  
 V - 1 DC - SIGN 가 HIV - 1 DC  
 , DC - SIGN HIV - 1  
 , DC - SIGN HIV - 1

DC 가 DC [Hock , Immunol. 83, 573 - 581, 1994], [de Saint - Vis , Immunity 9(3), 325 - 36, 1998], [Hart , 1997]. DC DC - SIGN mRNA DC - SIGN 가 ( 1 ).  
 DC - SIGN T DC DC - SIGN DC - T  
 DC T . , DC - SIGN DC - T

Figure 387, 713 - 717, 1997]. 2  
 가 TCR 가 APC MHC - DC TCR  
 TCR T DC MHC - DC CK1 DC T-  
 DC T , T , MHC [ Adema , Nat  
 , T , MHC , TCR

: KIM185( 2 ; Andrew , Eur.J.Immunol. 23, 2217 - 2222, 1993), AZN - L19( 2 ), NKI - L15( L ; Keizer , Eur.J.Immunol. 15, 1142 - 1147, 1985), A B2( 1 ; Da Silva , J.Immunol. 143, 617 - 622, 1989), CBR - IC3/1 CBR - IC3/2 ( ICAM - 3 ; de Fougerolles , J.Exp.Med. 177, 1187 - 1192, 1993), CD14 (WT14), CD4(WT4). BALB/c DC ICAM - 3 DC가 DC - SIGN AZN - D1 AZN - D2

9B :

Sallusto Lanzavecchia, J.Exp.Med. 179, 1109 - 1118 ; Romani , J.Exp.Med. 180, 83 - 93, 1994  
DC DC PBMC  
IL - 4 7 (Schering - Plough, Brussels, Belgium ; 1000  
U/Ml). 4 가 . 7 MHC, CD1a, p150,95 CD80  
DC (Lub , Mol.Biol.Cell  
8, 719 - 728, 1997) K562 10 µg PCR ICAM - 3 R1 (Dr. D. Simmons ) 2 µg PG  
K - hyg (te Riele , 1990) ICAM - 3 K562 (K562 - ICAM

- 3) Figdor , J.Immunol.Methods 68, 73 - 87, 1984 , PBMC  
T (> 90% CD3 )

9C :

(Pink Ziegler, 1979 ; Research Methods in immunology, L. Lefkovits B.Pernis ed s.(New York : Academic Press), pp169 - 180) Na <sup>125</sup>I(Amersham, Buckinghamshire, UK)  
(1% NP - 40, 50 mM (pH 7.8), 150 mM NaCl, 1 mM CaCl<sub>2</sub>, 1 mM Mg Cl<sub>2</sub>, 1 mM PMSF, 0.02 mg/Ml ) DC 4 , 1 . 4 15 13,000 g 3 Protein A -  
CL - 4B (Pharmacia Fine Chemicals, Piscataway, NJ) mAb  
Lammlie (Lammlie, Nature 227, 680 - 685, 1970) ,  
SDS - PAGE ,  
Eurosequence BV(Groningen, The Netherlands)

9D : DC - SIGN cDNA

RNA (Chomczynski Sacchi, Anal Biochem 162  
(1), 156 - 9, 1987) gp120 C cDNA RT - PCR DC  
RNA PCR gp120 C (5' - - > 3') ( M  
98457)(Curtis , 1992) XF29, AGAGTGGGTGACATGAGTG ; XR1265, GAAGTTCTGCTACGCAGGAG. PCR pGEM - T ( Promega, Madison WI) gp  
120 C (Curtis , 1992). cDNA  
pRc/CMV(pRc/CMV - DC - SIGN) , DEAE COS7 pRc/CMV - D  
C - SIGN [Seed and Aruffo, Proc.Natl.Acad.Sci.U.S.A 84, 3365 - 3369, 1987].

9E :

TransFluorSpheres(488/645 nm, 1.0 µg ; Molecular Probes, Eugene, O R) ICAM - 1 Fc ICAM - 3 Fc [Geijtenbeek , 1999 ]. , 20 µl (50 mM MES 5 mg/Ml) 50 µl TransFluorSpheres 가 . 30 µl EDAC(1.33 mg /Ml) 가 100 mM PBS(50 mM , 0.9% NaCl pH 7.4)  
3 150 µl PBS, 0.5% BSA (w/v) (15 µl) 0.5 Ml PBA PB  
Fc Fab (6 µg/Ml) 37 2 S, 0.5% BSA 4 0.5 Ml IgG1 Fc (ICAM - 1 Fc, VCAM - 1 Fc ; 250 ng/Ml)  
, 100 µl PBS, 0.5% BSA  
4 . ICAM - 1 Fc ICAM - 3 Fc IgG1 Fc (Dr.D.Simmons ) Geijtenbeek ( )  
Tris - - BSA (20 mM Tris - HCl pH 8.0, 150 mM NaCl, 1mM CaCl<sub>2</sub>, 2 mM MgCl<sub>2</sub>, 0.5% BSA ; 5 × 10<sup>6</sup> cells/Ml) . 50,000 mAb (20 µg/Ml) / ,  
10 , 96 V (20 beads/cell)  
/ 가 37 30 100 µl TSA % FACScan(Becton and Dickinson & Co., Oxfo rd, CA) 가

9F :

DC ICAM - 3 . DC ICAM - 3  
 $(2 \times 10^6 \text{ cells/M}\ell)$  37 . (Molecular Probes, Eugene, OR ; 50  $\mu\text{g/M}\ell$ ) . DC ICAM - 3  
(Molecular Probes, Eugene, OR ; 40  $\mu\text{g/M}\ell$ ) 1 . , DC ICAM - 3  
 $(50 \times 10^3 \text{ cells/M}\ell)$  37 .  
(0.5%) , FACScan(Becton and Dickinson & Co., Oxford, CA)

DC T . DC( $50 \times 10^3 \text{ cells}$ ) 10  
, DC - SIGN AZN - D1 AZN - D2 (10 g/M $\ell$ ) / .  
Calcein - A (Molecular Probes, Eugene, OR ; 25  $\mu\text{g/10}^7 \text{ cells/M}\ell$ , 30 , 37 ) PBL(1  
 $\times 10^6 \text{ cells}$ ) 가 37 .  
Oxford, CA) T DC %  
가 .

9G : DC T  
(20 g/M $\ell$ ) T ( $100 \times 10^3$ ) DC (1.5  $\times 10^3$ ) 가 .  
4 4 16 [  $^3\text{H}$  ] - (1.52 TBq/mmol,  
0.5  $\mu\text{Ci/well}$  ; Amersham, Buckinghamshire, UK) 가 .

9H : DC HIV - 1

HIV - 1<sub>Ba-L</sub> (MDM) MDM  
7 , p24 ELISA(Diagnostics Pasteur, Marnes la Coquette, France) TCID<sub>50</sub>  
,  $10^4 \text{ /M}\ell$  20 , DC( $50 \times 10^3$ ) 2  
HIV - 1<sub>Ba-L</sub> 가 ( $10^5$  10 $^3$  ) , PHA/IL - 2 PBMC  
( $50 \times 10^3$ ) . DC - T 3 5 p24 ELISA  
(Diagnostics Pasteur, Marnes la Coquette, France) p24 IL - 2(10 U/M $\ell$ ) PH  
A(10  $\mu\text{g/M}\ell$ ) PBMC .

9I :

1 (8  $\mu\text{m}$ ) 100% (10 ) , PBS , , 60  
(10  $\mu\text{g/M}\ell$ ) , ABC - AP Vectastain (V  
ector Laboratories, Burlingame, CA)

[ 1]

	DC - SIGN *	DC - SIGN mRNA ‡
	-	-
7 DC	+++	+
PBL	-	-
T	-	-
B	-	-
B ( )\$	-	n.d.
	-	-
CD34 <sup>+</sup>	-	n.d.
PBMC( #)	-	-
T †	-	-
† †	-	-

\* : . - < 20 , + ++ > 400 (AZN - D1 )

‡ : RNA DC - SIGN XF29 XR1265 RT - PCR

\$ :

# : 2 PHA(10 µg/Mℓ) IL - 2(10 U/Mℓ) .

† : T : HSB, PEER, CEM Jurkat

† † : : THP - 1, MM6 U937

n.d. :

(57)

1.

, C , , C DEC - 205 , 가

2.

1 , T , ,

3.

1 / 2 , T , ICAM - 2 , ICAM - 3 C , T ,

4.

1 3 , , , , , , 가 /

5.

HIV , HIV ( , gp120)  
 HIV , C

6.

T HIV가 ,  
 C

7.

, , , 가 /  
 ,

1) C ;

2)

8.

7 , C

9.

7 8 ,  
 ,

10.

1 9 , C ; D -  
 A ; N - ; L - ;  
 ; D - C ; gp120 ;  
 ;

11.

1 9 , C 1 가 ,

12.

10        11            ,            C  
   1                      가     C  
   ;   /                    ,

13.

1        9            가     C  
   ;   /                    ,

14.

13            ,            가 AZN - D1        AZN - D2

15.

13        14            가            ,            ,            /

16.

1)            C            ;  
 2)

17.

16            ,            C

18.

16        17            ,  
                   ,

19.

16        18            ,            C            ;            D  
   A            ; N -        ; L -        ;            A  
   ;            ;            - D -        ; gp120  
                   C            ,

20.

,    13            14

21.

, / , 13 14

22.

3 14 ; / , , 1 . 가 C / , 1

23.

a) 13 14 ;

b) 가 ; 가 ,

c) , , / .

24.

23 , 가 , , / .

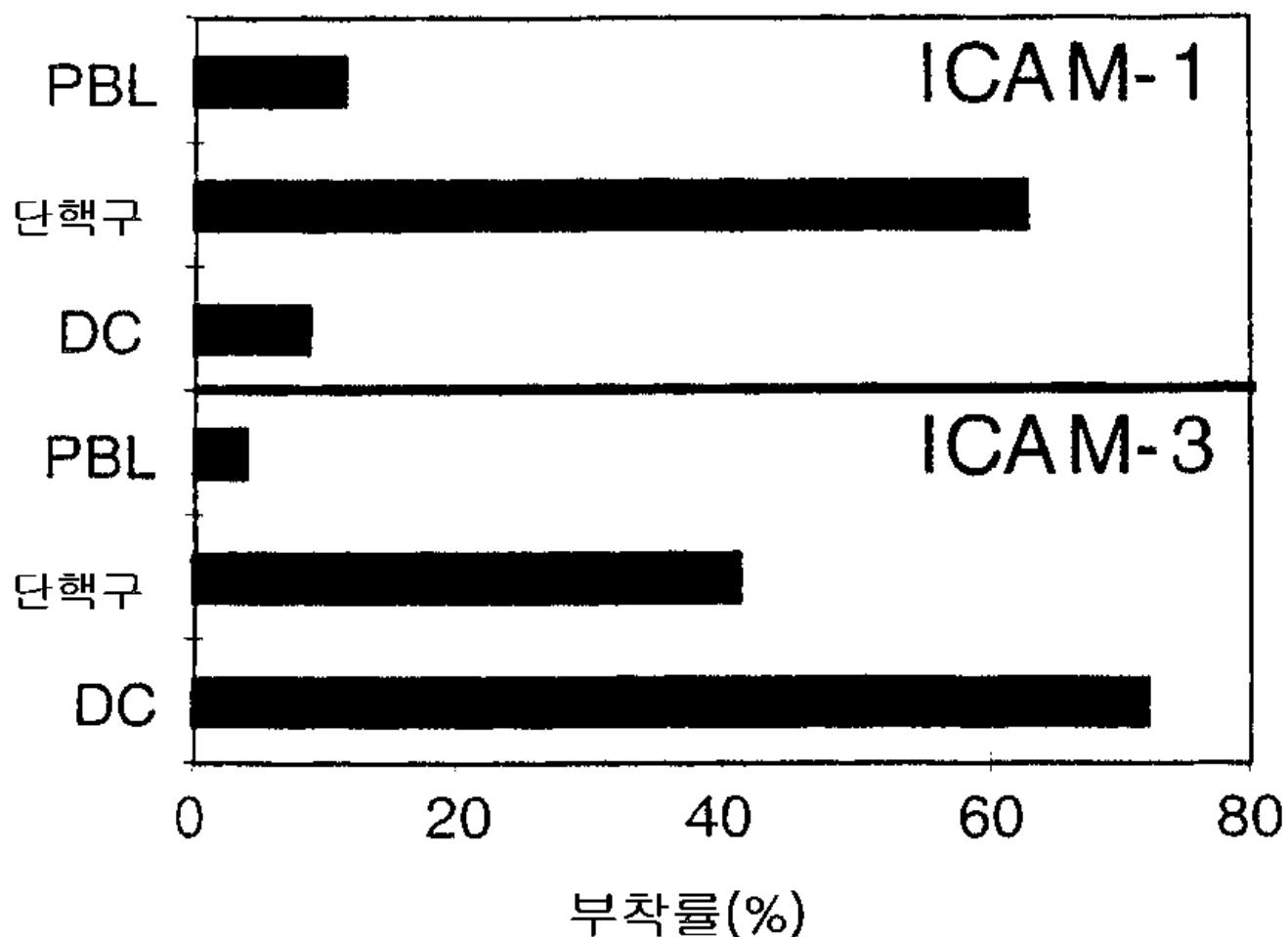
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23 24 , 가 , , / .

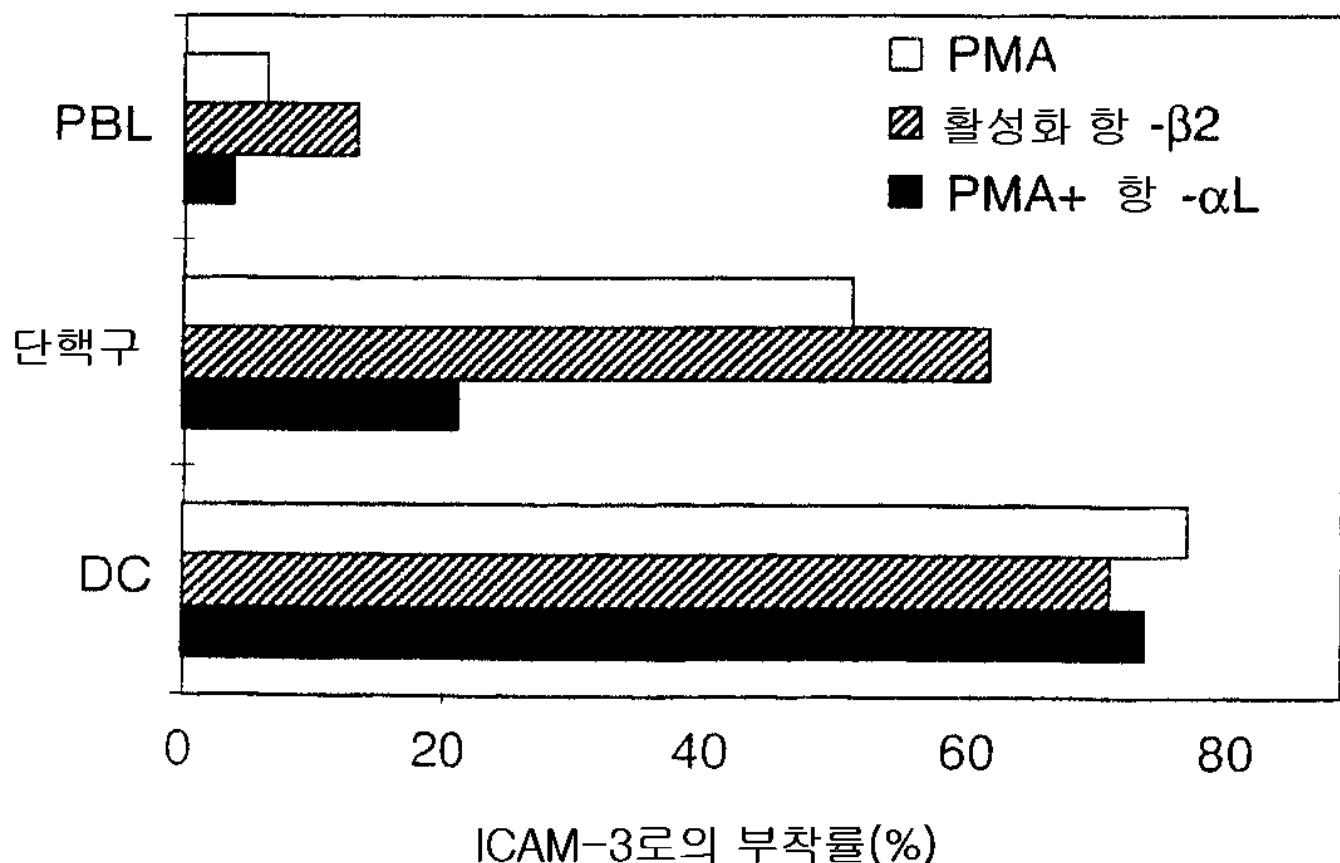
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23 24 .

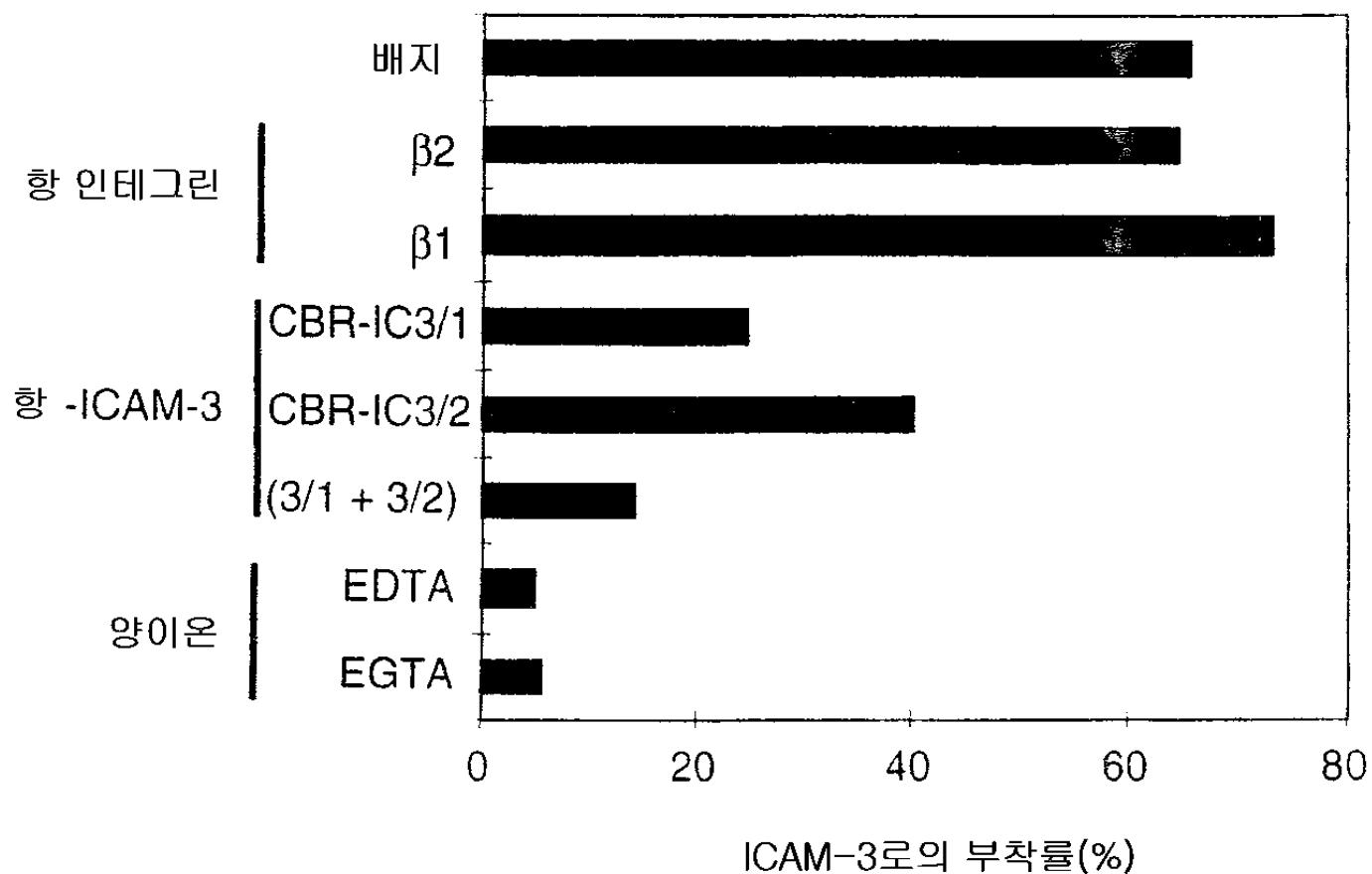
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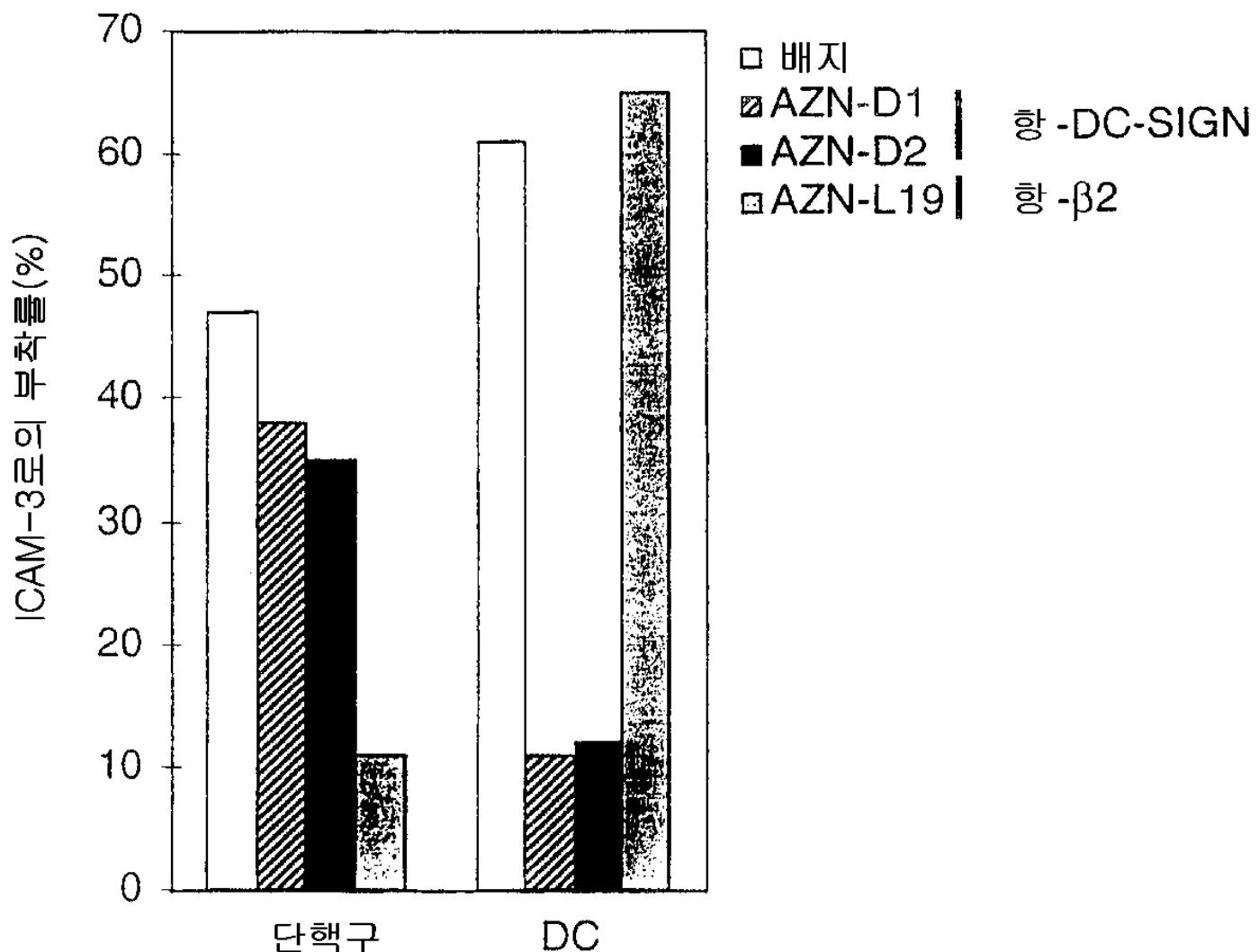
1b



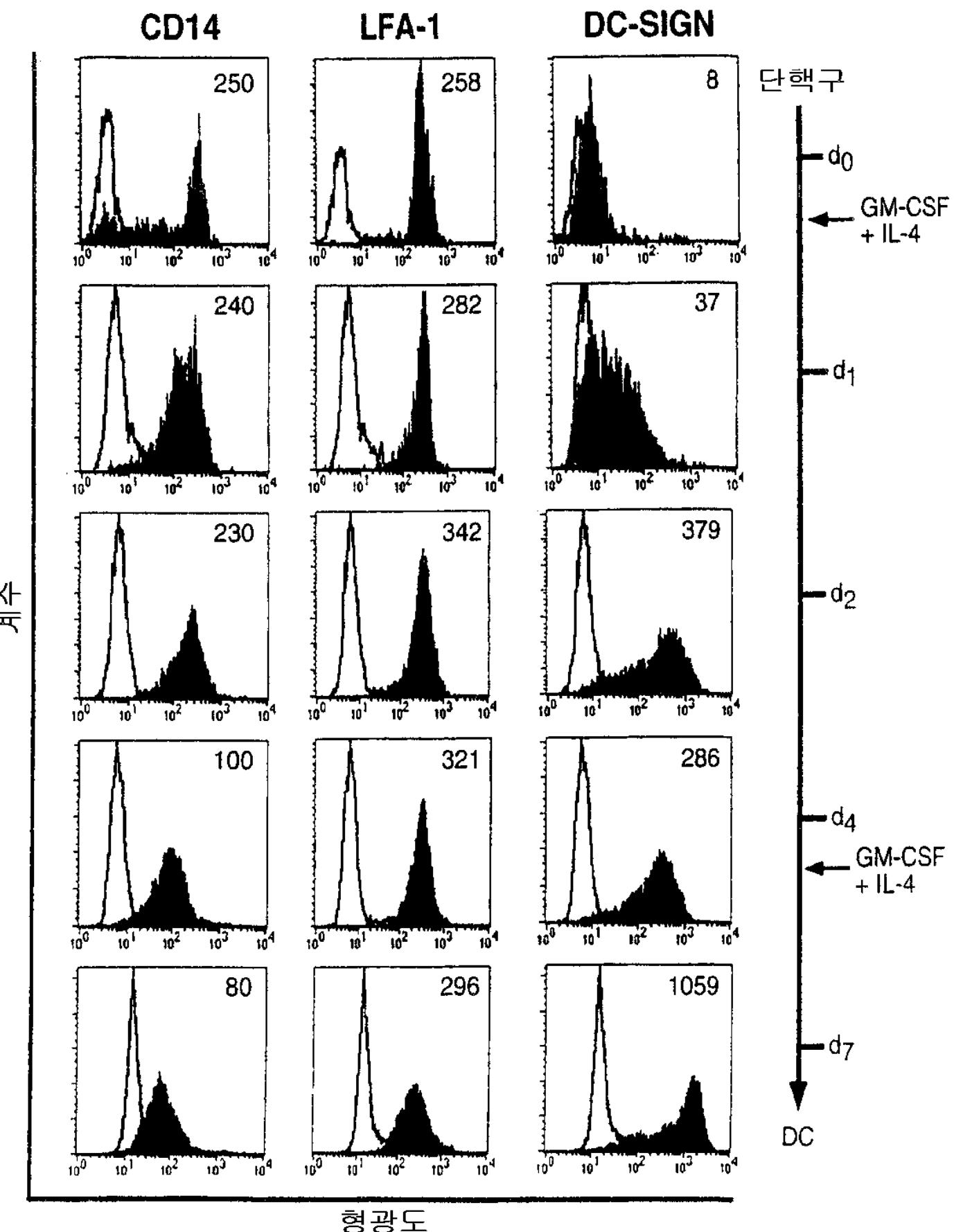
1c



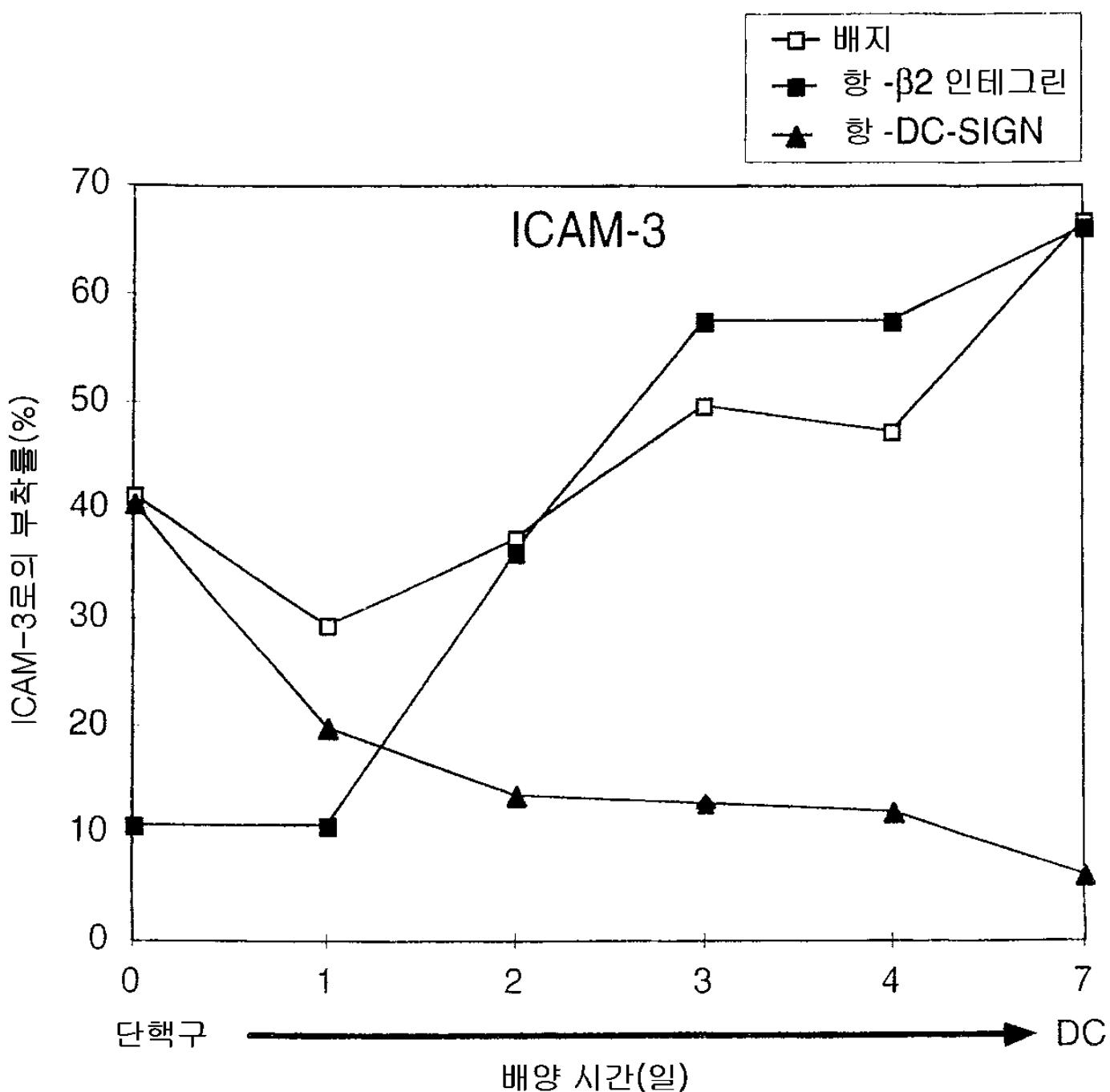
2a



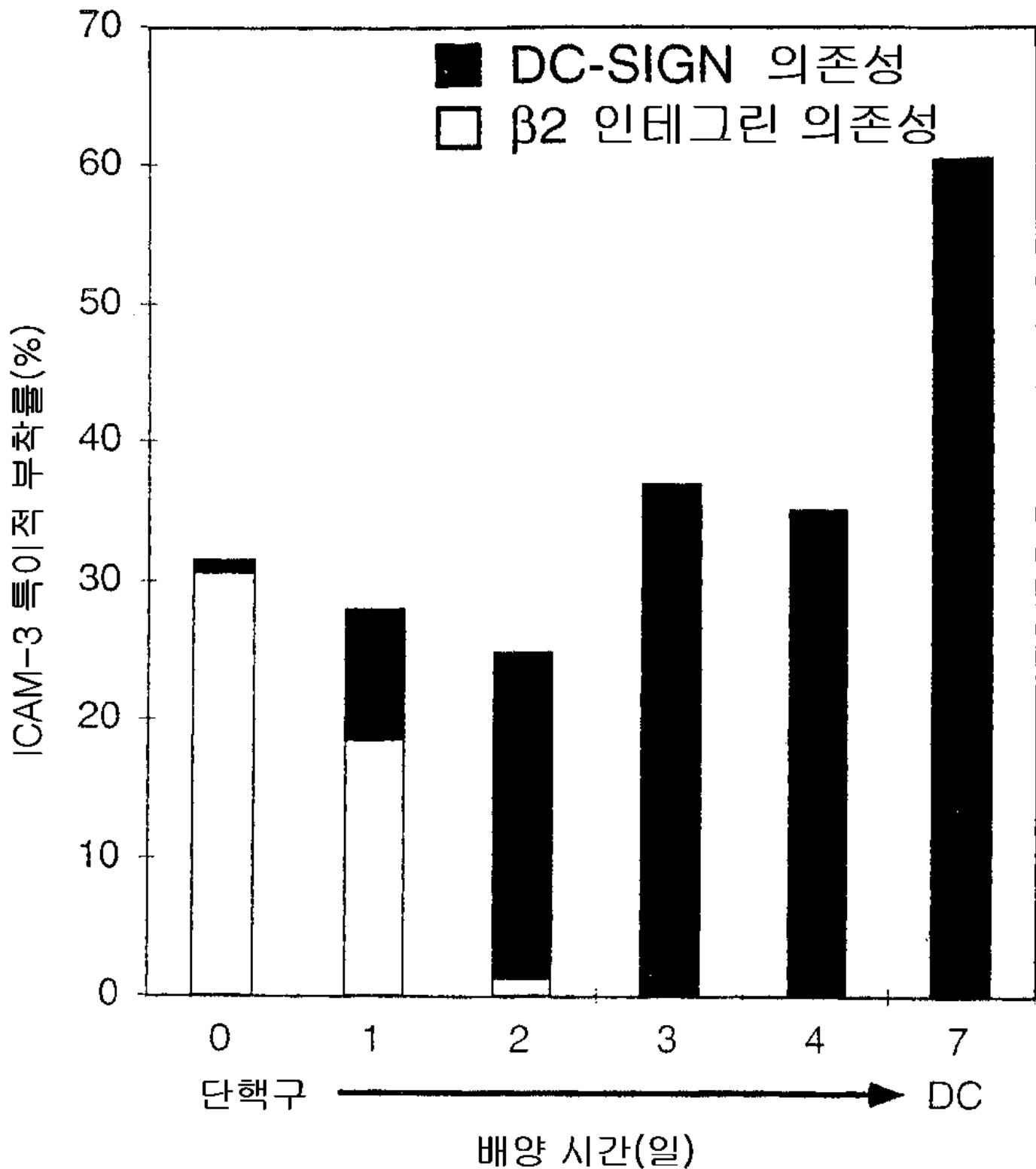
2b



2c



2d



3a

kDa      1      2      3

205 —

►  $\alpha L$

►  $\alpha M$   $\alpha X$

►  $\beta 2$

126 —

89 —

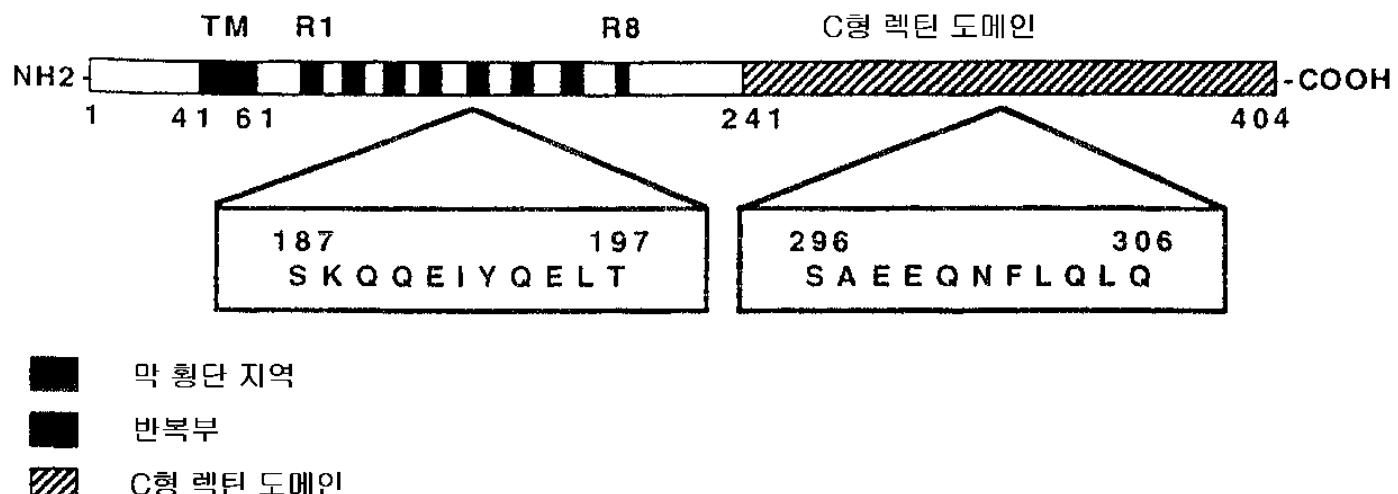
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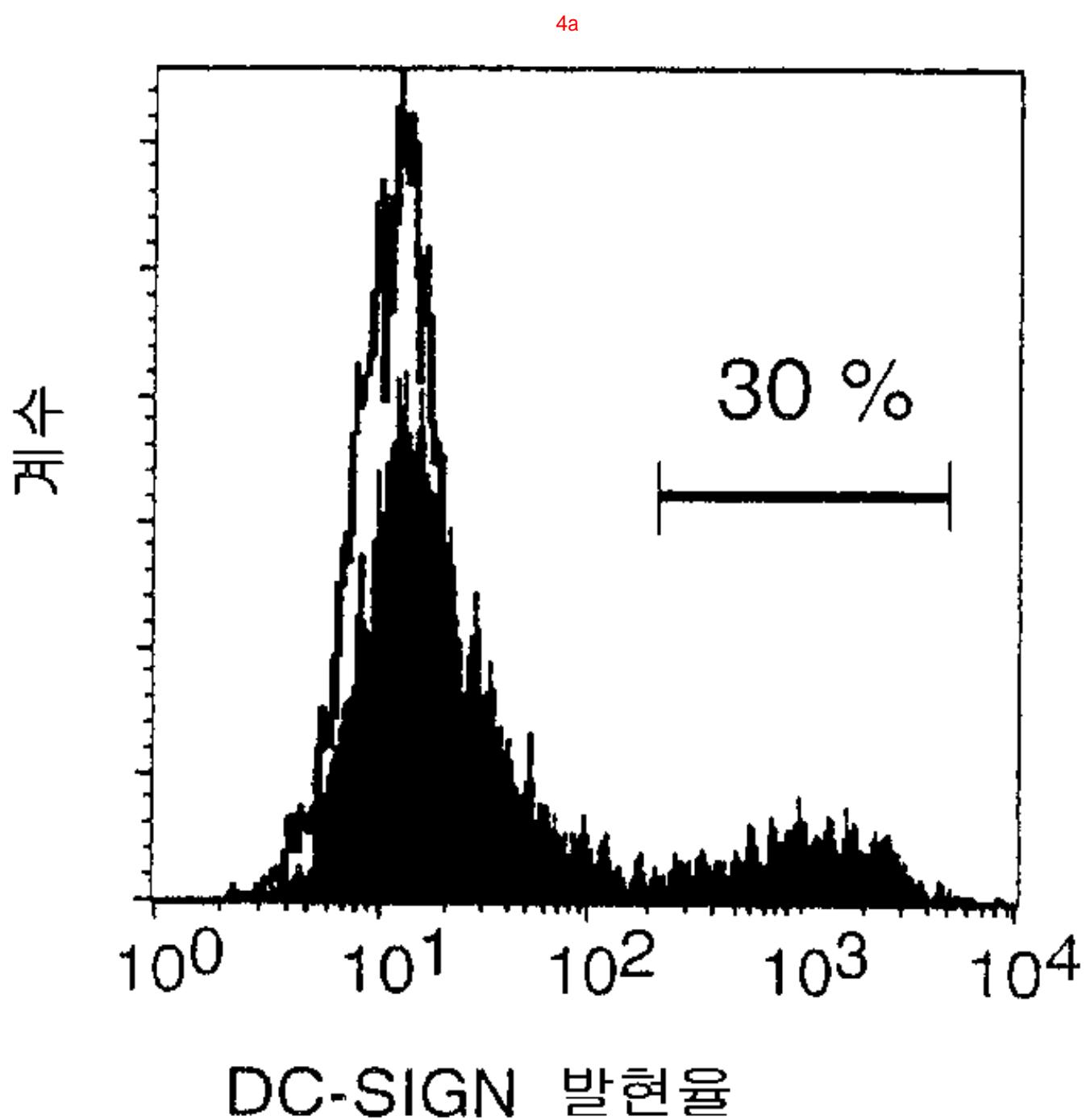
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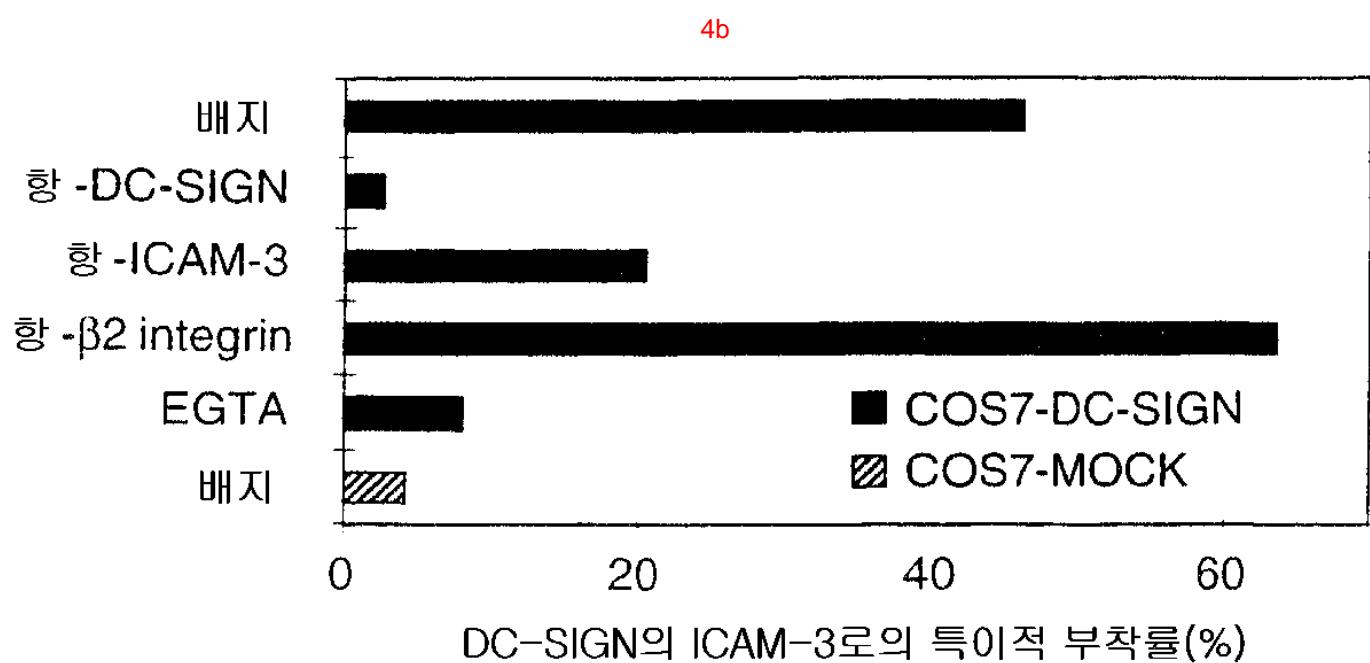
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20 —

3b

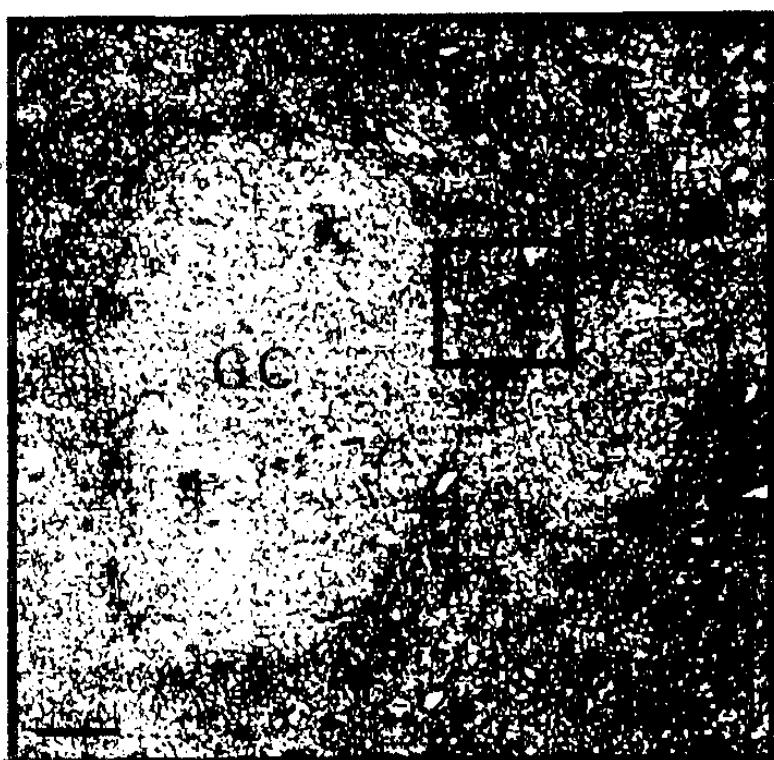




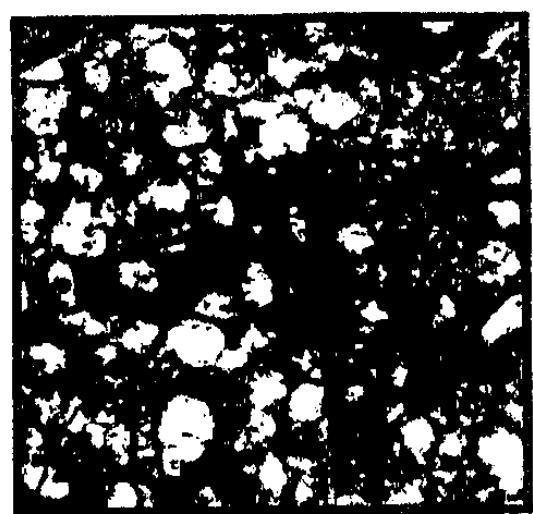


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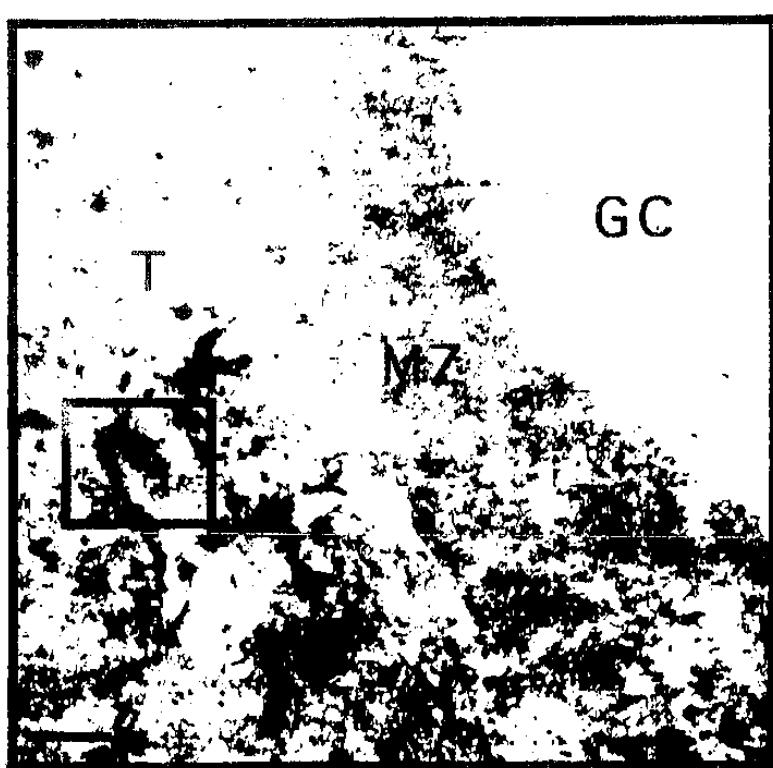
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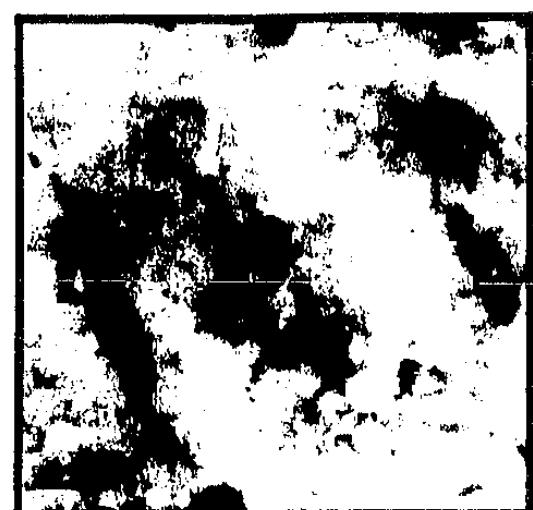
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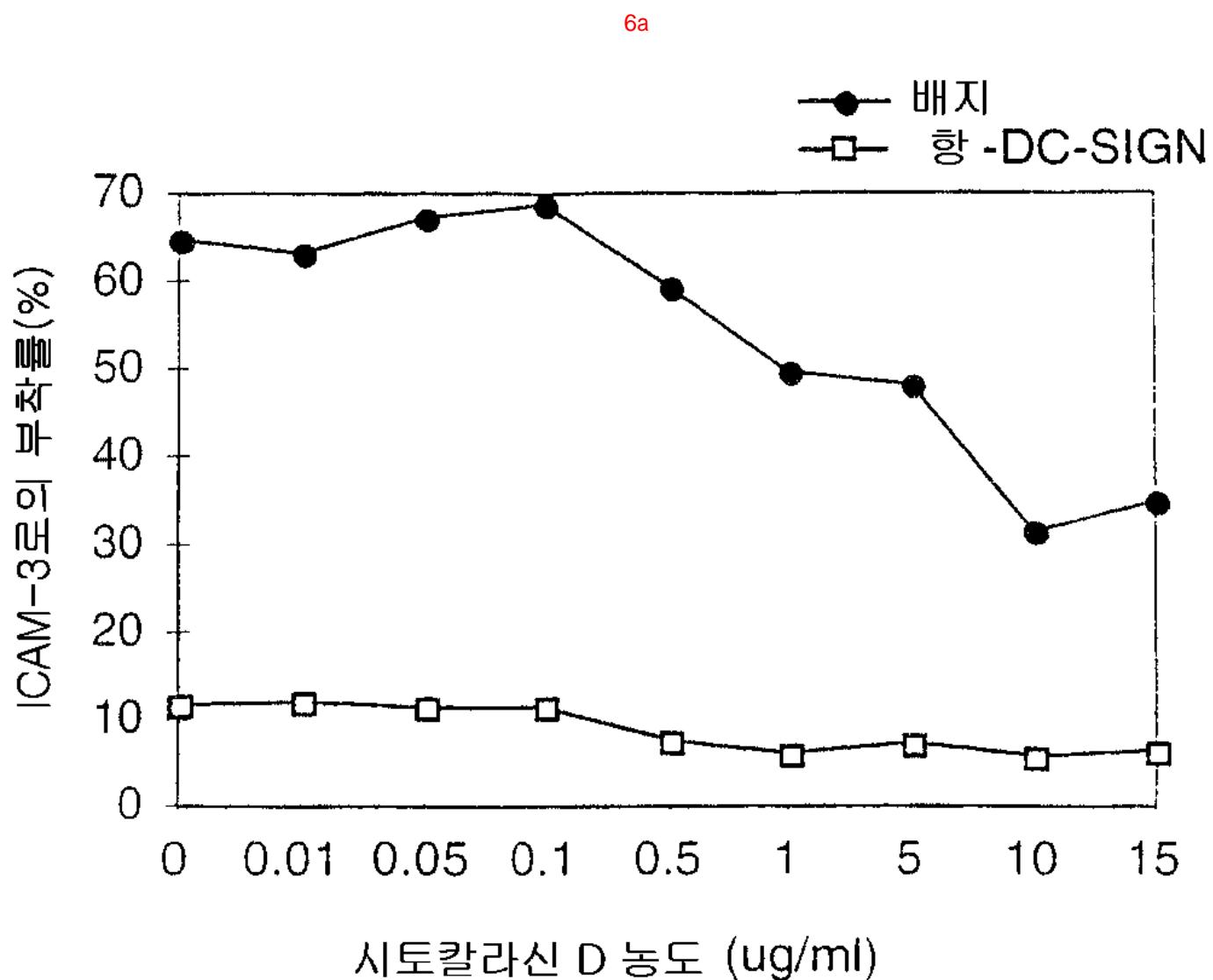


C

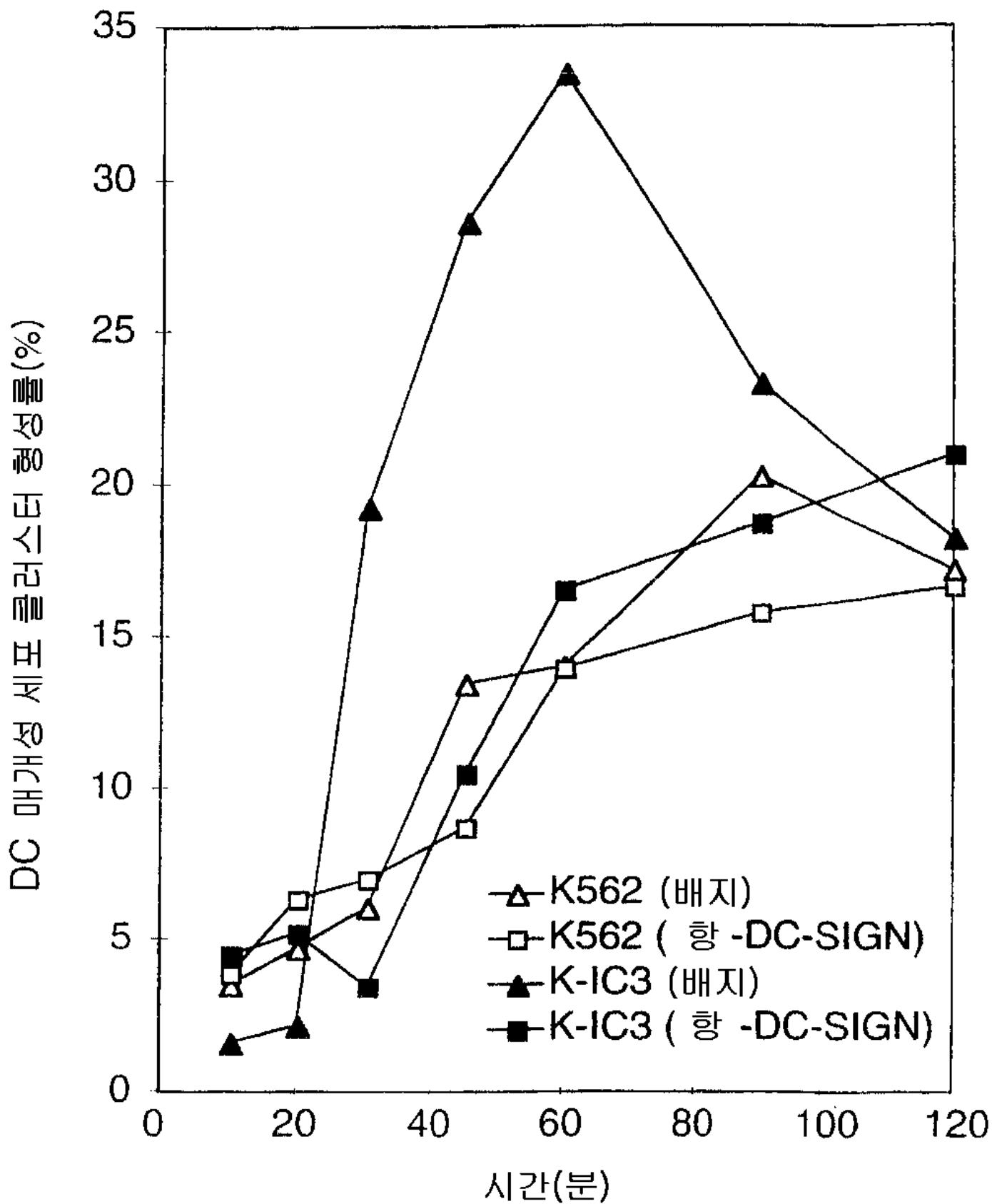


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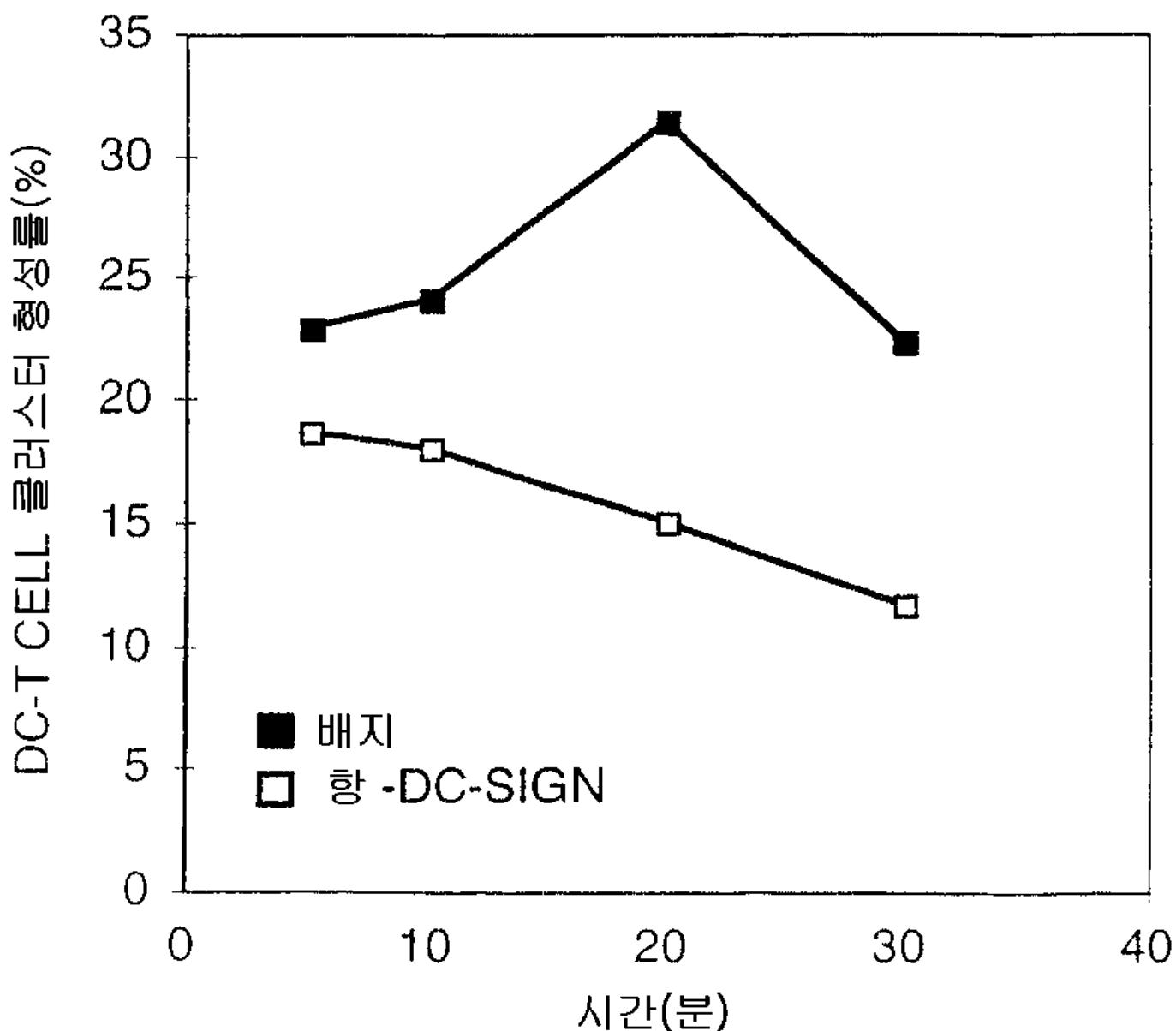




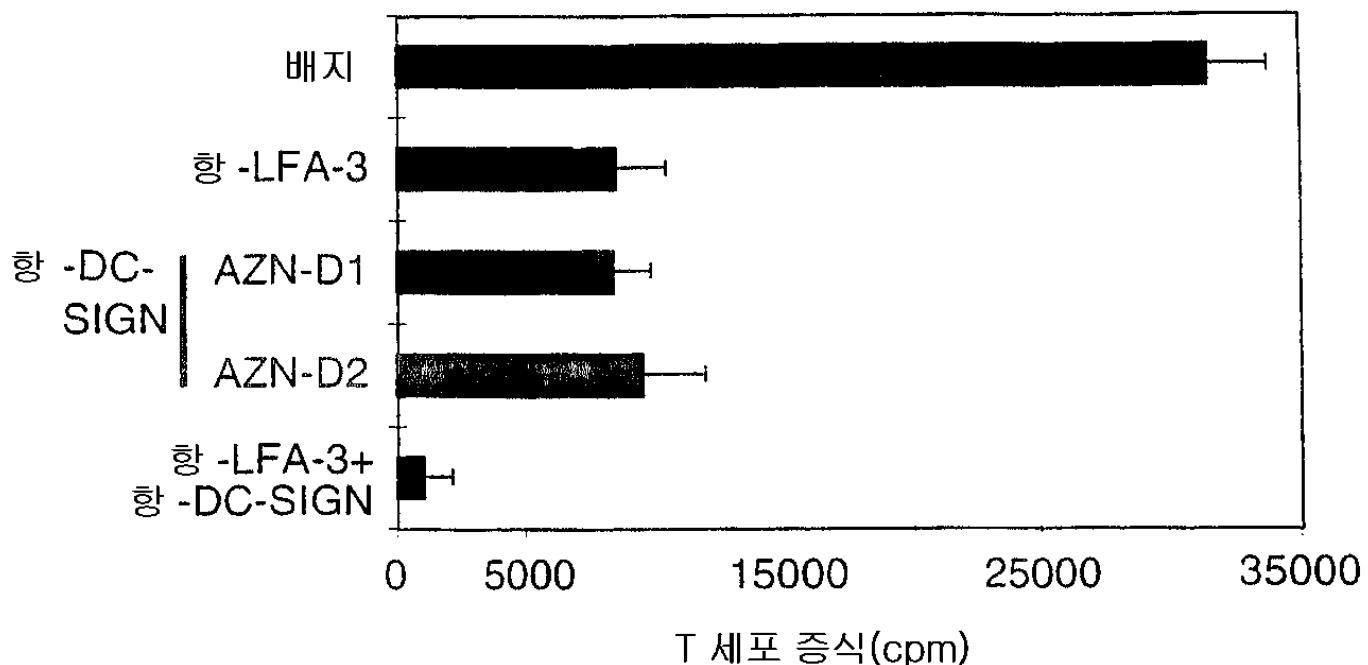
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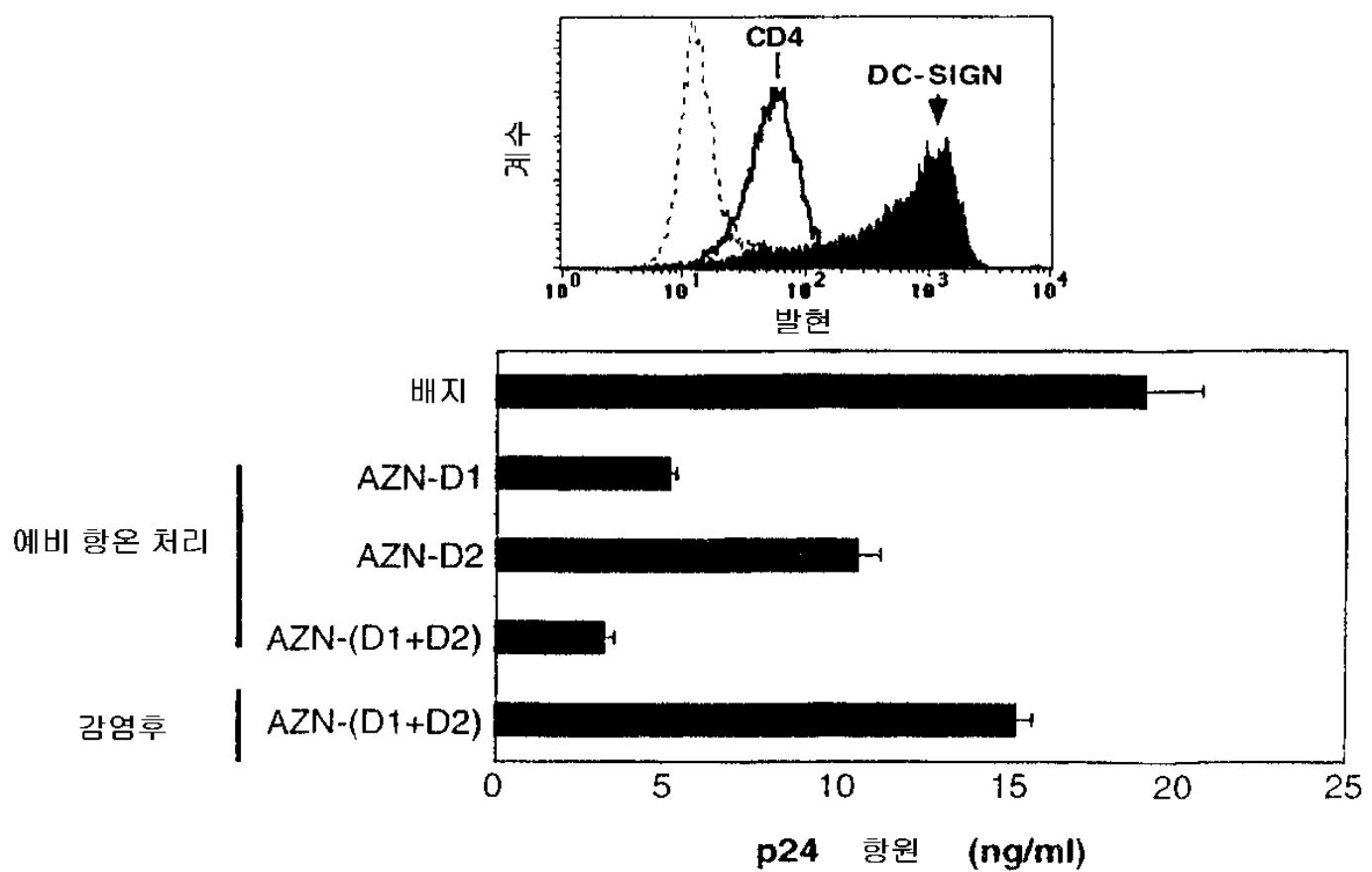
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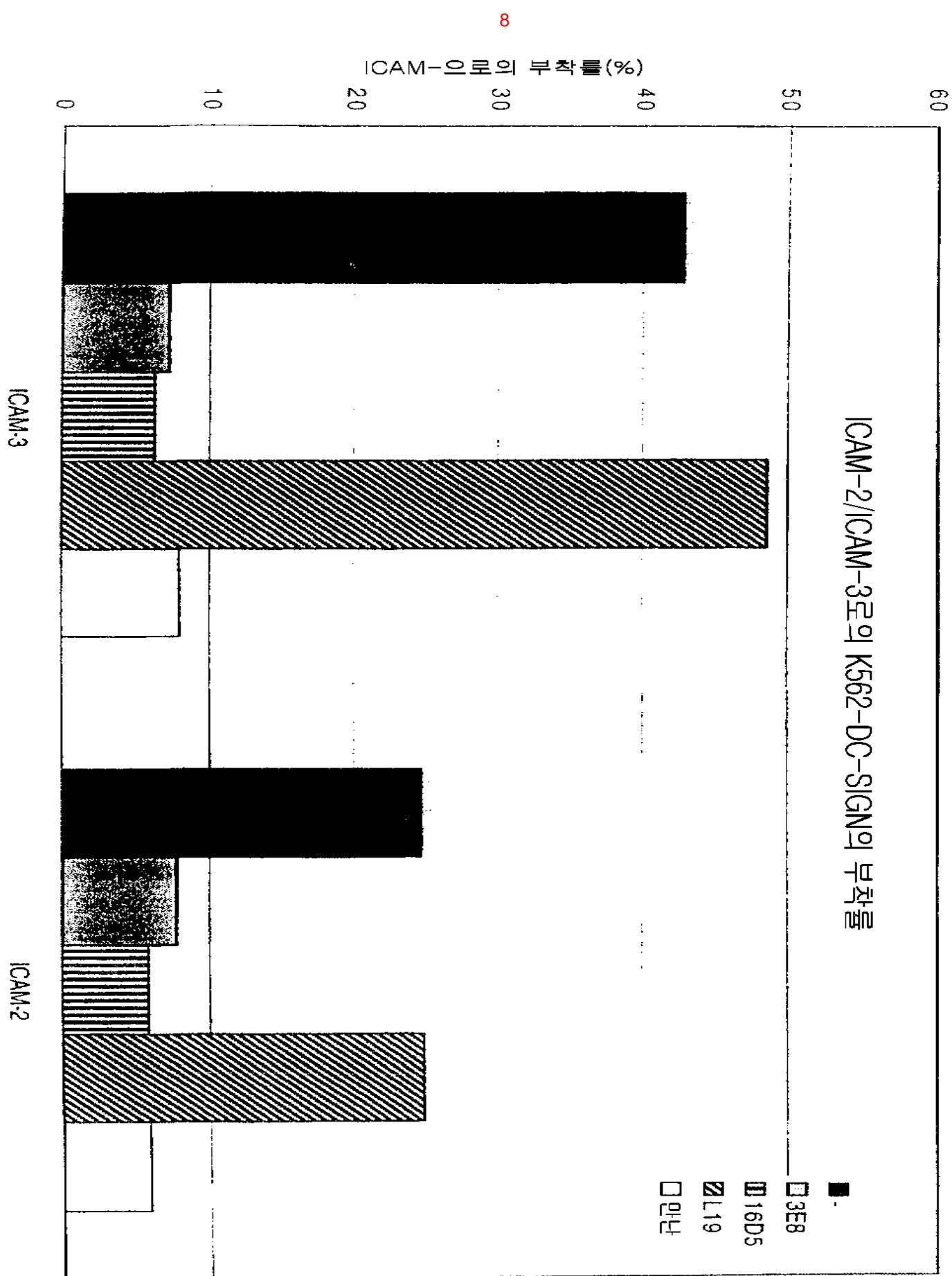


6d



7





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Met ser asp ser lys glu pro arg leu gln gln leu gly leu leu glu glu gln leu  
61/21 91/31  
AGA GGC CTT GGA TTC CGA CAG ACT CGA GGA TAC AAG AGC TTA GCA GGG TGT CTT GGC CAT  
arg gly leu gly phe arg gln thr arg gly tyr lys ser leu ala gly cys leu gly his  
121/41 151/51  
GGT CCC CTG GTG CTG CAA CTC CTC TCC TTC ACG CTC TTG GCT GGG CTC CTT GTC CAA GTG  
gly pro leu val leu gln leu leu ser phe thr leu leu ala gly leu leu val gln val  
181/61 211/71  
TCC AAG GTC CCC AGC TCC ATA AGT CAG GAA CAA TCC AGG CAA GAC GCG ATC TAC CAG AAC  
ser lys val pro ser ser ile ser gln glu gln ser arg gln asp ala ile tyr gln asn  
241/81 271/91  
CTG ACC CAG CTT AAA GCT GCA GTG GGT GAG CTC TCA GAG AAA TCC AAG CTG CAG GAG ATC  
leu thr gln leu lys ala ala val gly glu leu ser glu lys ser lys leu gln glu ile  
301/101 331/111  
TAC CAG GAG CTG ACC CAG CTG AAG GCT GCA GTG GGT GAG CTT CCA GAG AAA TCT AAG CTG  
tyr gln glu leu thr gln leu lys ala ala val gly glu leu pro glu lys ser lys leu  
361/121 391/131  
CAG GAG ATC TAC CAG GAG CTG ACC CGG CTG AAG GCT GCA GTG GGT GAG CTT CCA GAG AAA  
gln glu ile tyr gln glu leu thr arg leu lys ala ala val gly glu leu pro glu lys  
421/141 451/151  
TCT AAG CTG CAG GAG ATC TAC CAG GAG CTG ACC TGG CTG AAG GCT GCA GTG GGT GAG CTT  
ser lys leu gln glu ile tyr gln glu leu thr trp leu lys ala ala val gly glu leu  
481/161 511/171  
CCA GAG AAA TCT AAG ATG CAG GAG ATC TAC CAG GAG CTG ACT CGG CTG AAG GCT GCA GTG  
pro glu lys ser lys met gln glu ile tyr gln glu leu thr arg leu lys ala ala val  
541/181 571/191  
GGT GAG CTT CCA GAG AAA TCT AAG CAG CAG GAG ATC TAC CAG GAG CTG ACC CGG CTG AAG  
gly glu leu pro glu lys ser lys gln gln glu ile tyr gln glu leu thr arg leu lys  
601/201 631/211  
GCT GCA GTG GGT GAG CTT CCA GAG AAA TCT AAG CAG CAG GAG ATC TAC CAG GAG CTG ACC  
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661/221 691/231  
CGG CTG AAG GCT GCA GTG GGT GAG CTT CCA GAG AAA TCT AAG CAG CAG GAG ATC TAC CAG  
arg leu lys ala ala val gly glu leu pro glu lys ser lys gln gln glu ile tyr gln  
721/241 751/251  
GAG CTG ACC CAG CTG AAG GCT GCA GTG GAA CGC CTG TGC CAC CCC TGT CCC TGG GAA TGG  
glu leu thr gln leu lys ala ala val glu arg leu cys his pro cys pro trp glu trp  
781/261 811/271  
ACA TTC TTC CAA GGA AAC TGT TAC TTC ATG TCT AAC TCC CAG CGG AAC TGG CAC GAC TCC  
thr phe phe gln gly asn cys tyr phe met ser asn ser gln arg asn trp his asp ser  
841/281 871/291  
ATC ACC GCC TGC AAA GAA GTG GGG GCC CAG CTC GTC GTA ATC AAA AGT GCT GAG GAG CAG  
ile thr ala cys lys glu val gly ala gln leu val val ile lys ser ala glu gln  
901/301 931/311  
AAC TTC CTA CAG CTG CAG TCT TCC AGA AGT AAC CGC TTC ACC TGG ATG GGA CTT TCA GAT  
asn phe leu gln leu gln ser ser arg ser asn arg phe thr trp met gly leu ser asp  
961/321 991/331  
CTA AAT CAG GAA GGC ACG TGG CAA TGG GTG GAC GGC TCA CCT CTG TTG CCC AGC TTC AAG  
leu asn gln glu gly thr trp gln trp val asp gly ser pro leu leu pro ser phe lys  
1021/341 1051/351  
CAG TAT TGG AAC AGA GGA GAG CCC AAC AAC GTT GGG GAG GAA GAC TGC GCG GAA TTT AGT  
gln tyr trp asn arg gly glu pro asn asn val gly glu glu asp cys ala glu phe ser  
1081/361 1111/371  
GGC AAT GGC TGG AAC GAC GAC AAA TGT AAT CTT GCC AAA TTC TGG ATC TGC AAA AAG TCC  
gly asn gly trp asn asp asp lys cys asn leu ala lys phe trp ile cys lys ser  
1141/381 1171/391  
GCA GCC TCC TGC TCC AGG GAT GAA GAA CAG TTT CTT TCT CCA GCC CCT GCC ACC CCA AAC  
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1201/401  
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< 110> KONINKLIJKE UNIVERSITEIT NIJMEGEN

< 120> COMPOSITION AND METHOD FOR MODULATING DENDRITIC CELL - T CELL INTERACTION

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ggtcccctgg	tgctgcaact	cctctccTTc	acgctttgg	ctgggctcct	tgtccaagtG	180
tccaaggTCC	ccagctccat	aagttagggaa	caatccaggC	aagacgcata	ctaccagaac	240
ctgacccAGC	ttaaagctgc	agtgggtgag	ctctcagaga	aatccaagct	gcaggagatC	300
taccaggAGC	tgaccCAGCT	gaaggctgca	gtgggtgagC	ttccagagaa	atctaagCTG	360
caggagatCT	accaggagCT	gacCCGGCTG	aaggctgcag	tgggtgagCT	tccagagaaa	420
tctaagCTGC	aggagatcta	ccaggagCTG	acctggCTGA	aggctgcagt	gggtgagCTT	480
ccagagaaAT	ctaagatgca	ggagatctac	caggagCTGA	ctcggctgaa	ggctgcagtG	540
ggtgagCTTC	cagagaaATC	taagcagcag	gagatctacc	aggagctgac	ccggctgaag	600
gctgcagtgg	gtgagCTTCC	agagaaatct	aagcagcagg	agatctacca	ggagctgacc	660
cggctgaagg	ctgagCTGGG	tgagCTTCCA	gagaaatcta	agcagcaga	gatctaccag	720
gagctgacCC	agctgaaggc	tgcagtgaa	cgcctgtGCC	accCCTGTCC	ctgggaatgg	780
acattctcc	aaggAAACTG	ttacttcATG	tctaactccc	agcggAACTG	gcacgactCC	840
atcaccgcct	gcaaAGAAGT	ggggggccCAG	ctcgTCGtaa	tcaaaAGTGC	tgaggagcag	900
aacttcctac	agctgcagTC	ttccagaagt	aaccgttca	cctggatggg	actttcagat	960
ctaaatcagg	aaggcacgtg	gcaatgggtg	gacggctcac	ctctgtGCC	cagcttcaag	1020
cagtattgga	acagaggaga	gccccacaac	gttggggagg	aagactgcgc	ggaatttagt	1080
ggcaatggct	ggaacgacga	caaAtgtaat	cttgccaaat	tctggatCTG	caaaaAGTCC	1140
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