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**THE BETA1 CYTOPLASMIC DOMAIN REGULATES THE  
LAMININ-BINDING SPECIFICITY OF THE ALPHA7X1 INTEGRIN**

by

Ming-Guang Yeh

DISSERTATION

Submitted in partial satisfaction of the requirements for the degree of

DOCTOR OF PHILOSOPHY

in

Oral Biology

in the

GRADUATE DIVISION

of the

UNIVERSITY OF CALIFORNIA, SAN FRANCISCO



Date

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**ABSTRACT**

THE  $\beta 1$  CYTOPLASMIC DOMAIN REGULATES THE  
LAMININ-BINDING SPECIFICITY OF THE  $\alpha 7X1$  INTEGRIN

Ming-Guang Yeh

Chairperson of the Supervisory Committee: Professor Randall H. Kramer

Department of Stomatology, Oral Biology Division.

The laminin-binding integrin  $\alpha 7$  is alternatively spliced in the putative ligand-binding region to yield either the  $\alpha 7X1$  or the  $\alpha 7X2$  form of the receptor. Similarly, the partner  $\beta 1$  integrin cytoplasmic domain is converted from  $\beta 1A$  to  $\beta 1D$ . To determine whether  $\beta 1D$  modulates the activity of the  $\alpha 7$  receptor, we transfected MCF-7 cells expressing mouse  $\alpha 7X1$  with  $\beta 1D$  cDNA. Cells expressing mouse  $\alpha 7X1$  coupled with  $\beta 1A$  and failed to adhere to laminin-1, whereas transfectants expressing  $\alpha 7X1$  and  $\beta 1D$  showed strong adhesion. Contrarily,  $\alpha 7X2$  adhered to laminin-1 readily when coupled with  $\beta 1A$ . We also generated human  $\alpha 7X1$  transfectants in which the expression of  $\beta 1A$  or  $\beta 1D$  was regulated by the tetracycline promoter system.  $\beta 1D$  overexpression induced adhesion of  $\alpha 7X1$  cells to laminin-1, whereas induction of  $\beta 1A$  expression did not. Attachment to other ligands (e.g., collagen type I) was unaltered. Interestingly,  $\alpha 7X1$  whether complexed with  $\beta 1A$  or  $\beta 1D$  displayed the same level of poor adhesion to laminin-2/4 or

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strong adhesion to laminin-10/11. These findings indicate that  $\alpha 7$  function is regulated not only by X1/X2 in its extra cellular domain but also by  $\beta 1$  cytoplasmic splice variants. Expression of  $\beta 1D$  appears to alter the integrin ligand specificity of  $\alpha 7X1$  by enhancing its binding to laminin-1 but not to laminin-2/4 or laminin-10/11. Functional regulation of  $\alpha 7\beta 1$  by developmentally regulated splicing events may be important during myogenic differentiation and repair, because the integrin participates in multiple activities related to cell proliferation and survival, motility, and formation of stable myotendinous and neuromuscular junctions.



Randall H. Kramer, Ph.D.

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## **LIST OF ABBREVIATIONS**

**BSA, bovine serum albumin**

**cDNA, complementary DNA;**

**Col, collagen**

**DMEM, Dulbecco's modified Eagle's medium**

**DMSO, dimethyl sulfoxide**

**Dox, doxycycline**

**dpc, days post conception**

**GUS, beta-glucuronidase**

**LN, laminin**

**mAb, monoclonal antibody**

**MTJ, myotendinous junction**

**NMJ, neuromuscular junction**

**pAb, polyclonal antibody**

**PAGE, polyacrylamide gel electrophoresis**

**PCR, polymerase chain reaction**

**SDS, sodium dodecyl sulfate**

**UDG, uracil DNA glycosylase**

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## Chapter 1

### **Introduction and the Goals of Study**

#### *The importance of integrins*

The dynamic interaction between myoblasts and laminins is important for skeletal muscle development, regeneration, and myotube stability and survival (Vachon *et al.*, 1997; Gullberg *et al.*, 1999; Colognato and Yurchenco, 2000). The muscle-specific laminin-binding  $\alpha7\beta1$  integrin mediates myoblast motility on laminin substrates (Kaufman *et al.*, 1991; Yao *et al.*, 1996a; Yao *et al.*, 1996b; Crawley *et al.*, 1997; von der Mark *et al.*, 2002). In mature skeletal muscle, the  $\alpha7$  receptor is associated with costameres and the myotendinous and neuromuscular junctions (Bao *et al.*, 1993; Belkin *et al.*, 1996; Martin *et al.*, 1996; van der Flier *et al.*, 1997). The physiological importance of  $\alpha7$  is demonstrated in  $\alpha7$ -null mutant mice, which develop a form of muscular dystrophy with myotendinous junction (MTJ) defects (Mayer *et al.*, 1997), and in humans with mutations in  $\alpha7$  gene, which develop congenital myopathies (Hayashi *et al.*, 1998; Pegoraro *et al.*, 2002).

#### *Splice variants of integrin $\alpha7\beta1$*

The  $\alpha7$  integrin subunit is alternatively spliced in both the extracellular (X1 and X2) (Ziobar *et al.*, 1993) and the cytoplasmic (A, B and C) domains (Song *et al.*, 1993). One report showed that there was also a D splice variant of  $\alpha7$  in adult human striated muscle

(Leung *et al.*, 1998). However, the function of the D isoform has not been widely studied (Vignier *et al.*, 1999). NMJs have all A, B, and C isoforms of  $\alpha 7$ ; the MTJs have both A and B isoforms of  $\alpha 7$ ; and the non-junction sites have only C isoform of  $\alpha 7$  (Hodges *et al.*, 1997). However,  $\alpha 7C$  splice variant was shown to be non-existent in vivo in one report (Vignier *et al.*, 1999). The purpose of this study is to know the functionality of the  $\alpha 7X1$  and  $\alpha 7X2$  extracellular splice isoforms. In terms of adhesion to laminin-1, both cytoplasmic A and B isoforms of  $\alpha 7$  are equally competent (Yao *et al.*, 1996b). This thesis addresses the difference between  $\alpha 7X1$  and  $\alpha 7X2$  in the context of the B cytoplasmic tail.

The  $\beta 1$  subunit also has several cytoplasmic splice variances-A, B, C, and D isoforms. They are all derived by alternative splicing from exactly the same position in cytoplasmic tail.  $\beta 1A$  is ubiquitously expressed.  $\beta 1B$  mRNA is coexpressed at a low level together with  $\beta 1A$  mRNA in many tissues, but the protein has been detected only in skin and liver (Balzac *et al.*, 1993).  $\beta 1B$  is found only in humans and does not localize to focal contacts and exhibits a dominant negative effect on adhesive events specific to only  $\beta 1$  integrins (Cali *et al.*, 1998; Armulik *et al.*, 2000).  $\beta 1C$  has two subtypes:  $\beta 1C-1$  (Languino and Ruoslahti, 1992) and  $\beta 1C-2$  (Svineng *et al.*, 1998).  $\beta 1C-1$  was shown to be expressed in placenta, normal platelets and several blood cells lines (Languino and Ruoslahti, 1992). Both  $\beta 1C-1$  and C-2 were shown to be growth inhibitory (Fornaro *et al.*, 1998; Fornaro *et al.*, 1999; Meredith *et al.*, 1999).  $\beta 1D$  is present only in cardiac and skeletal muscles in vertebrates. In early embryogenesis, the only  $\beta 1$  isoform present in

striated muscle is  $\beta 1A$ . Late in development,  $\beta 1D$  is expressed and replaces  $\beta 1A$  completely in adult striated muscle (van der Flier *et al.*, 1995; Zhidkova *et al.*, 1995; Belkin *et al.*, 1996). The detailed temporal expression pattern is discussed elsewhere under the section of “integrin profile during myogenesis”. The switch from  $\beta 1A$  to  $\beta 1D$  subunit in cardiac muscle cells coincides with the expression of  $\alpha 7$  in mouse (van der Flier *et al.*, 1997).  $\beta 1D$  is present in myotendinous junctions and costameres in skeletal muscles, and in intercalated discs of cardiac muscle. The amount of  $\beta 1D$  in different types of skeletal muscle (fast, slow, and mixed type) is similar, but cardiac muscle expresses about five times as much as skeletal muscle (van der Flier *et al.*, 1997). This thesis addresses the different effects of  $\beta 1A$  and  $\beta 1D$ .

This complex differentiation-dependent pattern of  $\alpha 7$  and  $\beta 1$  splicing allows the generation of a unique set of variant  $\alpha 7\beta 1$  complexes that are structurally, and presumably functionally, distinct.

### *Structure of $\alpha 7$ integrin*

The  $\alpha$  subunits of integrins can be classified into two major groups. The first one contains a 180 amino acid insert between the second and the third homology repeat (see below). This insert is called I domain and is homologous to the collagen binding site found in von Willebrand factor. The crystal structure of this domain has been solved (Lee *et al.*, 1995). This group includes  $\alpha 1$ ,  $\alpha 2$ ,  $\alpha M$ ,  $\alpha L$ , and  $\alpha X$ . The second group does not have the I-domain;  $\alpha 3$ ,  $\alpha 4$ ,  $\alpha 5$ ,  $\alpha 6$ ,  $\alpha 7$ ,  $\alpha IIb$ , and  $\alpha V$  belong to the non-I-domain group.

Besides the presence of I-domain, integrin  $\alpha$  subunits are highly conserved. The N-terminus of extracellular domain of all  $\alpha$  subunits has 7 homologous repeats (see Figure 1).

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Figure 1.

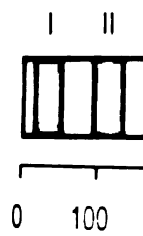


Figure 1.

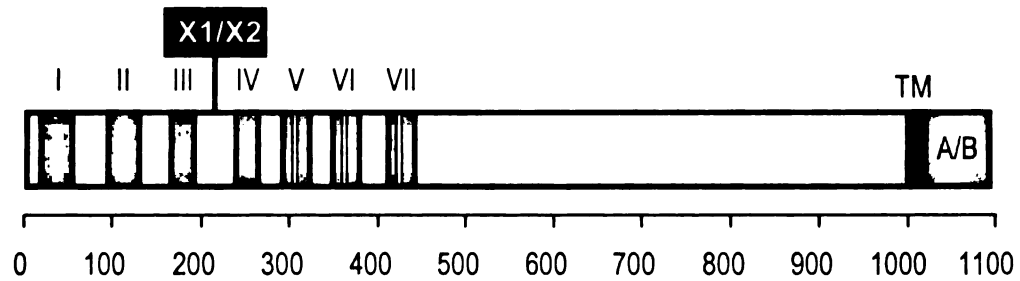
identified al

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**Figure 1.**



**Figure 1. The  $\alpha 7$  integrin subunit.** The relative positions of the identified alternatively spliced sites in the extracellular domain (X1 and X2) and the cytoplasmic domains (A and B) are indicated. Also shown are the seven homology repeat domains (I-VII, gray boxes) and divalent-cation-binding sites (white slots), and the transmembrane domain (TM).

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transmembrane region.

According to one model, the 7 homology repeats region forms a 7-bladed  $\beta$ -propeller (Springer, 1997). According to this model, each blade is composed of 4 invariable regions and 4 alternate variable regions. The nomenclature for these regions has also been coined according to this model. Basically, only the invariable regions are designated alphabetically from N to C terminus. There are 4 invariable regions in each blade, so it starts from A, then followed by variable region, then invariable region B, then variable region, then invariable region C and so on. To distinguish the invariable regions from each blade, a number is put before the alphabet to denote its blade position. For example, 3C is the third invariable region in the homologous repeat III (see Figure 2). As shown in figure 2, regions 3B, 3C, and 3D are the 3 invariable regions of blade III, and 4A is the first invariable region of blade IV. In the transition from 3D to 4A, a highly conserved motif (CUP) of this blade constitute one seventh of the cup region which forms the top of propeller and holding an arginine residue from  $\beta_1$ . Recently, this model has been confirmed by X-ray crystallography (Xiong *et al.*, 2001). Crystal structure of the extracellular segment of integrin  $\alpha V\beta_3$  in complex with an Arg-Gly-Asp ligand has also been reported (Xiong *et al.*, 2002). The crystal structure reveals that the ligand peptide inserts into a crevice formed between the propeller of  $\alpha_7$  and the  $\beta_A$  domain of  $\beta_1$ . More precisely, the arginine side chain inserts into a narrow groove at the top of the propeller domain, formed primarily by the homology repeat III and IV. The asparagine of the ligand peptide contains a carboxylate group, which protrudes into a cleft between the  $\beta_A$  loops (A'- $\alpha_1$  and C'- $\alpha_3$ ) of  $\beta_1$ . The rest of the  $\alpha$  subunits consist of  $\beta$ -sandwich domain, transmembrane region, and cytoplasmic domain.

Figure 2.

msAlpha7 X2B  
msAlphaV  
Consensus

msAlpha7 X2B  
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msAlpha7 X2B  
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msAlpha7 X2E  
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**Figure 2.**

N-terminus protein sequence alignment of  $\alpha$ V and X2B

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msAlpha7 X2B	(1)	MARI	PRCD	FLPP	PGIYYL	ITTS	LLAGLFI	PPAIA	AFNLDVM	G																												
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		Section 2																																				
	(41)	41	50	60	70	80																																
msAlpha7 X2B	(41)	AIRKE	GEPG	SLSFG	SV	ALHRQ	LQPRP	QSW	LLVGAP	QALAL																												
msAlphaV	(38)	PAEY	AGPE	GSYFG	AV	DFEFP	STSS-	RM	ELLVGAP	KANTT																												
Consensus	(41)		G	GS	FGFAV				FLLVGAP	A																												
		Section 3																																				
	(81)	81	90	100	110	120																																
msAlpha7 X2B	(81)	PGQQAN	RTGGL	FACPL	LS	LEETDC	YRVDI	DRGANV	QKES	--																												
msAlphaV	(77)	QPGIVE	GGQVL	KCECS	SS	RRCQPI	EFD	STGNR	RDYAK	DPL																												
Consensus	(81)			L	S		D			KD																												
		Section 4																																				
	(121)	121	130	140	150	160																																
msAlpha7 X2B	(119)	--KEN	QWLG	SVRS	QGP	GKIV	TC	AHRYE	SRQR	VDQALET																												
msAlphaV	(117)	EFKS	HQWFG	ASVRS	KQ--	DKI	AC	APLYH	WRTE	IKQERE	P																											
Consensus	(121)	K	QW	G	SVRS		KIL	CA	Y	R	M	Q	E																									
		Section 5																																				
	(161)	161	170	180	190	200																																
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msAlphaV	(155)	-----	VG	TCF	LD	Q	D	G	T	K	T	V	E	A	P	C	R	S	K	N	I	D	A	D	G	Q	G	F										
Consensus	(161)					I	D			EW	C	K										D	G	F														
		Section 6																																				
	(201)	201	210	220	230	240																																
msAlpha7 X2B	(197)	CQQGT	AATF	SPDS	HYIV	FG	AP	CTYN	NW	KGLL	FVTN	ID	SSDP																									
msAlphaV	(185)	CQGGF	SIDE	T	KADR	VLL	G-	G	PC	SE	YW	Q	Q	L	I	S	D	Q	V	A	E	I	I	S														
Consensus	(201)	CQ	G	A	FS		LL	AP	G	S	F	W	G	L		NI																						
		Section 7																																				
	(241)	241	250	260	270	280																																
msAlpha7 X2B	(237)	--DQL	VYK	T	LDP	ADR	LT	G	P	A	G	D	E	T	L	N	S	Y	L	G	F	S	I	D	S	G	K	G	L	M								
msAlphaV	(224)	KYD	P	N	V	Y	S	I	K	Y	N	N	Q	L	A	T	R	T	A	Q	A	I	F	D	D	S	Y	L	G	S	V	A	V	G	-	D	F	N
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X1/X2 splice region

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The X1/X2 alternative splicing in the  $\alpha 7$  mRNA occurs in the variable region between the III and IV homology repeat domains near the putative ligand-binding site, where it may define ligand specificity, affinity, or ligand-binding competence (Irie *et al.*, 1995; Kamata *et al.*, 1995; Tozer *et al.*, 1999). Aligning the  $\alpha 7$  sequence to the  $\alpha V$  sequence published in the crystal structure paper with sequence alignment tool (Vector NTI, version 6 from InforMax), the X1/X2 alternative splice region spans the invariable region 3C to the 4A (see Figure 2 ). Amino acid residues in the  $\alpha$ IIb subunit those are critical for ligand binding to integrin  $\alpha$ IIb $\beta$ 3 are identified in the  $\beta$ -propeller model (Kamata *et al.*, 2001). Using alanine scanning strategy, they found several hot spots for ligand binding. Most of them are within the 3B to 4A regions. According to alignment by Vector NTI program, this region includes the X1/X2 splice region.

#### *Developmentally regulated expression of $\alpha 7 \beta 1$ variants*

The formation of a multinucleated muscle fiber is an extraordinary dynamic process involving multiple cell adhesive interactions consisting of homotypic and heterotypic cell-cell interactions as well as interactions with extracellular matrix (ECM) molecules. Early studies of integrins on myogenic cells revealed that  $\beta 1$  integrins localized to focal contacts, costameres, neuromuscular junctions, and myotendinous junctions. One major ECM component in the basement membrane (BM) of striated muscle tissue is laminin. Integrin  $\alpha 7 \beta 1$  has been shown to be the receptor for this ligand in skeletal muscle (Vachon *et al.*, 1997). As been mentioned earlier,  $\alpha 7 \beta 1$  integrin has splice isoforms whose expression is developmentally regulated. Data gathered from

studies on mouse myogenesis is summarized by Gullberg *et al.* (Gullberg *et al.*, 1998) and by Brancaccio *et al.* (Brancaccio *et al.*, 1998). The major focus of this study is the  $\alpha 7$  X1/X2 and  $\beta 1$  A/D splice variants and the developmental expression patterns of these isoforms are schematized in the following figure (Figure 3).

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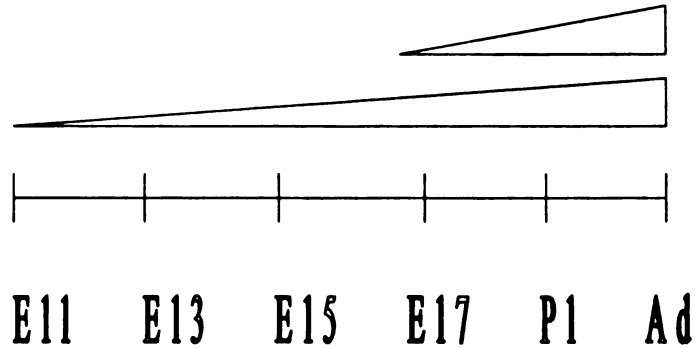
**Figure 3.**

Timing of expression of integrin in developing mouse tissue, based on western and immunochemistry, adopted from Brancaccio *et al.* (1998).

**beta1D expression**

**skeletal muscle**

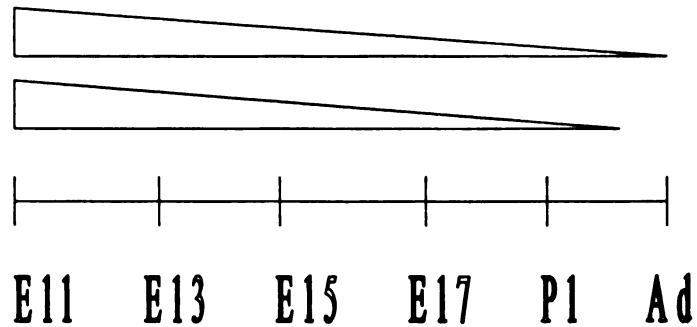
**cardiac muscle**



**beta1A expression**

**skeletal muscle**

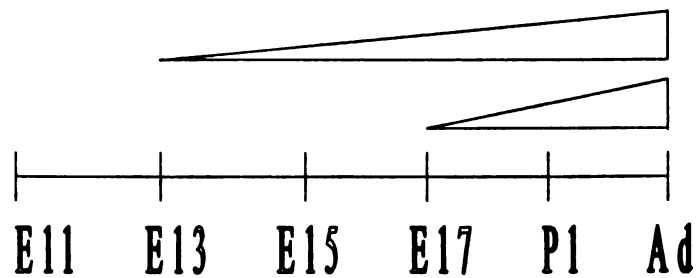
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**alpha 7 expression**

**skeletal muscle**

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### *Spatial and temporal distribution of laminin isoforms*

The basement membrane (BM) surrounding muscle cells can be classified into three domains. First, the domain where muscle end-points attach to tendons is called the myotendinous junction (MTJ). During development and growth, the MTJ is subject to dynamic changes. According to electron microscopic studies on developing MTJs in avians, basement membrane deposition is one of the first steps in MTJ formation (Tidball and Lin, 1989). Later, junctional folds and myofibrillar insertion occur. In the adult, the MTJ is the major site for the transmission of force from the muscle cell interior to the surrounding ECM. Second, the specialized structures where motor neurons attach are called neuromuscular junctions (NMJ). Third, the basement membrane surrounding the majority of the sarcolemma, excluding the MTJs and the NMJs, is referred to as the sarcolemma BM.

Laminin is a heterotrimer of  $\alpha$ ,  $\beta$  and  $\gamma$  chains. The laminin family currently consists of at least 12 heterotrimers that are formed from five  $\alpha$  chains, three  $\beta$  chains and three  $\gamma$  chains (Timpl and Brown, 1994; Timpl, 1996; Colognato and Yurchenco, 2000). The detection of temporal and spatial distribution of laminin heterotrimers is complicated by the presence of multiple chains and not much information has been reported. In developing and mature mouse muscle, the expression of laminins is complex, and is summarized in Table I (Gullberg *et al.*, 1999). Due to the lack of a full battery of antibodies to detect human laminin  $\alpha$  chains (even the specificity of detecting mAb 4C7 has previously been erroneously attributed to  $\alpha 1$ ) and to the difficulty of sampling, knowledge about laminin isoforms in human muscle has been only fragmentary (Pedrosa-

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Domellof *et al.*, 2000). In human muscle, laminin  $\alpha 1$  chain is expressed at myotendinous junctions as early as 8 weeks of gestation and is evident at 22 weeks (Pedrosa-Domellof *et al.*, 2000). In the MTJ of adult human, only a rudimentary amount of the laminin  $\alpha 1$  chain is detected. Similarly, in mouse muscle formation, the  $\alpha 1$  chain is present at the MTJ from E11 through the late fetal stage, but not at the sarcolemma or neuromuscular junction basement membrane (Patton *et al.*, 1997). The laminin  $\alpha 5$  chain has a similar expression pattern as that of the  $\alpha 1$  chain. It was also detected at 8-10 weeks in the MTJ, but it is also expressed at other two BM compartments. Laminin  $\alpha 2$  chain has a totally different expression than  $\alpha 1$  or  $\alpha 5$ . It was detected at all the BM compartments from 12 weeks onward.

In summary, the early accumulation of the laminin  $\alpha 2$  and  $\alpha 5$  chains at the MTJ is followed by expression of these isoforms along the entire length of the developing myotube. On the contrary, the  $\alpha 1$  chain was restricted only to MTJs. In the adult,  $\alpha 1$  and  $\alpha 5$  staining was diminished at the MTJ but is still weakly detectable whereas  $\alpha 2$  chain was strongly expressed.

Table I. Temp

$\alpha 7$

$\alpha 7 X1^S$

$\alpha 7 X2^S$

$\beta 1D$

Laminin-1

Laminin-2 4

Laminin-10 11

\*Not detected

\*\*Weak expression

\*\*\*The more + si

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**Table I.** Temporal distribution of  $\alpha 7$ ,  $\beta 1D$ , and laminin isoforms in mice

	Early embryo	Fetus	Late Fetus	New Born	Adult
$\alpha 7$	-*	**	+++	+++++	+++++++
$\alpha 7X1^{\S}$	-	+/-***	++	+	+/-
$\alpha 7X2^{\S}$	+/-	+	+++	+++++	+++++++
$\beta 1D$	+/-	+	+++	++++	+++++++
Laminin-1	+++++	+++	++	+	+/-
Laminin-2/4	+	++	+++	++++	++++
Laminin-10/11	+++	+++	+	+	+/-

\*Not detected

\*\*Weak expression.

\*\*\*The more + sign, the stronger the expression.

$\S$ Detect mRNA by PCR

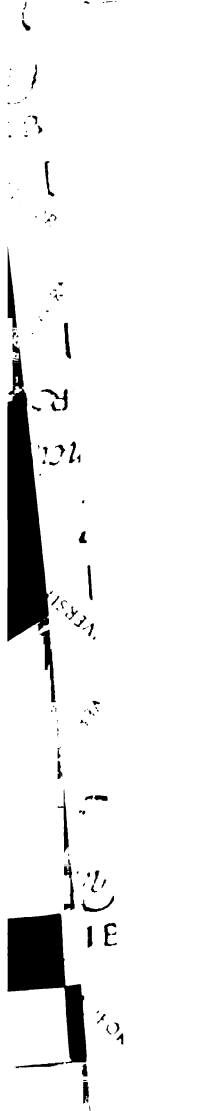
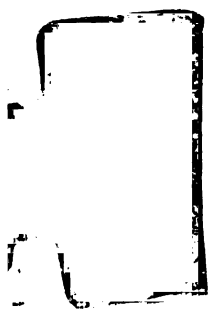
### *Goals of the study*

Integrins are capable of changing their function dynamically (Diamond and Springer, 1994; Humphries, 1996; Mould, 1996; Hughes and Pfaff, 1998). It has been proposed (reviewed in (Liddington and Ginsberg, 2002)) that certain intracellular events lead to alterations in integrin conformation, resulting in enhanced adhesion to ligand (inside-out signaling). In addition, the process of ligand binding may regulate the activity of the integrin receptor (outside-in signaling). In the case of inside-out signaling, there is strong evidence that the actin cytoskeleton and its linker proteins (e.g., talin) provide one mechanism that can regulate integrin activation. This process may result in changes in

integrin affinity or integrin-clustering events leading to increased avidity (Calderwood *et al.*, 2000). However, it remains controversial whether events such as lateral rearrangements that may modulate receptor avidity can play a role in integrin function (reviewed in Shimaoka *et al.*, 2002). As stated above, following myoblast differentiation into myotubes and mature muscle, the  $\beta 1$  integrin cytoplasmic alternative splice shifts from A to D. Given the consensus that the cytoplasmic domains of  $\beta$  integrins can regulate the conformation and activity of integrin subunits (Schwartz *et al.*, 1995; Hughes *et al.*, 1996; Crommie and Hemler, 1998; Calderwood *et al.*, 1999; Woodside *et al.*, 2001; Liddington and Ginsberg, 2002), it is possible that the  $\beta 1D$  isoform controls  $\alpha 7X1/X2$  function. To investigate this possibility, we examined the functionality of  $\alpha 7$  integrin in the presence of the  $\beta 1A$  or  $\beta 1D$  cytoplasmic domain. The ligands studied were laminin-1, laminin-2/4, and laminin-10/11 because expression of these laminin isoforms switches during the time period when  $\beta 1A$  switches to  $\beta 1D$  and because  $\alpha 7\beta 1$  is the major receptor for laminin. The cell line used for this study was the breast carcinoma cell line MCF-7, because this cell line neither express  $\alpha 7$  nor bind to laminin-1 and because  $\alpha 7X1$  and  $\alpha 7X2$  binds differently to laminin-1 when expressed in MCF-7. There are several non-integrin receptors also known to bind laminin. Dystroglycan is the most important one in this category. It is now apparent that dystroglycan plays a critical role in the pathogenesis of several muscular dystrophies and is involved in early development, organ morphogenesis, and synaptogenesis. Dystroglycan interacts with a wider repertoire of extracellular ligands other than laminin. The interaction of the cytoskeleton with dystroglycan is not as efficient as that with integrin. In the MCF-7 tissue culture system, the adhesion to ECM mainly is attributed to integrin but not dystroglycan or other

receptors in sho

laminin-binding





## Chapter 2

### **Materials and Methods**

#### *ECM proteins*

For cell adhesion studies, several different ligands were used as described previously. Laminin-1 isolated from the mouse EHS tumor was obtained from Gibco-BRL (Life Technologies, Invitrogen, Carlsbad, CA). Human placental merosin, consisting of laminin-2/4 isoforms purified by EDTA extraction followed by ion-exchange chromatography was purchased from Gibco-BRL. Human laminin-10/11, isolated from placenta by mild pepsin digestion and by affinity chromatography on mAb 4C7-coupled Sepharose, was also from Gibco-BRL. Collagen was obtained from Cohesion (Palo Alto, CA). Human vitronectin and fibronectin were gifts from Caroline Damsky (University of California, San Francisco; UCSF).

#### *Antibodies*

Antibodies against integrin subunits—rat anti-human  $\beta 1$  (mAb AIIB2), mouse anti-human  $\alpha 2$  (mAb VM1), rat anti-human  $\alpha 6$  (mAb GoH3), and rabbit anti-mouse  $\beta 1 D$  cytodomain (polyclonal antibody anti- $\beta 1 D$ )—were kindly provided by Caroline Damsky, Vera Morhenn (SRI International, Menlo Park, CA), Arnond Sonnenberg (Netherlands Cancer Center, Amsterdam), and Eva Engvall (Burnham Institute, La Jolla Cancer Research Center, CA), respectively. Mouse anti-human  $\alpha 3$  (mAb J143) was from the

American Type Culture Collection (ATCC) (Manassas, VA); rat anti-mouse  $\beta 1$  (mAb MB1.2) and mouse anti-human  $\beta 1$  (mAb 2000) were from Chemicon (Temecula, CA); hamster anti-mouse/human  $\beta 1$  (mAb Ha2/5) was from Pharmingen (BD Bioscience, San Diego, CA); rat anti-mouse  $\alpha 7$  (mAb CY8), mouse anti-human  $\alpha 7$  (mAb 9.1), rabbit anti-human  $\beta 1$  cytodomain (pAb 22778), and rabbit anti-mouse  $\alpha 7$  light chain (pAb 1211) were from our laboratory, as previously described (Yao *et al.*, 1996b; Yao *et al.*, 1997; Vizirianakis *et al.*, 2001). Fluorescein-conjugated secondary antibodies were obtained from Jackson ImmunoResearch Laboratories (West Grove, PA).

#### *Construction of Mouse $\beta 1D$ cDNA*

The full-length mouse  $\beta 1D$  cDNA was prepared by reverse transcription followed by polymerase chain reaction (PCR) from a mouse skeletal muscle cDNA library (Clontech, Palo Alto, CA). The two primers for PCR were: reverse primer, 5'-CTAGTCTAGAATTCAGAGACCAGCTTTACGTCCATAG and forward primer, 5'-CGCGGATCCGAATTCAAGATGAATTTGCAACTGGTTTCCTG. The sequence of full-length mouse  $\beta 1D$  was verified at the Biomolecular Resource Center (UCSF). The mouse  $\beta 1D$  cDNA was then ligated into *Bam*HI and *Xba*I sites of pcDNA3.1/Hygro (Invitrogen, Carlsbad, CA). Full-length mouse  $\beta 1A$  cDNA was obtained from Louis Reichardt (UCSF) and was cloned into the regulatable expression vector in the same manner as  $\beta 1D$  (see below).

## Cloning of human $\alpha 7X1$

Full-length human  $\alpha 7X2$  cDNA (Vizirianakis *et al.*, 2001) was used as a template for the generation of full-length human  $\alpha 7X1$ . The X2 exon was replaced with the X1 exon by uracil DNA glycosylase (UDG) excision. The generation of cohesive ends on the PCR product of the X1 exon was performed by UDG-mediated excision of uracil.

Specifically, we designed primers containing uracil that matched the corresponding T in the splicing regions. We then used these primers to generate the X1 exon by PCR from human  $\alpha 7$  cDNA using human fetus limb muscle RNA as template (forward primer: 5'

-AAGGGUACUGCCAGGGTGGAGCTCTGTG; reverse primer: 5'

-AGAAGCCAAAGUAGCUGTTGGCAGGGA). The full-length cDNA proximal to the splicing site was also generated by PCR (forward primer: 5'

-CCGGGATCCATGGCCGGGGCTCGGAGCCG; reverse primer: 5'

-AGTACCCUCCAATTATAGGTTCTGGGG). The primers for full-length cDNA distal to the splicing site were as follows: forward primer, 5'

-AGCTACTTTGGCUUCUCTATTGACTCGGGGAAAGGTCTG; reverse primer, 5'

-CCGCTCGAGCTAGGCGGTGCCTGGCCCT. The three PCR products were first

treated with UDG according to the manufacturer's protocol (Gibco-BRL) and were then

ligated. The ligation product was purified by gel electrophoresis followed by cloning into

pCDNA3.1/Hygro and verified by sequencing.

## *Generation of Cell Lines*

MCF-7 human carcinoma cells and the  $\alpha 7$ -transfected cells were cultured as described previously (Yao *et al.*, 1996a; Ziober *et al.*, 1997). MCF-7 cells do not adhere well to laminin-1 even though they express moderate levels of integrins  $\alpha 2\beta 1$ ,  $\alpha 3\beta 1$ , and  $\alpha 6\beta 1$ , which are potential receptors for this ligand. Previously we generated MCF-7 cells expressing full-length mouse  $\alpha 7$  cDNA containing either the X1 or X2 splice form, which formed heterodimers with the partner  $\beta 1A$  subunit. MCF-7 cells expressing the splicing isoform  $\alpha 7X2$  were active in binding laminin-1, whereas cells expressing  $\alpha 7X1$  lacked ligand-binding activity (Ziober *et al.*, 1997). The original  $\alpha 7X1$ -expressing population was further enriched by fluorescence-activated cell sorting (FACS) using mAb CY8 against mouse  $\alpha 7$  to yield the X1/A cell line used in the current experiments. The X2/A cell line was derived from MCF-7 cells transfected with mouse  $\alpha 7X2$  which heterodimerizes with the endogenous human  $\beta 1A$  (Yao *et al.*, 1996a).

### *Experimental approaches*

Expression of mouse  $\beta 1D$  in MCF-7 cells was generated in two different systems (see Flow-chart I). In the first, the X1/A-p cells were stably transfected with mouse  $\beta 1D$  cDNA by the calcium phosphate procedure (Mammalian Transfection Kit; Stratagene, La Jolla, CA). Following selection with hygromycin, the double transfectants ( $\alpha 7X1/\beta 1D$ ) were enriched by FACS after labeling with anti-mouse  $\beta 1$  antibody (mAb MB1.2; Chemicon). The enriched population was thereafter called X1/D. In addition, a total of 20 single cell clones were isolated either by single-cell sorting by FACS or by limiting dilution into individual wells of 96-well plates. The expression of mouse  $\beta 1D$  was

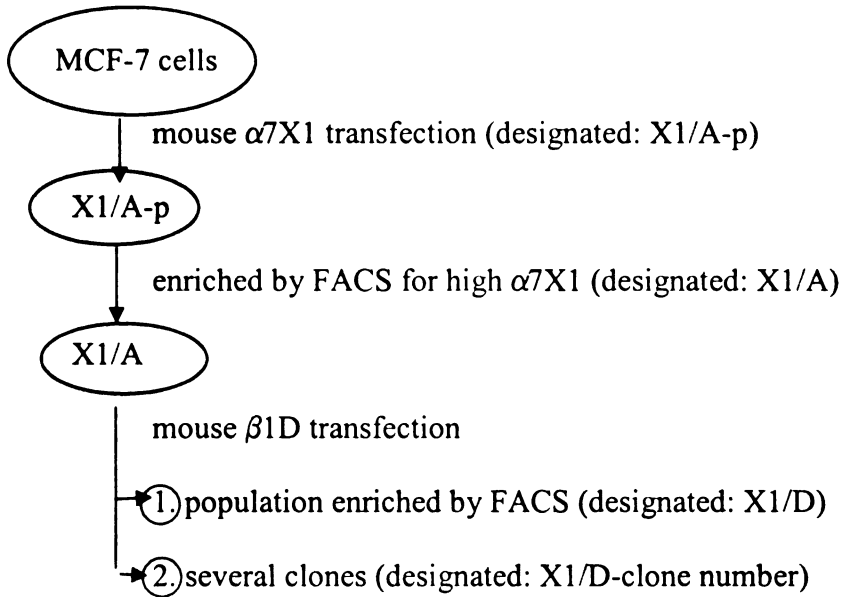
verified by immunoblot analysis using a  $\beta$ 1D-specific pAb from Eva Engvall and by flow cytometry using mAb MB1.2. A number of clones with a range of  $\alpha$ 7 and  $\beta$ 1D levels, as determined by FACS, were chosen for further study.

In the second approach, MCF-7 cells were transfected with pTet-Off (pUHD15-1 neo; Clontech), and the expression of tTA-vp16 was verified. The MCF-7 Tet-Off cells were then transfected with the full-length cDNA for either human  $\alpha$ 7X1 integrin (as described above) or human  $\alpha$ 7X2 integrin (Vizirianakis *et al.*, 2001). We used anti-human  $\alpha$ 7 mAb 9.1 to enrich the  $\alpha$ 7-expressing population by FACS. Retroviral transduction was then used to introduce the pRevTRE plasmids (Clontech) with mouse  $\beta$ 1A or mouse  $\beta$ 1D into the human  $\alpha$ 7 stable transfectants. In brief, Phoenix Ampho retrovirus packaging cells from the ATCC were transfected with pRevTRE  $\beta$ 1A/ $\beta$ 1D via calcium phosphate. The mouse  $\beta$ 1D or mouse  $\beta$ 1A cDNA was cloned downstream of the tetracycline responsive element and the minimal immediate early promoter of cytomegalovirus ( $P_{\min}$ CMV). Retroviral supernatants were then used to infect  $\alpha$ 7-expressing cells. To test the regulation of expression, cells were cultured with various concentrations of doxycycline. After retroviral induction, cells were screened by FACS with mAb MB1.2 against mouse  $\beta$ 1 ectodomain, and cell lines with high expression of mouse  $\beta$ 1 integrin in the absence of doxycycline and low background in the presence of doxycycline were selected for further study. MCF-7 Tet-Off cells with human  $\alpha$ 7X1/mouse  $\beta$ 1A, human  $\alpha$ 7X1/mouse  $\beta$ 1D, human  $\alpha$ 7X2/mouse  $\beta$ 1A, or human  $\alpha$ 7X2/mouse  $\beta$ 1D were designated HuX1/A, HuX1/D, HuX2/A, and HuX2/D,

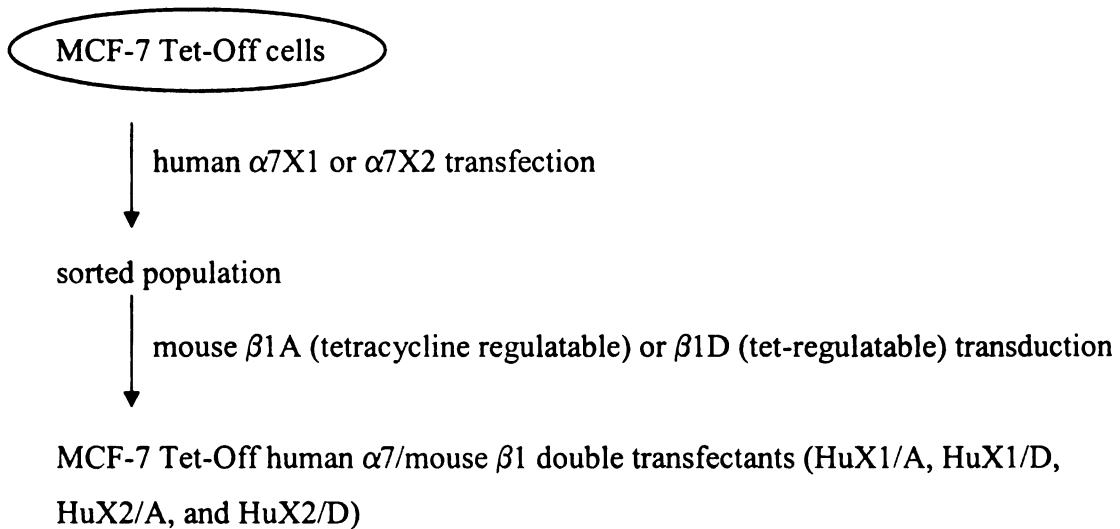
**Flow-Chart I.**

**EXPERIMENTAL APPROACH:**

**System 1**



**System 2**



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### *Flow Cytometry*

Standard procedures for flow cytometry were followed (Yao *et al.*, 1996a). Briefly, cells ( $10^6$ /ml) were incubated with predetermined optimal concentrations of primary antibodies, washed, and incubated with secondary FITC-conjugated fluorescein-labeled antibodies (affinity-purified goat anti-mouse or anti-rat antibodies; Jackson ImmunoResearch Laboratories). After washing, the cells were stained with propidium iodide and processed for flow cytometry on a FACScan (Becton Dickinson, San Jose, CA). Samples without primary or secondary antibody were always included as a control. Data are expressed as the mean fluorescence intensity (MFI) after subtraction of background staining produced by secondary antibody alone (<5% of signal). The concentration of primary and secondary antibodies was indicated at the beginning of material and methods section.

### *Cell Adhesion Assay*

Cell attachment was measured using a published protocol (Yao *et al.*, 1996a; Ziober *et al.*, 1997). Briefly, microtiter plates (96-well Immulon 1B plates; Thermo Labsystems, Franklin, MA) were coated with matrix proteins at the indicated concentrations in PBS for 1 h at 37°C. Single-cell suspensions were prepared and assayed in triplicate in 96-well plates with an incubation period of 30 to 60 min at 37°C. Adherent cells were fixed, stained with crystal violet, and solubilized in 2% SDS. Absorbance was read at 562 nm. Background cell adhesion to 1% BSA-coated wells (usually <5% of value) was subtracted from all readings. The effect of specific blocking antibodies was tested by preincubating the cells with an optimal blocking concentration of mAb on ice for 30 min

prior to the assay. Optional concentrations of the blocking antibodies were predetermined and used as previously described (Yao *et al.*, 1996a; Ziober *et al.*, 1997). The concentrations of integrin function-blocking mAb were as follows: 10  $\mu\text{g/ml}$  of purified GoH3 (anti- $\alpha 6$ ) and J143 (anti- $\alpha 3$ ). VM1 (anti-human  $\alpha 2$ ) ascites were used at a dilution of 1:400. Both CY8 (anti-mouse  $\alpha 7$ ) and AIIB2 (anti-human  $\beta 1$ ) ascites were used at a dilution of 1:300. Cytochalasin D was purchased from Sigma-Aldrich (St. Louis, MO) and dissolved in dimethyl sulfoxide (DMSO) at 5 mg/ml. Adhesion assays included control cells treated with DMSO alone (final concentration, 0.2%). The cells were processed as described above except that they were pretreated with 20  $\mu\text{M}$  cytochalasin D or DMSO alone for 20 min on ice before the assay was begun.

#### *Immunoprecipitation and Western Blotting of Cell Lysates*

Cell lysates were extracted with lysis buffer (1% Triton-100 in 50 mM Tris-HCl, pH 7.5, 1 mM  $\text{CaCl}_2$ , with 2 mM phenylmethylsulfonyl fluoride, 1 mM *N*-ethylmaleimide, and 2 mM leupeptin as protease inhibitors) for 90 min and centrifuged. The supernatants were precleared with agarose beads coupled with secondary antibodies overnight. For the purpose of depletion, three aliquots of 600  $\mu\text{g}$  of total protein of cell lysates were immunoprecipitated with antibody against  $\alpha 2$ ,  $\alpha 3$ , or  $\alpha 7$  for three consecutive rounds. The beads were pooled and washed with wash buffer (50 mM Tris-HCl, pH 7.4, 150 mM NaCl, 1 mM  $\text{MgCl}_2$ , 0.5% Nonidet P-40, and 0.1% BSA) three times and heated at 100 $^{\circ}\text{C}$  in SDS sample buffer. Samples were processed for SDS-PAGE under nonreducing conditions and then transferred to Immobilon-P membrane (Millipore, Bedford, MA). The membranes were incubated with rabbit antibody against



human integrin  $\beta 1$  (pAb 22778) (Martin *et al.*, 1996; Yao *et al.*, 1997) or mouse  $\beta 1$  (mAb 2000; Chemicon). In a different approach to assess the  $\alpha 7$  preference to  $\beta 1A$  or  $\beta 1D$ , equal amounts of cell lysate were immunoprecipitated with human  $\beta 1(A)$  or mouse  $\beta 1(D)$  antibody for three consecutive rounds. The samples were then processed as described above except under reducing conditions for immunoblotting and probed with rabbit antibody (pAb 1211) against mouse  $\alpha 7$  (Yao *et al.*, 1997).

### *Quantitative PCR*

HPLC purified salt-free primer for the target gene mouse  $\alpha 7X1$  (forward primer, GCAAGACCCCGCCTC; reverse primer, CTATCCTTGCGCAGAATGAC), for mouse  $\alpha 7X2$  (forward primer, ACCCTGCTGACCGGCTC; reverse primer, same as X1) and for total  $\alpha 7$  (forward primer, AACCAATGGCTGGGAGTCAG; reverse primer, ATCCCGAGTCTCAAAGCCT) were generated commercially (BioMolecular resource, UCSF). These primers were designed so that the melting temperature was  $10^0$  C lower than the specific probes and were optimized with regard to AmpliTaq Gold (ABI) and  $MgCl_2$  concentration at  $60^0$  C.

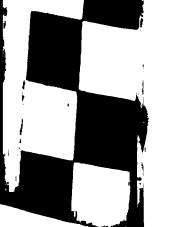
Total RNA was extracted from C2C12 myoblasts (ATCC) as described previously. Isolated total RNA integrity was electrophoretically verified by ethidium bromide staining and the purity was checked by an average optical density (OD)  $OD_{260}/OD_{280}$  nm absorption ratio of 1.8. 5  $\mu g$  of total RNA was reverse transcribed with 250 U MuL $\nu$  reverse transcriptase (Gibco-BRL) in a volume of 100  $\mu L$ , using 5  $\mu M$  random hexamer primers (Gibco) according to following protocol: 25  $^0$ C 10 second, 48

$^{\circ}\text{C}$  40 second,  $95^{\circ}\text{C}$  5 second, with dNTP 1 mM,  $\text{MgCl}_2$  7.5 mM and RNase inhibitor. cDNA of 25 ng and three 10 times serial dilutions were used to check the efficiency of probes for quantitative PCR. FAM(5'-) and TAMRA(3'-) tagged probes were commercially generated from ABI (Athens, GA). The probe specific for mouse  $\alpha 7\text{X1}$  was: TCCCGGTCCCTGCCAACAGC. The probe for mouse  $\alpha 7\text{X2}$  was: AGGACCAGCCGGAGACTTGACCTTG. The probe for total  $\alpha 7$  was: CAAGATTGTTACGTGTGCACACCGATATG. For all of the quantitative PCR in this experiment, relative expression was obtained in comparison to mouse GUS gene expression, thus eliminating the operator loading error for the differences in the amount of cDNA added to each reaction. Real-time master mix was prepared as follows:  $\text{MgCl}_2$  (7.5 mM), forward primer (480 nM), reverse primer (480 nM), probe (120 nM), dNTP (200  $\mu\text{M}$ ), 50 ng cDNA and 1.3 U AmpliTag Gold per 50  $\mu\text{L}$  reaction in buffer A. Each sample was in triplicate. The machine and software for quantitative PCR was PRISM 7700 and Sequence Detection Software version 1.6 (ABI). A four step experimental run protocol was used: (i) initial denaturation program (10 min at  $95^{\circ}\text{C}$ ); (ii) amplification and quantification program repeated 40 times (15 second at  $95^{\circ}\text{C}$ ; 60 second at  $60^{\circ}\text{C}$ ); (iii) melting curve program ( $60\text{-}95^{\circ}\text{C}$  with a heating rate of  $1^{\circ}\text{C}$  per second and a continuous fluorescence measurement); (iv) cooling program down to  $4^{\circ}\text{C}$  at the end of the 40th cycle. For the determination of relative expression from mRNA it is necessary to determine the threshold cycle (Ct) for each transcript. The Ct is defined as the point (in terms of cycle number) at which the fluorescence rises appreciably above the background fluorescence. Usually, this value is determined by the software automatically.

### *Growth curve analysis*

The MCF-7 Tet-Off double transfectants were kept in the media containing doxycycline (10 ng/ml). The day before the start of growth curve analysis, these cells were seeded at 6000 cells per well in the 96-well tissue culture plates. For each cell line, a triplicate was kept in doxycycline while the other triplicate was grown in the absence of doxycycline. The medium was replenished every day. At a specific time point, the cells were gently washed and fixed with 0.5 % formaldehyde. At the end of the experiment, all sample plates were gently washed, stained with crystal violet, and solubilized in 2% SDS. Absorbance was read at 562 nm as described in adhesion assay.

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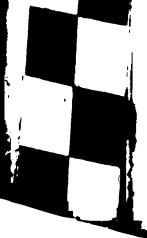
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## Characterization of $\beta 1D$ -expression cells

### *Generation of $\beta 1D$ -expression cells*

To test the hypothesis that alternatively spliced amino acid sequences in the  $\beta 1$  cytoplasmic domain may regulate the activity of the  $\alpha 7\beta 1$  integrin, we transfected human MCF-7 cells with mouse  $\alpha 7X1$  alone (X1/A-p cell line) or with both mouse  $\alpha 7X1$  and mouse  $\beta 1D$  cDNA (X1/D cell line). Following selection, the transfectants expressed high levels of  $\alpha 7X1$  and  $\beta 1D$  at their surface, as detected by flow cytometry using a mAb directed against  $\alpha 7$  (CY8) or mouse  $\beta 1$  (MB1.2) (Table II). Besides the population of double transfectants, a series of clones were also isolated by serial dilution and by single cell FACS sort (designated as X1/D-clone number). We also acquired high  $\alpha 7X1$ -expressing cells by sorting the high expresser of  $\alpha 7X1$  by FACS using the antibody against  $\alpha 7$  in X1/A-p cells. The cells derived were designated as X1/A. We previously reported that MCF-7 cells expressing  $\alpha 7X2$  with the endogenous human  $\beta 1A$  partner adhered readily to laminin-1, so this cell line was included in this study and was designated as X2/A. MCF-7 cells transfected with mouse  $\beta 1D$  or with  $\beta 1A$  were generated for control purposes and were designated as  $\beta 1D$  cells and  $\beta 1A$  cells, respectively.



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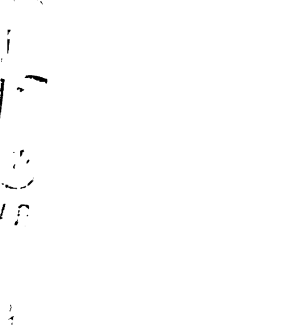
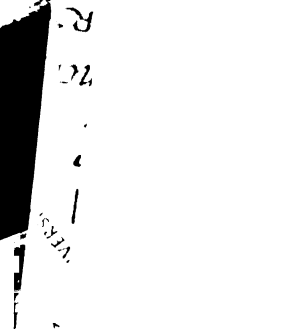
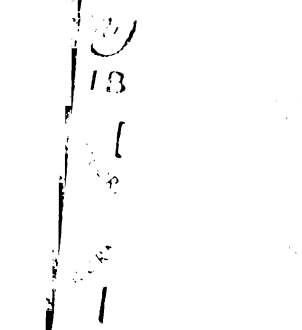
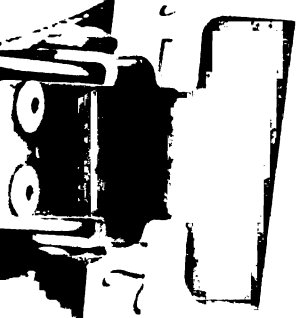
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*The expression of  $\beta 1D$  specifically augments the expression of  $\alpha 7X1$*

The surface expression profile of potential integrin receptors for laminins in MCF-7 cells and single or double transfectants is shown in table II. The expression of  $\beta 1D$  usually augments the expression level of  $\alpha 7X1$  as evidenced by FACS analysis. It is necessary to increase the expression level of  $\alpha 7X1$  in the single transfectant so that the level of  $\alpha 7X1$  in X1/A cells is comparable to that in X1/D cells. This is mandatory so that the adhesive properties can be compared. After enrichment of the expression of  $\alpha 7X1$ , the  $\alpha 7X1$  level of X1/A cells was comparable to that of X1/D cell as evidenced by FACS scan (Figure 4). Other  $\alpha$  subunits also varied; however, the variation was normally distributed and no specific trend was noticed (Appendix I). The morphology and growth rates were not changed noticeably.

*The expression of  $\beta 1D$  does not affect the general adhesive properties of  $\alpha 7X1$*

To test whether X1/D and X1/A cells displayed different adhesion profiles, we used standard adhesion assays to examine their ligand binding to the extracellular matrix components vitronectin and type I collagen. On these immobilized ligands, X1/D and X1/A cells showed a similar ligand dose-response pattern (Figure 5 A, B) over a range of ligand-coating concentrations. Thus, expression of  $\beta 1D$  did not alter the adhesion of  $\alpha 7X1$ -expressing cells to these components. Previous reports have shown that in MCF-7 cells, adhesion to vitronectin and type I collagen was mediated primarily by  $\alpha V\beta 1$  (Maemura *et al.*, 1995) and  $\alpha 2\beta 1$  integrins (Jones *et al.*, 1995; Maemura *et al.*, 1995),

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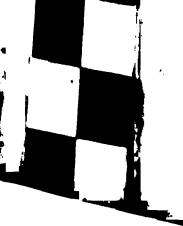
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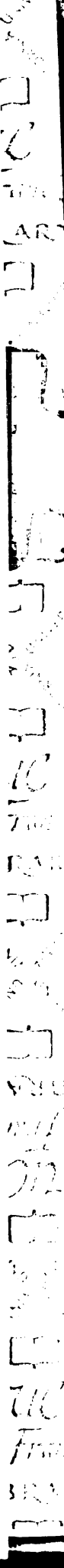
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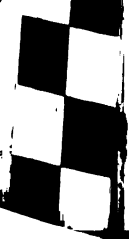




respectively. This result indicated that there was no change in the general adhesion activity toward type I collagen or vitronectin in the transfectants. The same procedures were carried out in one of the clones, X1/D-27, and the results were the same (Figure 4 and Figure 6 A, B).

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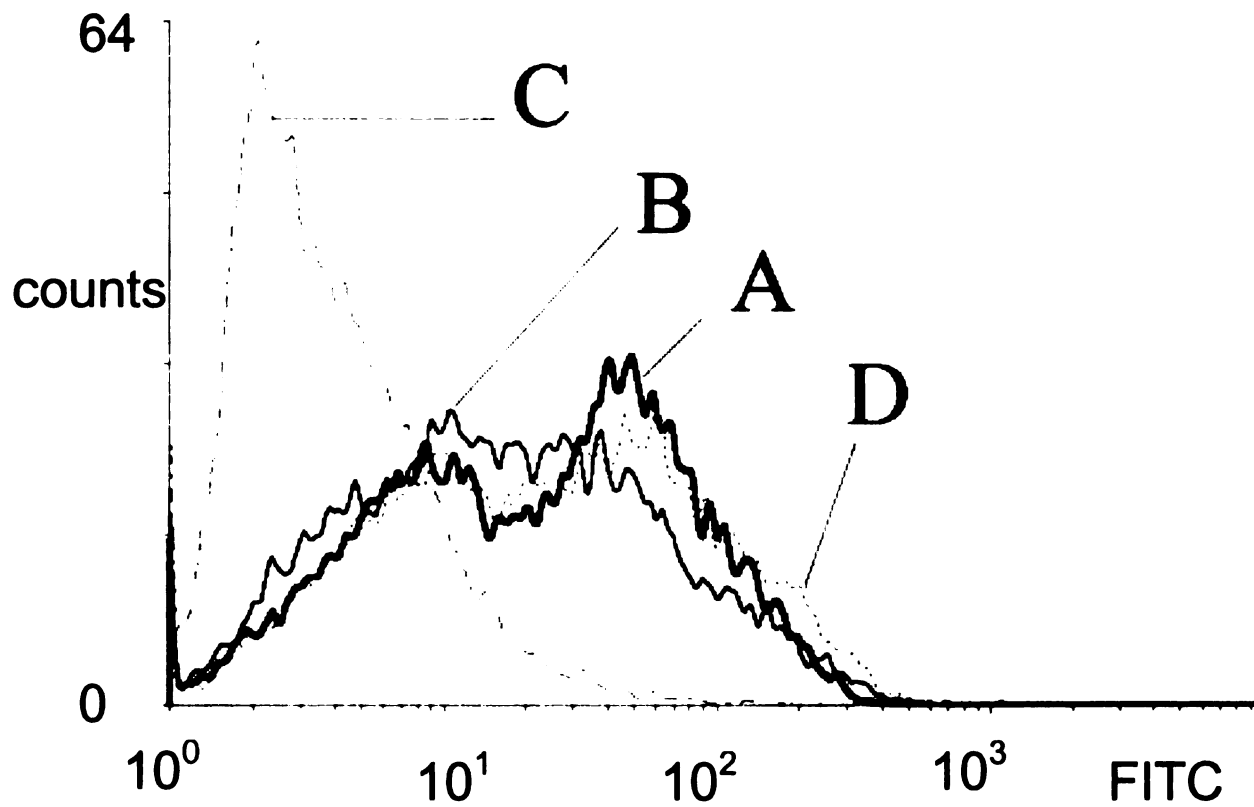
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**Figure 4.**  $\alpha$ 7X1 expression profile in MCF-7  $\alpha$ 7X1 and  $\beta$ 1D transfectants



- A. X1/A cells
- B. X1/D cells
- C. Control sample
- D. X1/D-27 cells

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**Table II.** Surface expression levels of  $\alpha 7/\beta 1$  in MCF-7 and transfectants

Cell line	Integrin pair <sup>*</sup>	Mean fluorescence intensity		
		$\alpha 7^a$	Hu $\beta 1A^b$	Ms $\beta 1D^c$
MCF-7	— $\beta 1A$	—	78	—
X1/A	Ms $\alpha 7X 1/\beta 1A$	43	86	—
X1/D	Ms $\alpha 7X 1/\beta 1D$	32	55	20
X2/A	Ms $\alpha 7X 2/\beta 1A$	40	58	—

<sup>\*</sup>The parental MCF-7 cells and transfectants were processed for FACS analysis as described in Material and Methods section. The Hu $\beta 1A$  is the endogenous integrin partner.

<sup>a</sup>mAb CY8, rat anti-mouse  $\alpha 7$ .

<sup>b</sup>mAb AIIB2, rat anti-human  $\beta 1$ .

<sup>c</sup>mAb MB1.2, rat anti-mouse  $\beta 1$ .

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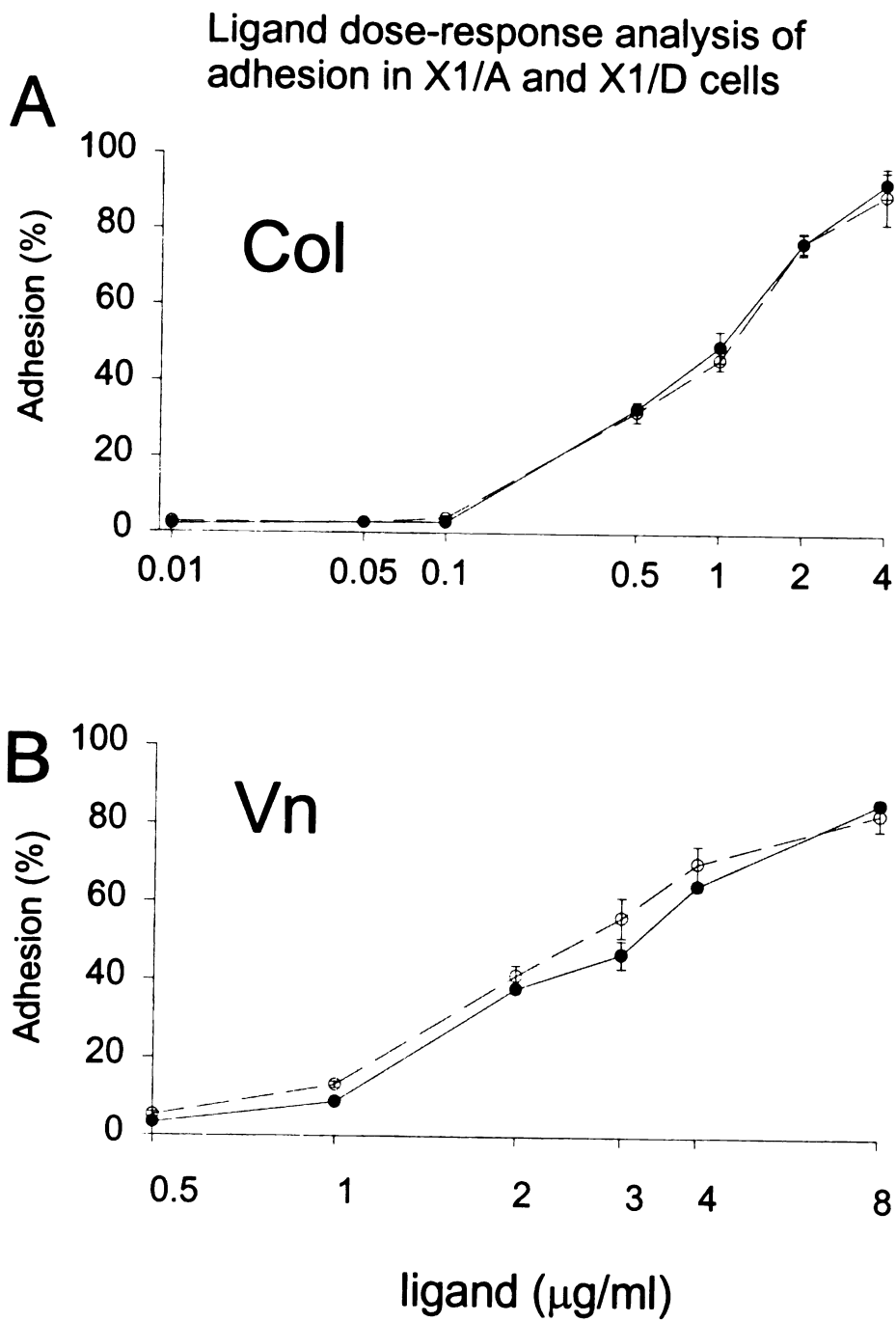
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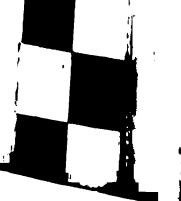
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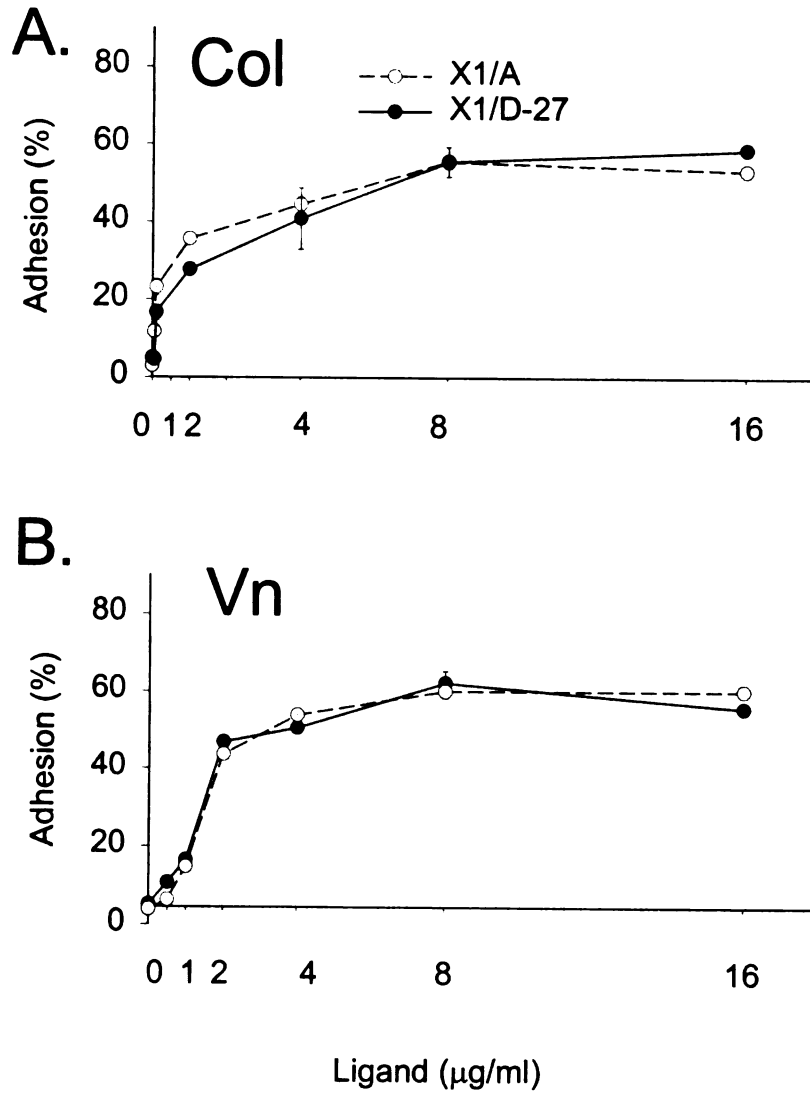
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**Figure 6.**

Ligand dose-response analysis of adhesion in X1/A and X1/D-27 cells



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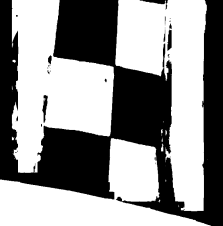
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## **There is no preference of $\alpha 7$ X1 in coupling with human $\beta 1$ A or mouse $\beta 1$ D**

### *Analysis of Integrin Complexes in the $\beta 1$ D Transfectants*

To determine what fraction of the  $\alpha 7$  subunit was partnered with endogenous  $\beta 1$ A or transfected  $\beta 1$ D, two different strategies were used. In the first, lysates prepared from X1/D cells were exhaustively immunoprecipitated with a mAb specific to either human  $\beta 1$  or mouse  $\beta 1$ , followed by immunoblotting for the partner  $\alpha 7$  (Figure 7A). The results show that the fraction of  $\alpha 7$  that heterodimerized with either the  $\beta 1$ D (lane 5) or the  $\beta 1$ A (lane 6) isoform was approximately equal. Thus, neither subunit appears to have any preferential affinity for  $\alpha 7$  during integrin heterodimerization.

In the second approach to estimate the relative pairing of the  $\alpha 7$  integrin subunit with endogenous  $\beta 1$ A or transfected  $\beta 1$ D, we used the X1/D-27 cell line which expresses high levels of both  $\alpha 7$ X1 and  $\beta 1$ D (Figure 7B). Cells were first processed for immunoprecipitation with anti- $\alpha 7$  antibody, followed by immunoblotting analysis with an antibody specific for either the human  $\beta 1$ (A isoform) or mouse  $\beta 1$ (D isoform). Again, similar levels of  $\beta 1$ A and  $\beta 1$ D were detected in the immunoprecipitates (Figure 11B, lane 3). There appeared to be an abundance of  $\beta 1$ D precursor (Figure 11B, lane 1; right panel) compared to the mature, fully processed form of the endogenous  $\beta 1$ A subunit. In contrast, nearly all of the endogenous  $\beta 1$ A subunit was present in the mature form (Figure 11B, lane 1; left panel).  $\alpha 2$  and  $\alpha 3$  subunits were associated with similar

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proportions of  $\beta$ 1A and  $\beta$ 1D (Figure 7C). In other studies, we examined the amount of  $\alpha$ 2 and  $\alpha$ 3 subunits associated with  $\beta$ 1A and  $\beta$ 1D by immunoprecipitation of  $^{35}\text{S}$ -methionine-labeled cells and similar results were observed. On the basis of these studies, we conclude that  $\alpha$ 7 as well as  $\alpha$ 2 and  $\alpha$ 3 subunits is able to complex with similar efficiencies to the  $\beta$ 1A and  $\beta$ 1D subunits.

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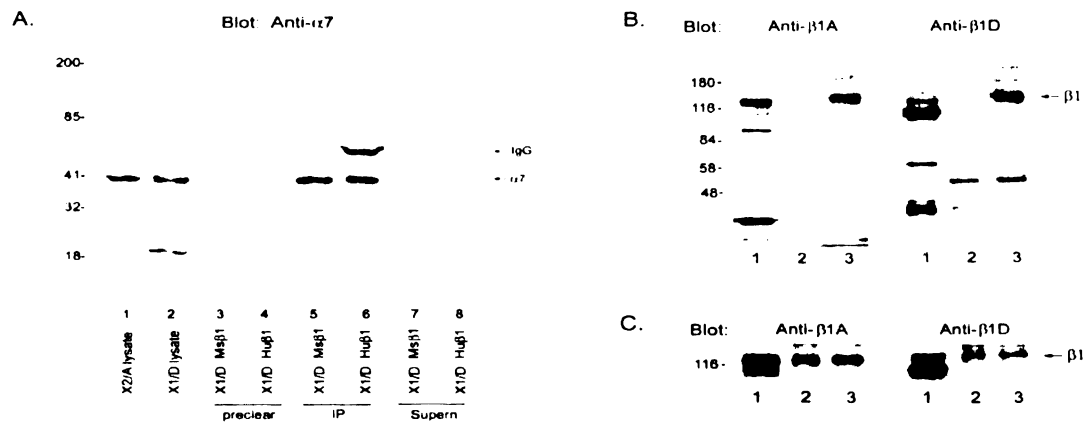
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**Figure 7.** Association of  $\alpha 7$  with  $\beta 1D$ .



(A) Immunodepletion of lysates with antibody against  $\beta 1$ . To determine the relative association of  $\alpha 7$  with human  $\beta 1$  or with mouse  $\beta 1$ , X1/D double transfectant cells were processed for immunodepletion experiments using antibodies against the  $\beta 1D$  (mouse) or  $\beta 1A$  (human) subunit. Cell lysates were prepared from X2/A cells (lane 1) or X1/D cells (lane 2) as controls for  $\alpha 7$  expression. X1/D cell lysate was subjected to a preclearing control (beads and secondary antibody only, lanes 3 and 4). X1/D cell lysate was exhaustively precipitated with mAb against mouse  $\beta 1$  (mAb MB1.2, lane 5) or against human  $\beta 1$  (mAb AIIB2, lane 6), and after depletion with mouse  $\beta 1$  (lane 7) or human  $\beta 1$  (lane 8) of X1/D cell lysates, supernatants were subjected to a final additional round of immunoprecipitation to confirm depletion. Immunoprecipitates were analyzed by SDS-PAGE under reducing conditions, followed by Western blotting with anti- $\alpha 7$  antibody and visualization by chemiluminescence. The position of the 38-kDa  $\alpha 7$  light chain is

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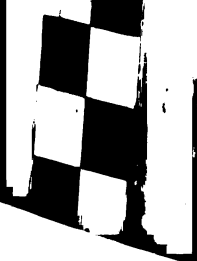
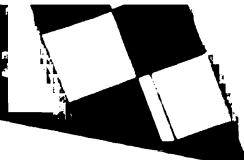
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indicated. (B) Immunodepletion of lysates with antibody against  $\alpha 7$ . Cell lysate from X1/D-27 cells (a clone of X1/D cells) was immunoprecipitated with a mAb to  $\alpha 7$ . Immunoprecipitates were processed for immunoblotting under nonreducing conditions and were probed with rabbit antibodies specific to  $\beta 1A$  (left) or  $\beta 1D$  (right) and by chemiluminescence. Lane 1, whole-cell lysate; lane 2, control anti-rat IgG-beads; lane 3, anti- $\alpha 7$ . The position of the mature  $\beta 1$  subunit is indicated. (C) Same procedures as in (B) except that the lysates (lane 1) were processed for immunoprecipitation with anti- $\alpha 2$  integrin (mAb VM1, lane 2) and anti- $\alpha 3$  integrin (mAb J143, lane 3) and blotted with antibodies against  $\beta 1A$  (left) or  $\beta 1D$  (right).

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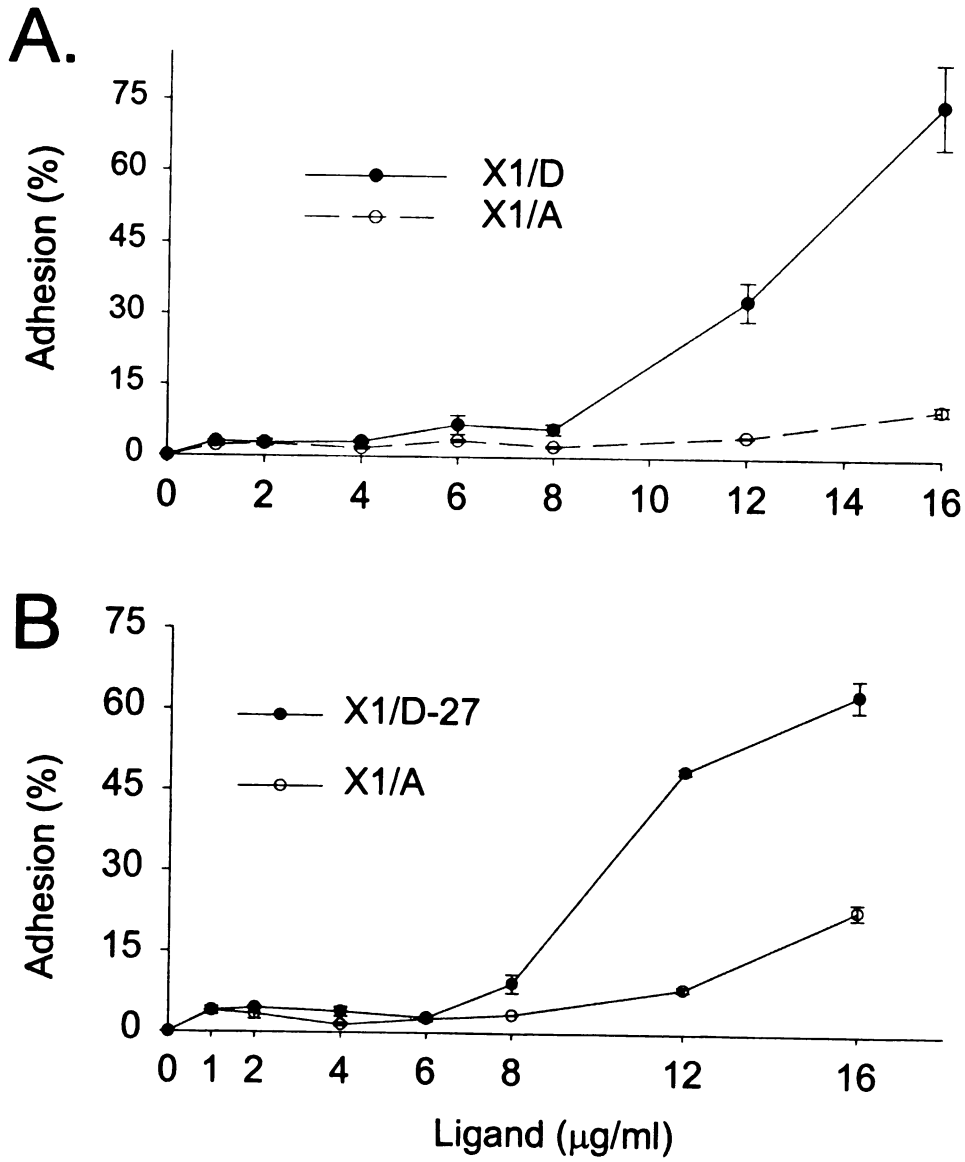
## **The expression of $\beta$ 1D specifically enhanced the adhesion of $\alpha$ 7X1 to laminin-1**

### *The expression of $\beta$ 1D enhanced the adhesion of $\alpha$ 7X1 to laminin-1*

We previously reported that MCF-7 cells expressing  $\alpha$ 7X1 with the endogenous human  $\beta$ 1A partner adhered poorly to laminin-1 (Ziober *et al.*, 1997). However, following transfection with the cDNA of the mouse  $\beta$ 1D subunit, X1/D cells acquired the capacity to strongly bind to laminin-1 (Figure 8A). This increase in adhesion was evident at coating concentrations above 8  $\mu$ g/ml. We also generated 20 single cell clones by limiting dilution and by single-cell sorting. A representative clone (X1/D-27) was chosen for the same assay, and the results were similar (Figure 8B). Since the stimulatory effect of  $\beta$ 1D on adhesion was restricted to laminin-1 and not vitronectin or collagen I, this suggested that laminin-binding integrins such as  $\alpha$ 7X1, but not other potential  $\beta$ 1 partner subunits, were involved. The poor adhesion of X1/A cells to laminin-1 was not related solely to the level of  $\alpha$ 7X1 expressed at the surface, since this cell line had a significantly higher expression level of  $\alpha$ 7X1 than did X1/D cells (Table II).

Figure 8.

Ligand dose-response analysis of X1/A, X1/D and X1/D-27 on laminin-1



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*Enhanced Adhesion of  $\beta$ 1D Transfectants to Laminin-1 Is  $\alpha$ 7 Specific*

MCF-7 expresses several integrin subunits that are potential receptors for laminin-binding (appendix I). To establish that the  $\beta$ 1D-induced increase in laminin-1 adhesion was attributable specifically to  $\alpha$ 7X1, we performed standard adhesion assays with blocking mAbs against other expressed  $\alpha$  subunits (Figure 9). The  $\alpha$ 7X1-expressing cells (X1/A) showed weak adhesion to laminin-1, and adhesion could be diminished further by treatment with a cocktail of blocking mAbs to  $\alpha$ 2,  $\alpha$ 3, and  $\alpha$ 6 integrins. In contrast, the X1/D cells expressing  $\beta$ 1D showed enhanced adhesion to laminin-1, and most of the adhesion to laminin-1 was inhibited by a blocking mAb to  $\alpha$ 7. Since the  $\beta$ 1D cDNA was from mouse, the human  $\beta$ 1 functional perturbing antibody, AIIB2, did not inhibit the adhesion. The other antibody (Ha2/5) which blocks both human and mouse  $\beta$ 1 almost completely inhibits the adhesion to laminin-1.

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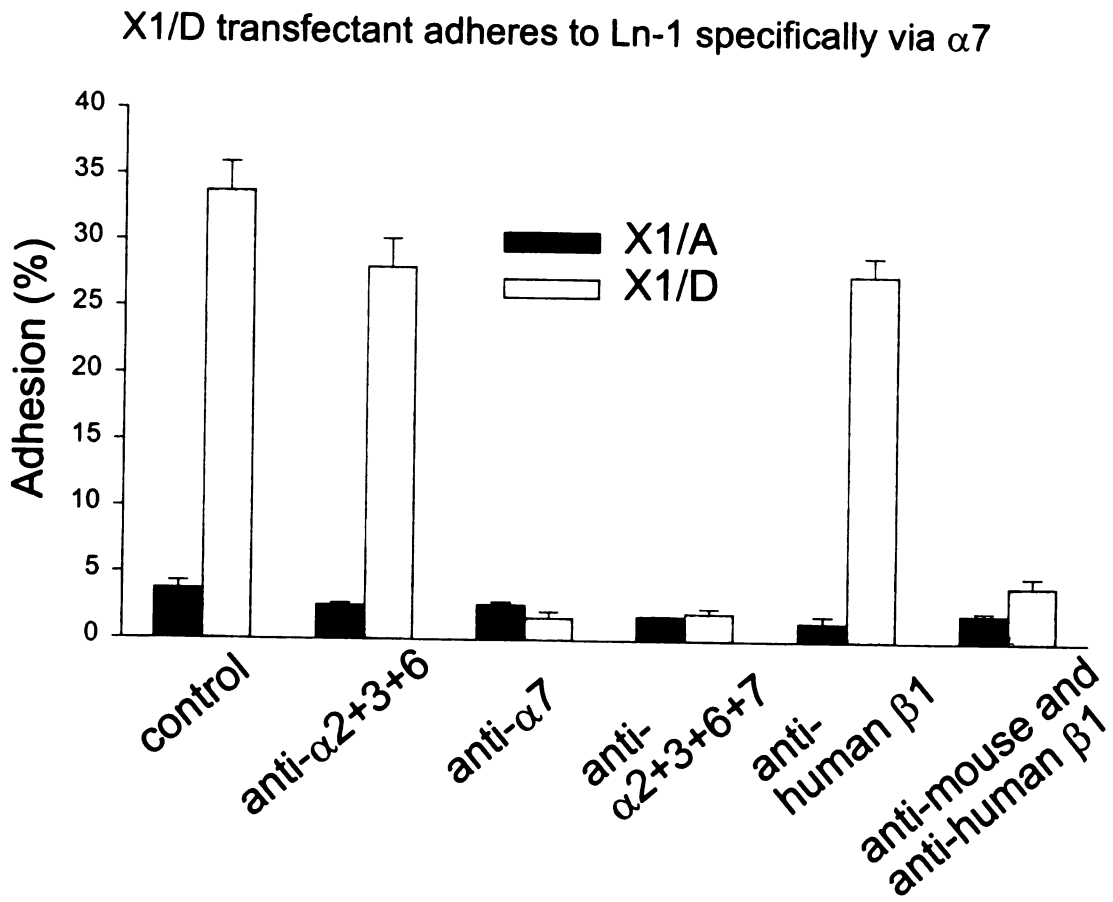
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### *Adhesion to Laminin-1 Correlates with $\alpha 7 X 1$ and $\beta 1 D$ Levels*

To further define the relationship between  $\beta 1 D$  and  $\alpha 7$  activity, we compared the relative adhesion to laminin-1 of a large series of  $\alpha 7 / \beta 1 D$ -expressing clones and cell lines. A three-dimensional scatter plot summarizing these results indicates that adhesion to laminin-1 strictly correlated not only with the expression of  $\alpha 7$ , but also with the relative level of expression of  $\beta 1 D$  (Figure 10). The parental cells with endogenous  $\beta 1 A$  (MCF-7) or cells transfected with  $\alpha 7 X 1$  alone (X1/A) both showed poor adhesion to laminin-1. The  $\alpha 7 X 1$  and  $\beta 1 D$  double transfectants (X1/D and clones 10, 12, 13, 17, 21, and 27) showed a strong correlation between levels of  $\alpha 7$  and  $\beta 1 D$  expression. Thus, the expression of mouse  $\beta 1 D$  in MCF-7 cells transfected with  $\alpha 7 X 1$  consistently enhanced the expression of  $\alpha 7$  in this system. The level of  $\alpha 7$  increased in parallel with that of the transfected mouse  $\beta 1 D$  and correlated with correspondingly greater adhesion to laminin-1. The surface expression level of relevant integrins was shown in appendix I. Furthermore, transfection of cells with mouse  $\beta 1 A$  or  $\beta 1 D$  in the absence of  $\alpha 7$  to yield the  $\beta 1 A$  or  $\beta 1 D$  cell lines, did not result in enhanced adhesion to laminin-1. This result indicated that the overexpression of mouse  $\beta 1 A$  or  $\beta 1 D$  was unable to augment adhesion activity of endogenous  $\alpha 2$ ,  $\alpha 3$ , or  $\alpha 6$  integrins for laminin-1. In other studies, we generated human  $\alpha 7 X 1$ - and  $\alpha 7 X 2$ -expressing MCF-7 Tet-Off cells and found that these transfectants showed a similar adhesion profile to laminin-1, with  $\alpha 7 X 1$  showing poor binding to laminin-1 and  $\alpha 7 X 2$  showing strong binding (Figure 11).

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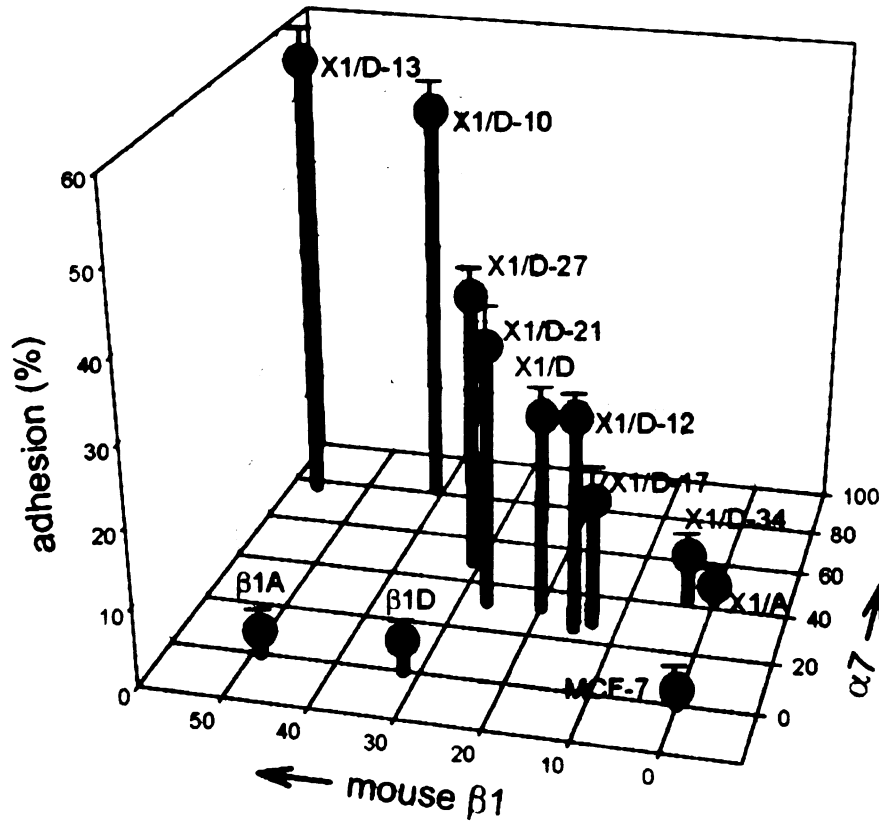


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**Figure 10.** Adherence to laminin-1 correlates with the surface expression level of  $\beta$ 1D



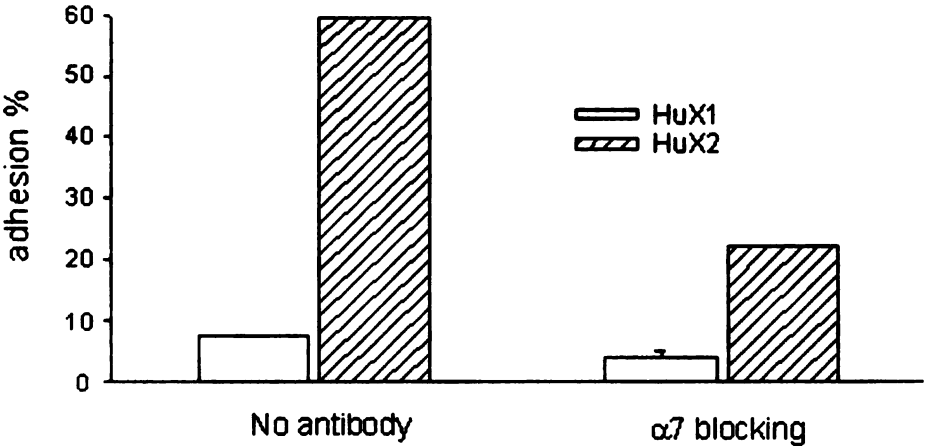
MCF-7 transfectants and clones derived from X1/D double transfectants was compared in standard adhesion assay as described in Figure 5. Parental cells (MCF-7), transfectants expressing  $\alpha$ 7X1 (X1/A), mouse  $\beta$ 1A ( $\beta$ 1A), or mouse  $\beta$ 1D ( $\beta$ 1D) all showed poor adhesion to laminin-1. The  $\alpha$ 7X1 and  $\beta$ 1D double transfectants (X1/D and its derived cell clones) showed increased adhesion to laminin-1 that correlated closely with the relative level of  $\beta$ 1D and  $\alpha$ 7X1 integrin expression.

**Figure 11.** Adhesion to laminin-1 in human X1 and human X2 cells

$\alpha 7$  FITC level

HuX1	114
HuX2	80

Adhesion to laminin-1 in HuX1 and HuX2 cells



### *Functionality of Human $\alpha7X1$ in Cells Expressing Tetracycline-regulatable $\beta1D$*

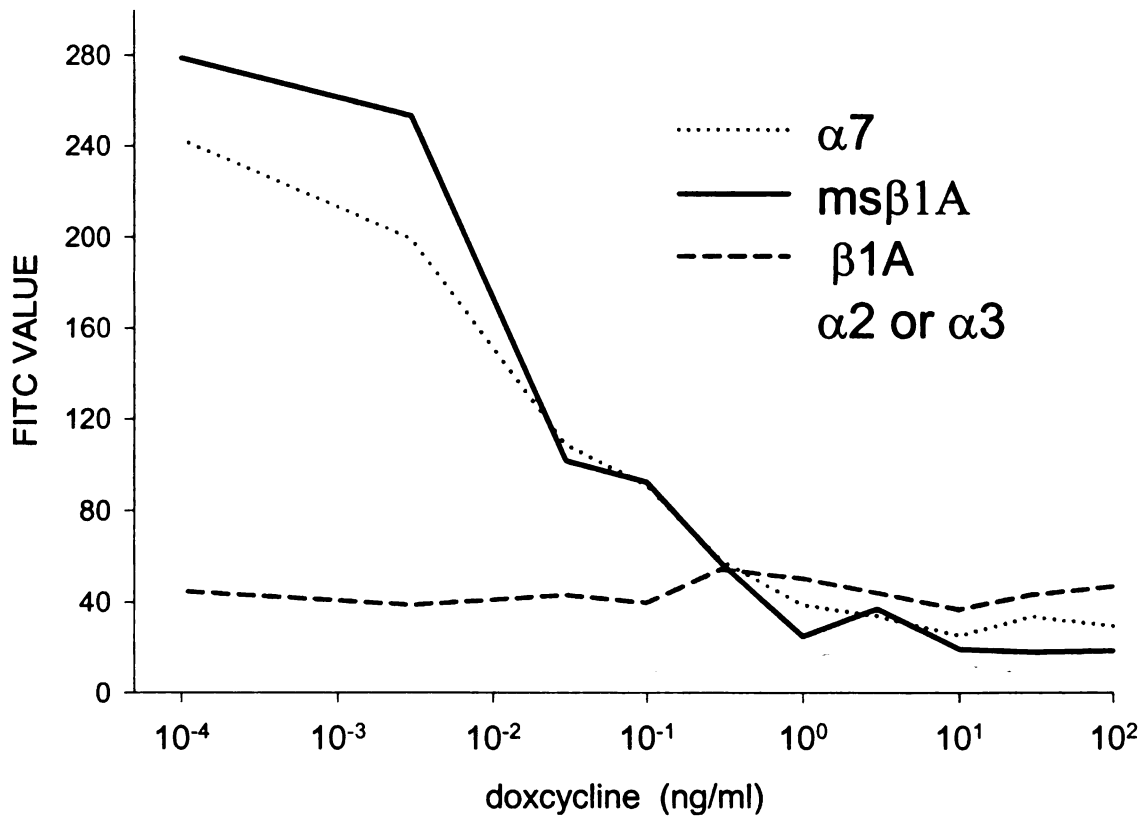
In a parallel approach to establish the role of integrin  $\beta1$  cytoplasmic variants on the expression and function of human  $\alpha7$ , we created stable cell lines constitutively expressing either the human  $\alpha7X1$  or  $\alpha7X2$  isoform and tetracycline-regulatable mouse  $\beta1A$  or  $\beta1D$  integrin. These cell lines were designated as HuX1/A, HuX1/D, HuX2/A, and HuX2/D.

### *Characterization of HuX1/A, HuX1/D, and HuX2/A cells*

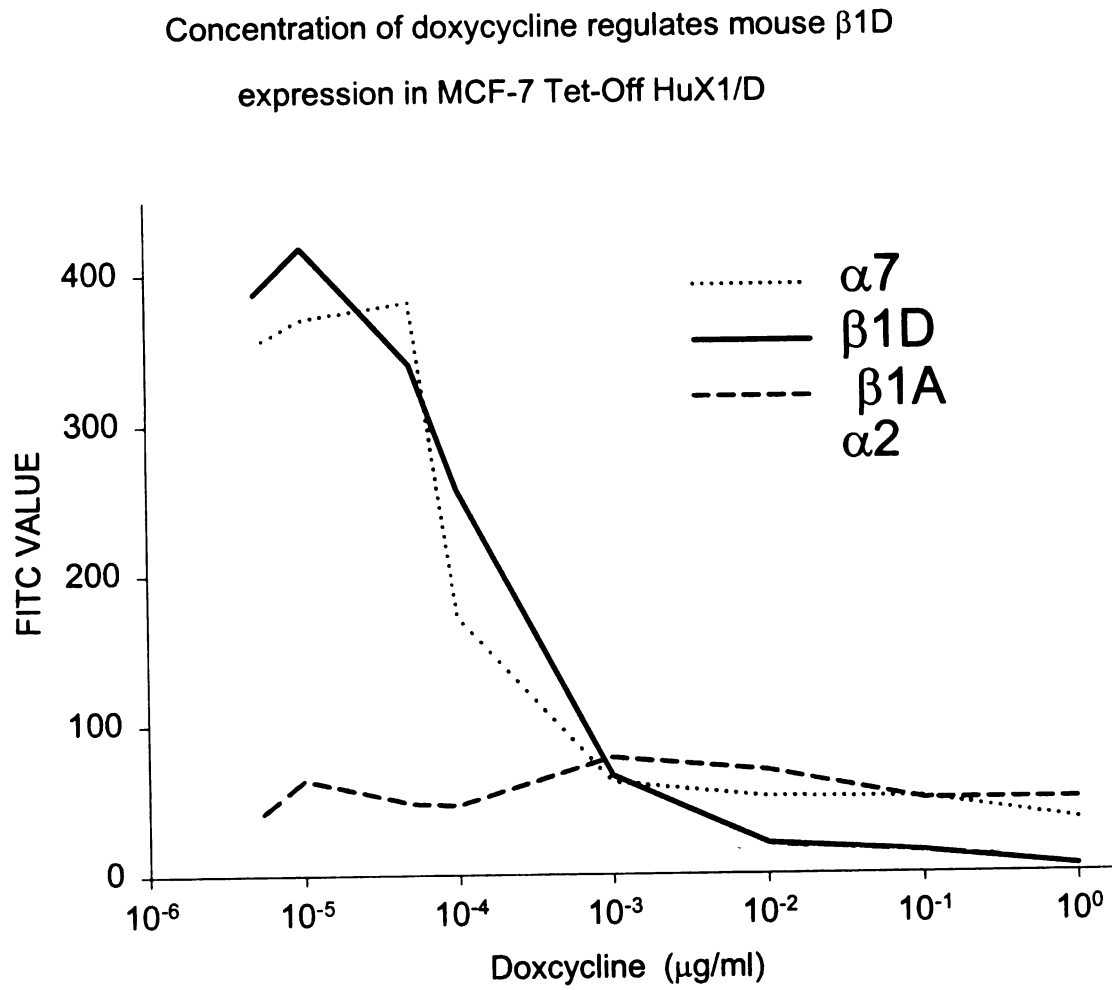
To regulate the expression of the mouse  $\beta1$  integrin subunit, cells were cultured in medium containing increasing concentrations of doxycycline, and integrin expression was then measured by FACS using mAb to  $\alpha7$  integrin and mouse  $\beta1$  integrin. Several clones of Tet-Off double transfectants were screened for doxycycline regulation of mouse  $\beta1A$  or  $\beta1D$  expression. Cells expressed high level of mouse  $\beta1$  in the absence of doxycycline and low background in the presence of doxycycline were chosen for further study. Figures 12, 13, and 14 show the dose-response curve of doxycycline in modulating regulatable mouse  $\beta1$  expression as assayed by FACS. In all three cells, the expression of human  $\alpha7$  was also modulated. If the concentration of doxycycline was further reduced, the increase in integrin expression eventually reached a plateau. Analysis of integrin profiles revealed that, besides  $\alpha7$  and  $\beta1D$  subunits, other integrins (human  $\alpha2$ ,  $\alpha3$ , and  $\beta1$ ) remained relatively stable as the regulatable mouse  $\beta1$  was modulated.

**Figure 12.**

Concentration of doxycycline regulates mouse  $\beta$ 1A expression in MCF-7 Tet-Off HuX1/A



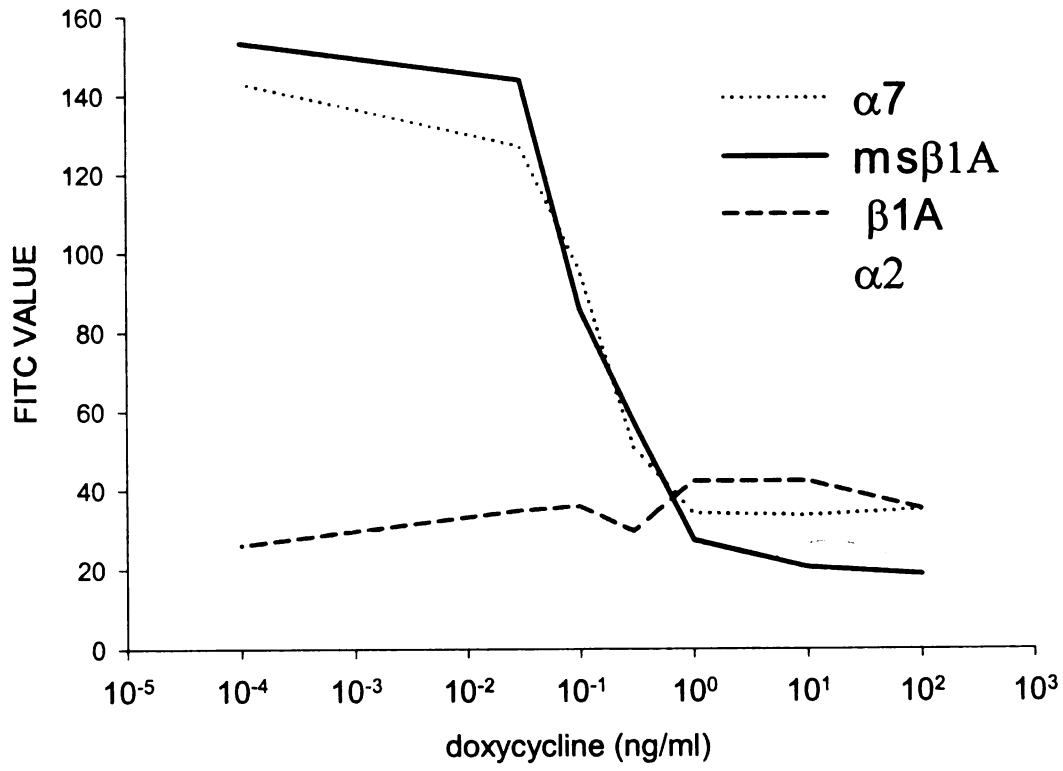
**Figure 13.**





**Figure 14.**

Concentration of doxycycline regulates mouse beta 1A expression in MCF-7 Tet-Off HuX2/A



*The tet-regulatable expression of  $\beta 1D$  but not  $\beta 1A$  enhances the adhesion to laminin-1*

Doxycycline suppressed the expression of mouse  $\beta 1$  integrin in a dose-dependent manner (Figure 15A) and at the lower range of doxycycline shown the integrin expression was inversely proportional to doxycycline concentration. The sensitivity to regulation by doxycycline repression of HuX1/A and HuX1/D differed slightly; the HuX1/D required about twice the doxycycline concentration to suppress mouse  $\beta 1$  integrin expression compared with that required for the HuX1/A. Expression of mouse integrin was completely suppressed with a doxycycline concentration of  $>1$  ng/ml for  $\beta 1D$  and 0.65 ng/ml for  $\beta 1A$ . By modulating the levels of doxycycline, comparable levels of mouse  $\beta 1$  integrin expression were reached for both cell lines (Fig. 15A). Due to the difference in sensitivity to doxycycline repression, we chose to use 0.1 and 0.3 ng/ml of doxycycline for HuX1/A and HuX1/D respectively for adhesion assays on laminin-1. When cultured at this low concentration of doxycycline, the integrin profiles of these cells were quantified as detected by FACS and is summarized in Table III. Interestingly, in the Tet-Off cells, high levels of human  $\alpha 7$  and the mouse  $\beta 1$  level are maintained as compared to that in the MCF-7 transfectants. The mouse  $\beta 1$  appears to be the dominant subunit compare to the endogenous human  $\beta 1A$ . As the level of mouse  $\beta 1A$  or  $\beta 1D$  integrin was increased following induction, we observed a corresponding increase in  $\alpha 7$  surface expression (Figure 15B). The levels of induced expression of  $\alpha 7$  were 5 to 6 times that of the background expression at high doxycycline concentrations.

After the optimization of doxycycline concentration in regulating the surface expression level of mouse  $\beta 1$ , we then analyzed the influence of  $\alpha 7X1$  and  $\beta 1$  expression on the adhesion of doxycycline-regulatable cells to laminin-1 in standard adhesion assays (Figure 15C) at the doxycycline concentration chosen above (0.1 ng/ml and 0.3 ng/ml for HuX1/A and HuX1/D respectively). Mouse  $\beta 1A$ -expressing cells showed poor adhesion at all levels of  $\beta 1A$  integrin expression even though  $\alpha 7X1$  was increased several fold over the range of doxycycline treatment (shown in Figure 15B). In contrast, induction of  $\beta 1D$  at low doxycycline concentrations resulted in a strong adhesive response to laminin-1 that corresponded to elevated  $\beta 1D$  and  $\alpha 7X1$  expression levels. In fact, the elevation in adhesion to laminin-1 correlated closely to the increase in  $\alpha 7X1$  expression.

**Table III.** Surface expression level of  $\alpha 7/\beta 1$  in MCF-7 Tet-Off double transfectants

Cell line	Integrin pair $\alpha 7/\beta 1$	Mean fluorescence intensity*		
		Human $\alpha 7^a$	Human $\beta 1A^b$	Mouse $\beta 1A/D^c$
HuX1/A	Human $\alpha 7X1$ /mouse $\beta 1A^c$	125	45	130 <sup>d</sup>
HuX1/D	Human $\alpha 7X1$ /mouse $\beta 1D^c$	110	36	120 <sup>c</sup>
HuX2/A	Human $\alpha 7X2$ /mouse $\beta 1A^d$	86	40	116 <sup>d</sup>

\*The transfected cell lines were processed for FACS analysis using the indicated antibody as detailed in Material and Methods Section.

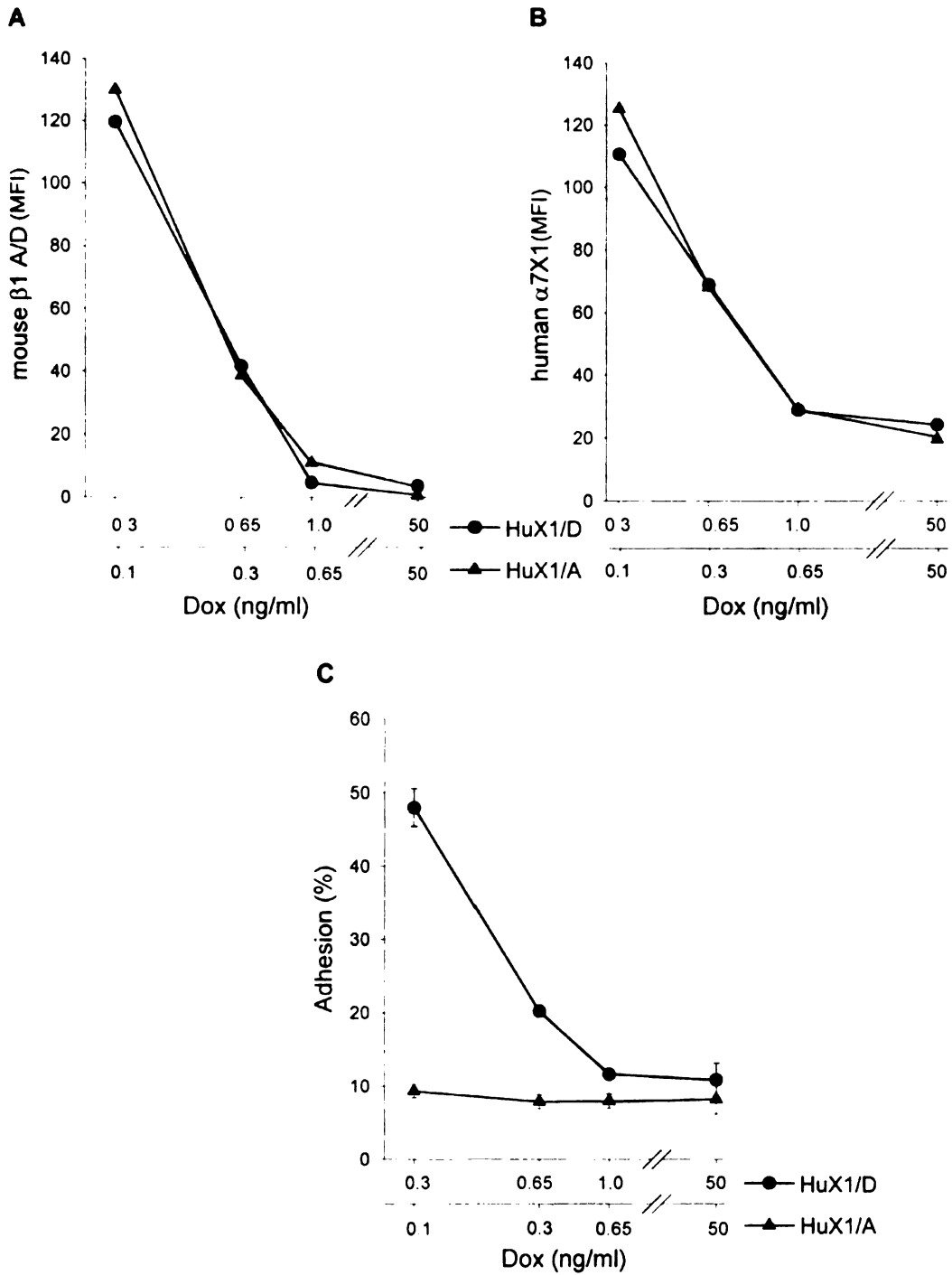
<sup>a</sup>mAb CY8, rat anti-mouse  $\alpha 7$ .

<sup>b</sup>mAb AIB2, rat anti-human  $\beta 1$ .

<sup>c</sup>mAb MB1.2, rat anti-mouse  $\beta 1$ .



**Figure 15.**  $\beta$ 1D expression promotes adhesion of human  $\alpha$ 7X1 Tet-Off cells to laminin-1



Previously we showed that  $\alpha 7X2\beta 1$  is fully functional to bind laminin-1 (Yao *et al.*, 1996a; Ziober *et al.*, 1997). Since it was shown that the  $\beta 1D$  tail binds better to talin than  $\beta 1A$  tail does (Belkin *et al.*, 1997; Pfaff *et al.*, 1998), then we asked the question whether the expression of exogenous mouse  $\beta 1D$  enhance the adhesion to laminin-1. To answer this question, we analyzed the adhesive efficiency of Tet-Off cells expressing  $\alpha 7X2/A$  and  $\alpha 7X2/D$ . After verifying the expression of  $\alpha 7X2$  with FACS analysis by the mAb 9.1, these cells were induced to express either mouse  $\beta 1A$  or mouse  $\beta 1D$  by retroviral transduction. We found no significant difference in adhesion to laminin-1 between  $\alpha 7X2$  cells expressing either  $\beta 1A$  or  $\beta 1D$  (Figure 16). Both double transfectants adhered strongly to laminin-1 (70 to 80%). The functional blocking antibody against human  $\beta 1$  did not reduce adhesion because the expression level of endogenous human  $\beta 1A$  was much lower than that of mouse  $\beta 1$  (see appendix II). This result indicated that  $\alpha 7X2\beta 1$  is fully functional for adhesion to laminin-1, and that exogenous  $\beta 1D$  did not enhance adhesion further.

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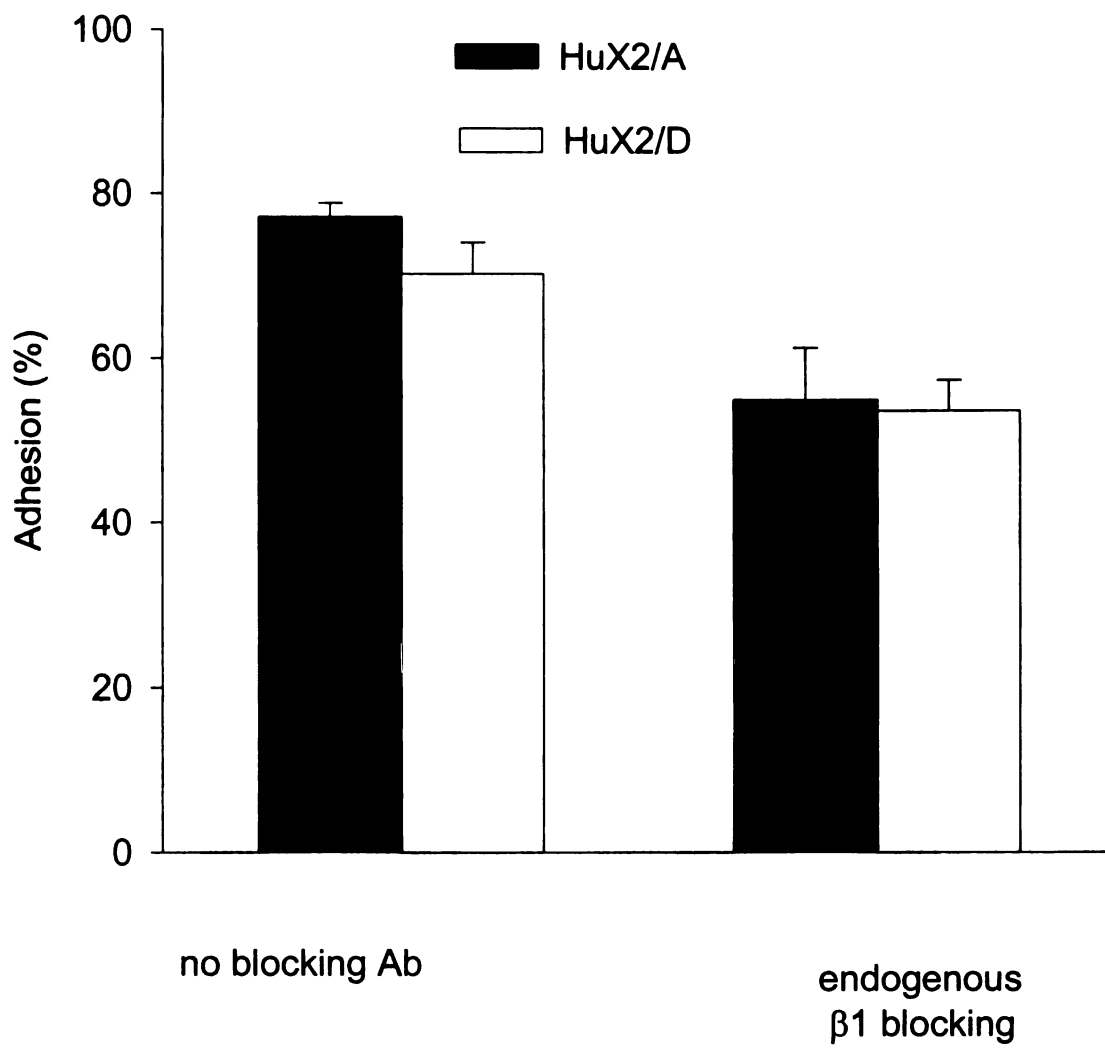
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**Figure 16.**

Adhesion to laminin-1 in HuX2/A and HuX2/D cells





### *Expression of $\beta 1D$ Does Not Alter Adhesion of $\alpha 7X1$ to Laminin-2/4 or Laminin-10/11*

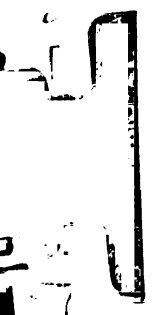
The above results indicate that  $\beta 1D$  was able to modify the binding of  $\alpha 7X1\beta 1$  to laminin-1. Recently, von der Mark and collaborators (von der Mark *et al.*, 2002) have reported that truncated forms of  $\alpha 7X1\beta 1$  and  $\alpha 7X2\beta 1$  bind differently to various laminin isoforms. We compared the relative adhesion efficiency of  $\alpha 7X1$  expressing cells in the presence of  $\beta 1A$  or  $\beta 1D$  to available laminin isoforms. We first tested the adhesion of  $\alpha 7\beta 1$ -expressing Tet-Off cells on preparations of human placental merosin. These cells have much higher expression level of  $\alpha 7$  and the levels of  $\alpha 2$ ,  $\alpha 3$ , and  $\alpha 6$  are very low (see appendix II and the data in characterization of HuX1/A, HuX1/D, and HuX2/A section). Previous analyses have indicated that these preparations contain primarily human laminin-4 ( $\alpha 2$ - $\beta 2\gamma 1$ , S-merosin) with lesser amounts of laminin-2 ( $\alpha 2$ - $\beta 1\gamma 1$ , merosin) (Delwel *et al.*, 1994; Spinardi *et al.*, 1995; Yao *et al.*, 1996a). Thus, we tested the relative adhesion of HuX1/A, HuX1/D, and HuX2/A to human merosin (laminin-2/4) (Figure 17A). Interestingly, cells expressing  $\alpha 7X1/\beta 1A$  (HuX1/A) or  $\alpha 7X1/\beta 1D$  (HuX1/D) adhered poorly. In contrast, cells expressing  $\alpha 7X2/\beta 1A$  (HuX2/A) showed strong attachment. Adhesion by HuX2/A cells to laminin-2/4 was not sensitive to a mixture of mAbs to  $\alpha 2$ ,  $\alpha 3$ , and  $\alpha 6$  integrins, but adhesion was completely blocked by anti- $\alpha 7$  mAb 9.1 (Figure 17B).

In a similar fashion, we examined the adhesive capacity of these cell lines to laminin-10/11 (Figure 18A). In contrast to adhesion on laminin-2/4 substrates, HuX2/A adhered poorly to laminin-10/11, whereas HuX1/A and HuX1/D both attached well.



Treatment of cells with anti-human  $\alpha 7$  blocking mAb effectively inhibited adhesion confirming that this receptor mediates binding (Figure 18B). Taken together, these results show that replacing  $\beta 1A$  with  $\beta 1D$  confers activity of  $\alpha 7X1$  for laminin-1 but does not alter binding of the double transfectants to laminin-2/4 or laminin-10/11 substrates.

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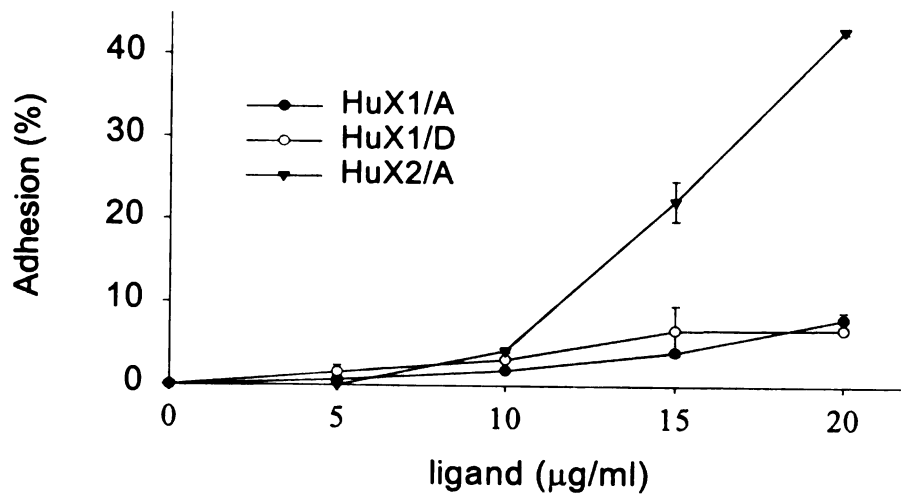
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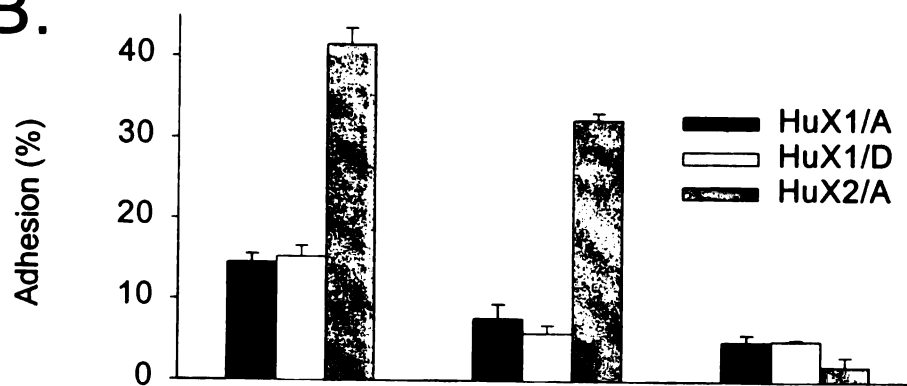
**Figure 17.**

Adhesion to LN 2/4 in MCF-7 Tet-Off transfectants

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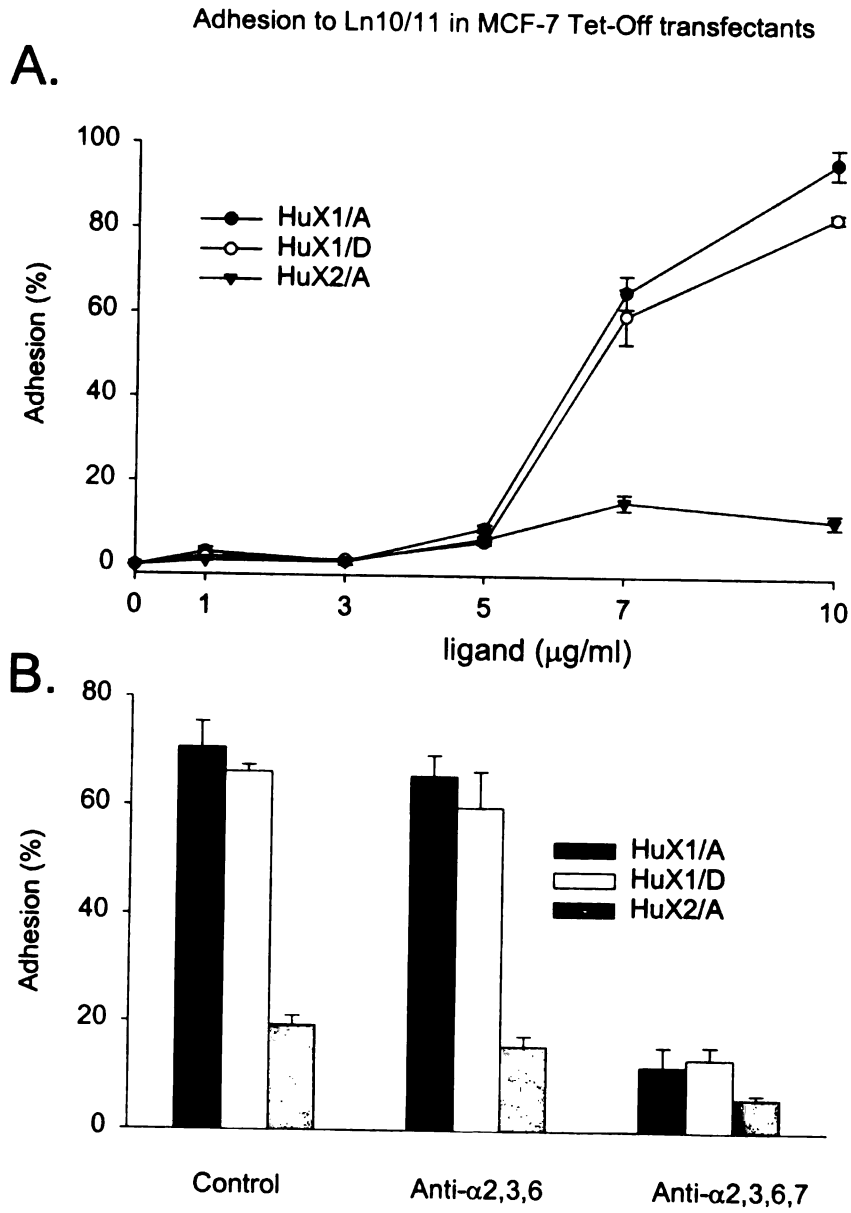
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**Figure 18.**



### *Cytochalasin D Modulates HuX1/D Adhesion to Laminin-1*

Integrin-cytoskeletal interactions have been shown to influence both integrin clustering and extracellular domain conformation. It is well established that integrins connect to the cytoskeleton through the cytoplasmic domain of the  $\beta$  subunit and evidence indicates that talin may play an important role in regulating integrin activity by binding to the  $\beta 1$  cytoplasmic domain (Liddington and Ginsberg, 2002). Furthermore, it has been shown that the  $\beta 1D$  tail binds talin better than the  $\beta 1A$  cytoplasmic domain does (Belkin *et al.*, 1997; Pfaff *et al.*, 1998). To test the potential importance of cytoskeleton linkage to the adhesive activity of  $\alpha 7X1$  expressing cells, we assayed binding of MCF-7 Tet-Off double transfectants to laminin-1 substrates in the presence of cytochalasin D, an inhibitor of actin polymerization. Treatment of cells with 20  $\mu M$  cytochalasin D did not affect the adhesion of the cell lines to type I collagen (Figure 19). Similarly, adhesion of HuX1/A and HuX2/A to laminin-1 was unaffected by cytochalasin D. However, the presence of the drug significantly reduced HuX1/D cell adhesion to laminin 1 by nearly 70%. This result suggests that the formation of polymerized actin during the adhesive process was necessary for the  $\beta 1D$  cytoplasmic domain to confer enhanced binding of  $\alpha 7X1\beta 1$  to laminin-1.

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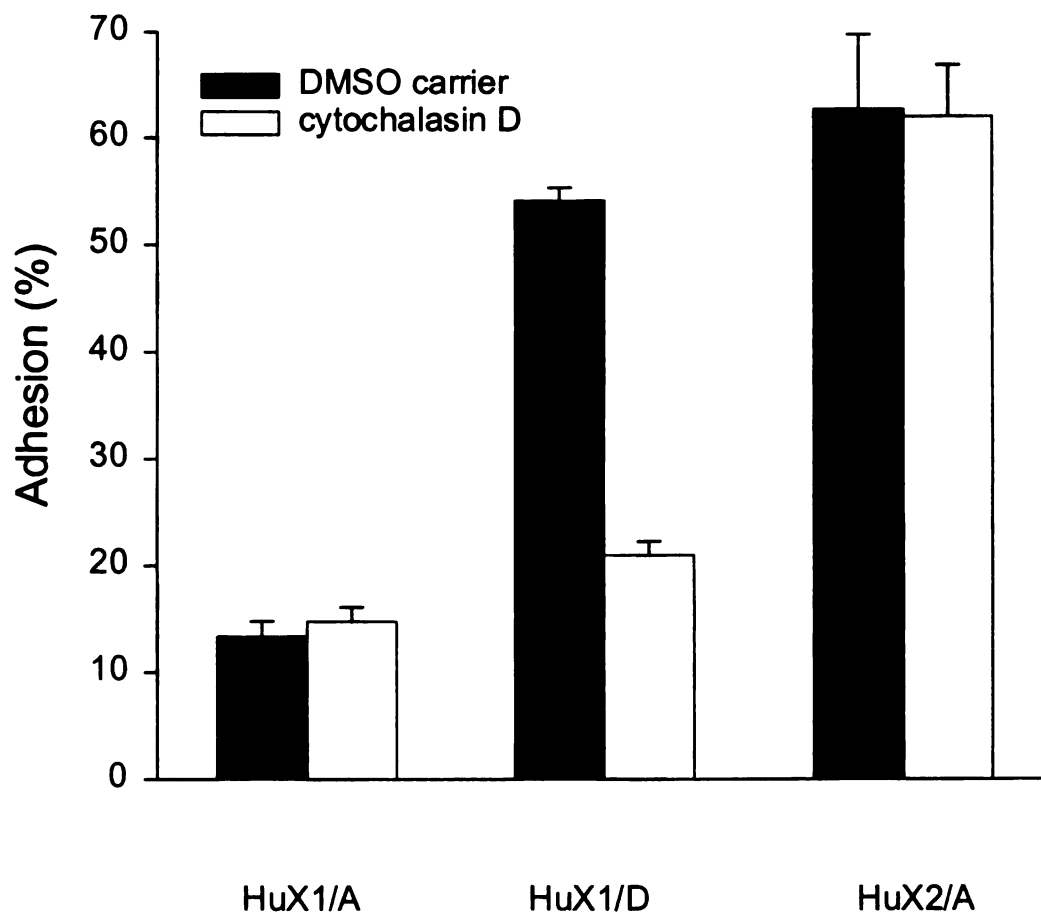


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**Figure 19.**

Cytochalasin D modulates the adhesion of HuX1/D to LN-1



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## **The potential role of $\beta$ 1D in growth arrest is not demonstrated when it is expressed in MCF-7 Tet-Off cells**

### *The use of doxycycline regulatable expression system*

The onset of  $\beta$ 1D expression during myodifferentiation coincides with the timing of myoblast withdrawal from the cell cycle. It has been shown that the expression of the exogenous  $\beta$ 1D integrin in C2C12 myoblasts and NIH 3T3 or REF 52 fibroblasts inhibits cell proliferation (Belkin and Retta, 1998) by nuclear injection of  $\beta$ 1D cDNA. This kind of experiment is technique sensitive and the conclusion is drawn from a limited number of experiment cells. To avoid the drawback of this kind of technique, we used the regulatable  $\beta$ 1D in MCF-7 cells to explore the role of  $\beta$ 1D in growth inhibition. The MCF-7 Tet-Off cells were kept in the media containing doxycycline to suppress the expression of mouse  $\beta$ 1 for 48 hours after the transfection of pRev-TRE  $\beta$ 1A or  $\beta$ 1D (see material and methods section). The expression vector carries hygromycin selection marker upstream of the tetracycline responsive element; thus the transfectants can be selected for stable transfectants by hygromycin even though the regulatable mouse  $\beta$ 1 is suppressed by doxycycline. The mouse  $\beta$ 1A or  $\beta$ 1D were never induced before the growth curve analysis was carried out.

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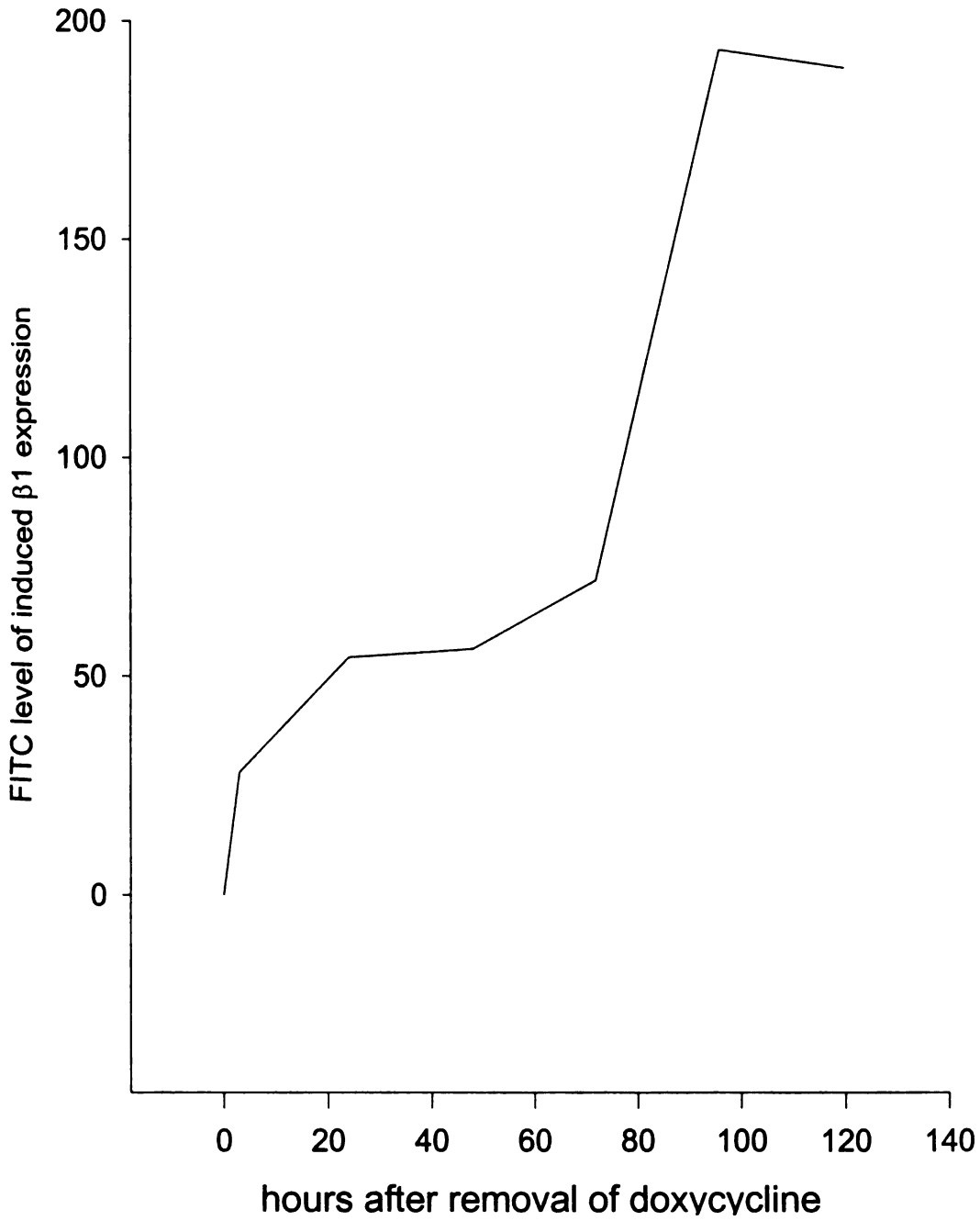
*$\beta 1A$ - and  $\beta 1D$ - expressing cells have the same growth rate*

Upon the removal of doxycycline, the level of surface expression of tet-regulatable  $\beta 1$  in MCF-7 Tet-Off cells starts to increase and reaches its maximum level in four days as detected by FACS scan (Figure 20). In the first 24 hours, there was a decent amount of mouse  $\beta 1$  expressed on cell surface. We then examined the growth curves of HuX1/A and HuX1/D cells for 8 days. Although the medium was replenished every day, we included the parental cells for growth curve analysis as a control. Each cell line was grown under doxycycline and hygromycin before experimental analysis. On day zero, the cells were counted and seeded in two series, one still cultured in doxycycline, the other in doxycycline-free media. At each time point, the cells were fixed and stained for OD. Surprisingly, we found that there were no differences in the rates of growth between doxycycline and doxycycline-free cultures in all three cell lines (Figure 21). Regardless of the expression of mouse  $\beta 1A$  or  $\beta 1D$ , the cells grew exponentially until day 6. After day 6, cells growth reached a plateau. The presence of 10 ng/ml of doxycycline did not alter the growth rate of parental cells nor did it alter the growth rate of transfectants. These results indicated that the low concentration of doxycycline has no toxic effect to MCF-7 cells. According to the doxycycline titration data, 10 ng/ml is more than enough to suppress the expression of regulatable  $\beta 1$ . Since there was no difference in growth rate in all groups, we concluded that expression of  $\beta 1D$  apparently did not alter the cell cycle progression.



**Figure 20.**

Kinetics of surface expression of  $\beta 1$  upon induction in MCF-7 Tet-Off cells



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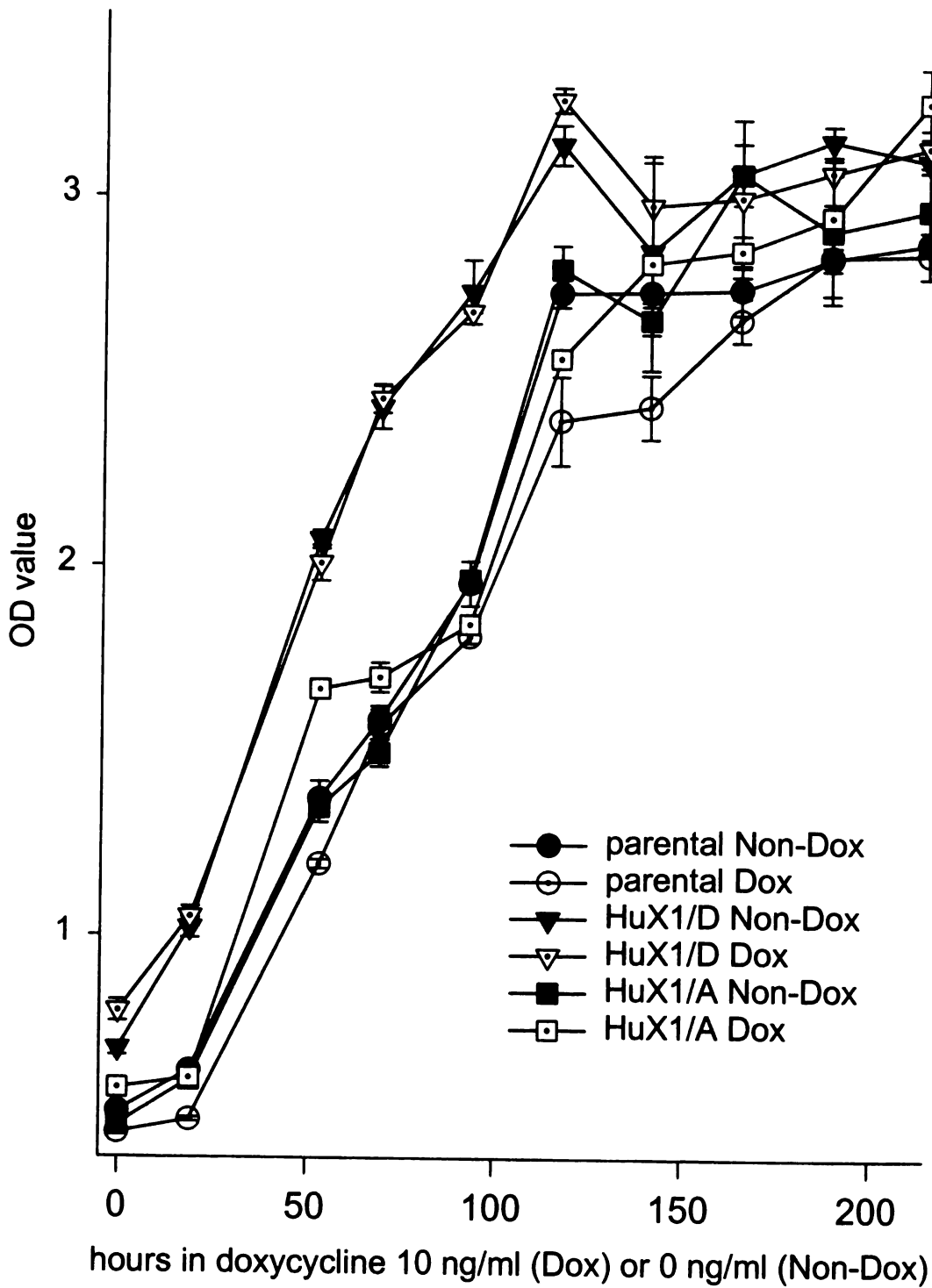
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Figure 21.

Growth curve of MCF-7 Tet-Off cells and transfectants

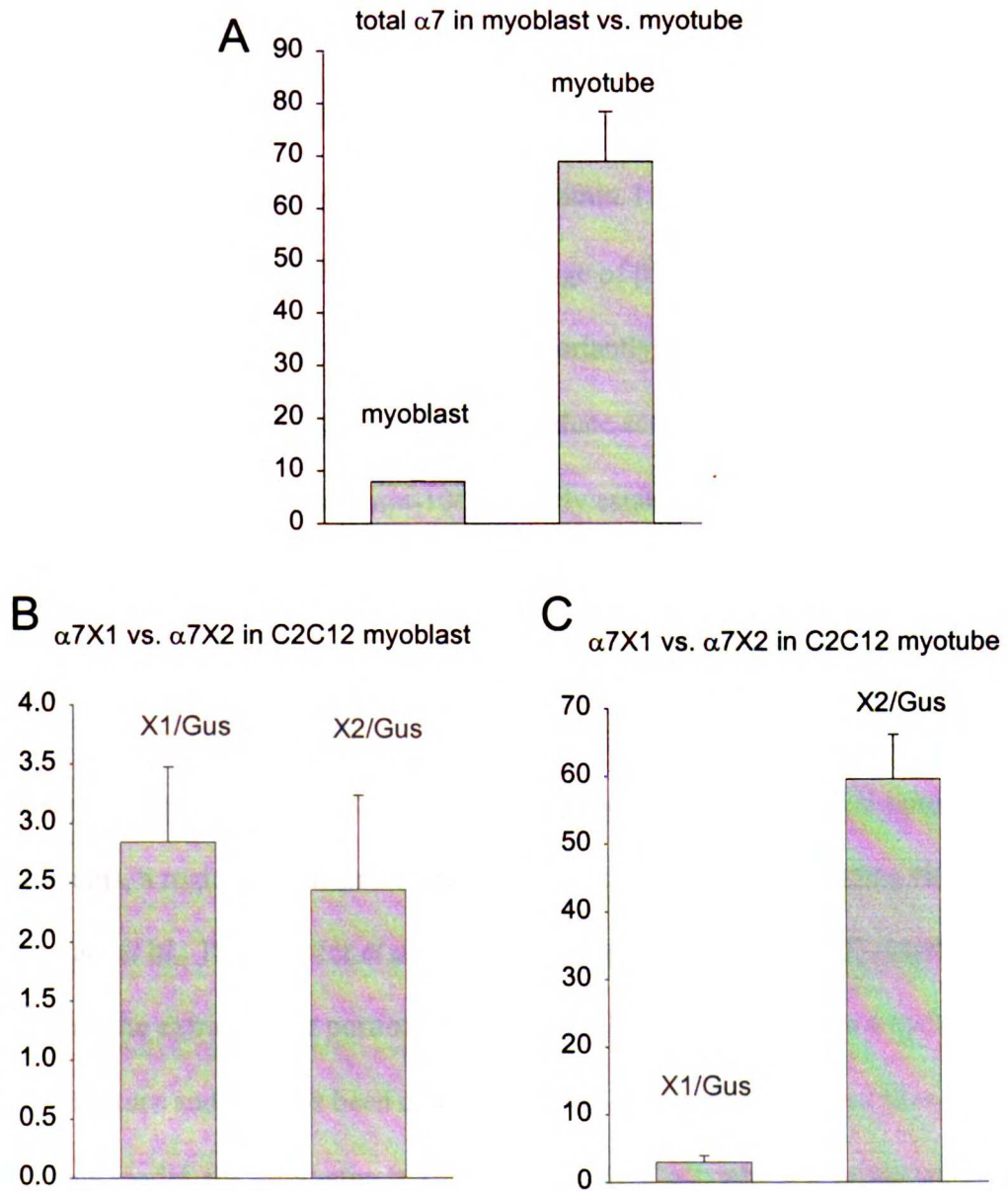




It has been shown in many reports that  $\alpha 7X1$  and  $\alpha 7X2$  are developmentally regulated. Since antibodies specific to  $\alpha 7X1$  or  $\alpha 7X2$  is not available, all the reports are based on low cycle PCR. Basically, it is a semi-quantitative analysis of mRNA. The validity of the result depends on how the experimental conditions are chosen and may vary from experiment to experiment. To get reproducible data about the expression profiles of  $\alpha 7X1$  and  $\alpha 7X2$  mRNA during myodifferentiation, we use quantitative PCR to compare the mRNA levels of total  $\alpha 7$ ,  $\alpha 7X1$ , and  $\alpha 7X2$  in C2C12 myoblasts and myotubes which were derived by in vitro myodifferentiation (Blau *et al.*, 1985). Basically, the myoblasts were cultured in 2% horse serum in DMEM for 6 days and the mRNA of differentiated myotubes was collected. The probes designed as reporter were verified for efficiency before utilization. The levels of mRNA of total  $\alpha 7$ ,  $\alpha 7X1$ , and  $\alpha 7X2$  were normalized reference to an internal house keeping gene (GUS). The results (Figure 22) were similar to what have been reported in low-cycle PCR. In myoblasts, the mRNA level of  $\alpha 7X1$  and that of  $\alpha 7X2$  were comparable. After myodifferentiation, the total  $\alpha 7$  mRNA was markedly increased in myotubes and this increase was mainly attributed to the increase in  $\alpha 7X2$ .

**Figure 22.**

Quantitative PCR of  $\alpha 7X1$  and  $\alpha 7X2$  in myoblasts and myotubes





## Conclusion and discussion

We demonstrate that when the  $\alpha 7X1$  alternatively spliced integrin subunit heterodimerizes with the muscle  $\beta 1D$  subunit, the ligand specificity is altered. When  $\alpha 7X1$  is coupled to  $\beta 1A$  it is unable to bind to laminin-1 but can efficiently mediate adhesion to laminin-10/11. Following the exchange of  $\beta 1A$  for  $\beta 1D$ , the receptor can then also promote cell binding to laminin-1. Importantly, these results show that the replacement of  $\beta 1A$  with  $\beta 1D$  does not simply induce activation of the receptor since the capacity of  $\alpha 7X1\beta 1A$  to bind laminin-10/11 clearly establishes that it is already activated. Furthermore, these data appear to represent the first example of direct modulation of ligand specificity of an integrin by the associated  $\beta 1$  cytoplasmic domain.

The alternative splicing of  $\alpha 7X1/X2$  occurs between the III and IV homology repeat domains, a region implicated in ligand specificity, affinity, or binding (Irie *et al.*, 1995; Kamata *et al.*, 1995; Tozer *et al.*, 1999). Springer (Springer, 1997) originally proposed that the extracellular portion of integrin  $\alpha$  chain is organized in a cyclic- $\beta$  propeller structure and this has been confirmed by X-ray crystallography (Xiong *et al.*, 2002). In this model, the alternatively spliced segment of  $\alpha 7$ , which is predicted to be located outside the “blade” structure joined by the two-homology repeat domains, may modulate folding of these adjacent  $\beta$  sheets, thereby controlling integrin conformation



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and function. We speculate that  $\alpha 7X1$ , when bound to the  $\beta 1D$  variant, forms an active laminin-1 ligand-binding pocket that resembles that formed by the constitutively active  $\alpha 7X2/\beta 1A$ . However, recent work (von der Mark *et al.*, 2002) indicates that a double truncated and soluble form of  $\alpha 7X1\beta 1$  complex lacking the cytoplasmic domains is unable to bind to laminin-1 in the presence of  $Mn^{+2}$ .

Exogenous  $\beta 1D$  expression in MCF-7 cells did not enhance cell adhesion to laminin-1 in the absence of  $\alpha 7$  and, in the presence of  $\alpha 7X1$  and  $\beta 1D$ , the enhancement of adhesion was mainly attributed to  $\alpha 7$  and not to the other moderately expressed  $\alpha 2$ ,  $\alpha 3$ , or  $\alpha 6$  integrins that could potentially bind to laminin-1. Since there is no preference in the coupling of these  $\alpha$  subunits with  $\beta 1A$  or  $\beta 1D$  ((Belkin *et al.*, 1997; Pfaff *et al.*, 1998); this study), our results indicate that  $\beta 1D$ -induced functional augmentation in MCF-7 cells is specific for the  $\alpha 7X1$  integrin. MCF-7 cells have  $\beta 1$  integrin receptors for other ligands, including  $\alpha 2\beta 1$  for collagen I (Jones *et al.*, 1995; Maemura *et al.*, 1995) and  $\alpha v\beta 1$  for vitronectin (Wong *et al.*, 1998). Adhesion to these two ligands was not altered by the expression of  $\beta 1D$ . In addition, the forced expression of  $\beta 1D$  in fully functional  $\alpha 7X2$  cells did not further enhance the adhesion. These results suggest that the  $\beta 1D$  partner subunit can selectively alter  $\alpha 7X1$  functionality.

It is well documented that affinity modulation can occur by inside-out signaling via transmembrane conformational transitions (Faull *et al.*, 1994; Schwartz *et al.*, 1995; Hughes and Pfaff, 1998; Longhurst and Jennings, 1998; Liddington and Ginsberg, 2002).

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Previous studies using cells transfected with  $\beta 1D$  indicated that the subunit induced a number of cellular alterations that included decreased spreading and migration, and increased binding of fibronectin and incorporation into matrix (Belkin *et al.*, 1997). These results are consistent with stabilized integrin-cytoskeleton interaction that appears to be mediated by increased binding of  $\beta 1D$  with talin and elevated levels of integrin activation. In the context of the current studies,  $\alpha 7X1$  appears to exist in a partially active conformation as indicated by the finding that the TS2/16 mAb could switch this integrin to a functional laminin-1 receptor (Ziober *et al.*, 1997). Formation of the  $\alpha 7/\beta 1D$  complex appears to induce a similar conversion from an intermediate level of activation to a fully active state. It is possible that association with  $\beta 1D$  modulates the ligand-binding affinity of  $\alpha 7$  for the laminin-1 ligand rather than altering its activation level. If this was the case, then  $\alpha 7X1/\beta 1A$  would assume an intermediate affinity state and formation of  $\alpha 7X1/\beta 1D$  generates a high affinity state. There is precedence for intermediate affinity states (Takagi *et al.*, 2002). Alternatively,  $\beta 1D$ -induced activation of  $\alpha 7X1$  may be related to post-receptor occupancy events such as receptor clustering and avidity regulation, although as Springer (Shimaoka *et al.*, 2002) has indicated the data supporting this form of integrin activity modulation remains controversial. Furthermore, the cellular environment can also modulate integrin function (Zhang *et al.*, 1996; Hughes *et al.*, 1997). Thus, in contrast to MCF-7 cells, when HT-1080 cells were transfected with  $\alpha 7$  splice variants, both X1 and X2 isoforms were fully active for binding to laminin-1 (Ziober *et al.*, 1997), indicating that factors related to cell-type specificity are important in regulating integrin function. Additional studies are needed to

further define the specific mechanism responsible for  $\beta$ 1D-induced functional modulation of  $\alpha$ 7X1.

Importantly, differences in the interaction of  $\beta$ 1 alternatively spliced cytoplasmic tails with the cytoskeleton have been reported. The muscle-specific  $\beta$ 1D splice variant has been shown to bind talin with higher affinity than  $\beta$ 1A (Belkin and Retta, 1998; Pfaff *et al.*, 1998). Conversely, the  $\beta$ 1A tail has been reported to bind filamin and  $\alpha$ -actinin more strongly than does  $\beta$ 1D (Belkin *et al.*, 1997; Pfaff *et al.*, 1998). Although the  $\beta$ 1D cytoplasmic domain-talin interaction may occur, this potential effect does not explain how  $\beta$ 1D induces binding to laminin-1 by  $\alpha$ 7X1 or why adhesion to other laminin isoforms by these receptors is unchanged. Also the  $\alpha$ 2 integrin subunit was shown to associate with  $\beta$ 1D, yet adherence to collagen I was unaltered. However, the experiments revealing that treatment of HuX1/D cells with cytochalasin D decreased binding to laminin-1, suggests a mechanism by which a conformational shift is transmitted to the extracellular domain following association of the  $\beta$ 1D tail with the cytoskeleton.

We investigated the mechanism underlying functional regulation of the alternatively spliced X1 and X2 variants by its partner  $\beta$ 1 integrin. The physiological importance of the  $\alpha$ 7 and  $\beta$ 1 splice variants is related to the varied roles of  $\alpha$ 7 $\beta$ 1 as the major laminin-binding receptor in skeletal muscle, where it has multiple functions not only during myoblast motility and differentiation but also in maintenance of mature fiber anchorage and assembly of its underlying basement membrane. When myoblasts begin to

differentiate into myotubes, the partner integrin of  $\alpha 7$  switches from the  $\beta 1A$  to the  $\beta 1D$  cytoplasmic form. Our results indicate that coupling of the  $\alpha 7X1$  variant with the  $\beta 1D$  isoform modifies the receptor's ligand binding specificity and forms a more promiscuous receptor that can mediate adhesion to laminin-1 substrates.

The mechanism by which the  $\beta 1D$  cytoplasmic domain can lead to enhanced binding of  $\alpha 7X1$  expressing cells to laminin-1 may be related to induced changes in integrin conformation. For example, it is possible that a conformational shift in the ligand-binding pocket could account for the modified ligand specificity we observed. It is interesting that  $\alpha 7X1$  when partnered to either  $\beta 1A$  or  $\beta 1D$  binds efficiently to laminin-10/11 yet exhibits activity to laminin-1 only when coupled to  $\beta 1D$ . Thus  $\alpha 7X1$  show an intrinsic binding activity to laminin 10/11 but cannot bind laminin-1 unless it is coupled to  $\beta 1D$ . In contrast,  $\alpha 7X2$  shows strong activity toward laminin-1 and laminin-2/4 but not to laminin-10/11. These results suggest a complex regulation of ligand specificity that must relate to structural components not only on the integrin receptor but also on the laminin isoforms as well.

The exchange of  $\beta 1A$  with  $\beta 1D$  may lead to enhanced ligand-binding affinity of  $\alpha 7\beta 1$  integrin. Most likely the conversion of  $\alpha 7X1$  to a permissive laminin-1-binding state by  $\beta 1D$  is a consequence of the ability of the cytoplasmic domain of  $\beta 1D$  to transmit conformational changes to the  $\alpha 7$  subunit that then regulate ligand specificity and/or affinity. Since  $\alpha 7X1$  when coupled with  $\beta 1D$  did not alter its poor binding to laminin

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2/4, this implies that the increase in activity toward laminin-1 is specific and presumably represents a subtle change in how the integrin regulates its ligand specificity. Clearly additional studies are needed to evaluate these possibilities.

The onset of  $\beta$ 1D expression during myodifferentiation coincides with the timing of myoblast withdrawal from the cell cycle. It has been shown that the expression of the exogenous  $\beta$ 1D integrin in C2C12 myoblasts and NIH 3T3 or REF 52 fibroblasts inhibited cell proliferation (Belkin and Retta, 1998) by nuclear injection of the  $\beta$ 1D cDNA. In contrast, our results showed that the parental cells and the transfectants have the same growth rate regardless of the expression of  $\beta$ 1D. Although we monitored cell surface expression level as a indicator of  $\beta$ 1D level inside the cell, it is unlikely that  $\beta$ 1D reaches the cell surface without exerting its effect on growth rate if it has any. MCF-7 is a tumor cell so it is possible that this cell can overcome the growth inhibitory effect. Usually this kind of adaptation would take some time to develop; however, our experiment was done in only a few days and these cells has never been induced to express  $\beta$ 1D before the experimental procedures. Moreover, Baudoin *et al.* knockout  $\beta$ 1A and knockin in  $\beta$ 1D exon (Baudoin *et al.*, 1998) and found the embryos were viable till 10.5 dpc. The embryonic fibroblasts can be cultured as cell line. Unless there was some compensation mechanism, the animal should not be viable at 10 dpc if  $\beta$ 1D inhibits growth.

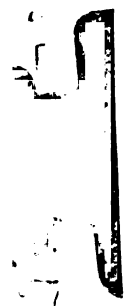
$\alpha$ 7 and  $\beta$ 1D are expressed during myoblast differentiation and are located at the



sarcolemma, MTJs, and NMJs in adult skeletal muscle (Bao *et al.*, 1993; Ziober *et al.*, 1993; Belkin *et al.*, 1996; Martin *et al.*, 1996). In mouse skeletal muscle,  $\alpha 7$  is expressed as early as E10.5 days of development, whereas the  $\beta 1D$  variant appears in the late stage embryo. Furthermore, there is a concurrent expression of both  $\alpha 7$  and  $\beta 1D$  during the first formation of sarcomeres and membrane infoldings at the MTJ (van der Flier *et al.*, 1997). In addition, when  $\beta 1D$  is detected in the mouse embryos at E17.5, it is enriched in MTJ along with the  $\alpha 7$  subunit. Conversely, laminin-1 is expressed during embryonic and fetal stages. Interestingly, laminin-1 is concentrated at the MTJ of human skeletal muscle and persists until birth (Gullberg *et al.*, 1999). Because the expression of the X1/X2 isoforms is developmentally regulated, we hypothesize that the functionality of  $\alpha 7$  is controlled in a tissue-specific manner by the coexpression of  $\beta 1D$ . Both X1 and X2 isoforms are expressed in myoblasts, but X2 is more abundant in mature muscle, implying that this form of the receptor is present at permanent anchorage sites. It has been reported that in the hind limb following early muscle differentiation,  $\alpha 7X1$  shows a relative increase in expression but that postnatally the X2/X1 ratio also increases (Hodges *et al.*, 1997). We propose that the X1 isoform is important during dynamic adhesion related to muscle development and repair (motility, fusion, and matrix assembly), whereas the X2 variant performs more stable adhesion functions (mature junctional assemblies) (Ziober *et al.*, 1997). This is consistent with the dynamics of the expression of  $\alpha 7$  and  $\beta 1$  alternatively spliced variants during development (Ziober *et al.*, 1993) and muscle regeneration (Kaariainen *et al.*, 2001; Kaariainen *et al.*, 2002). Laminin isoform expression also changes during muscle development and repair (reviewed in (Colognato and Yurchenco, 2000)) and so the expression of integrin receptor and potential ligand is

coordinately coupled.

In summary, we have demonstrated that the cytoplasmic domain of the  $\beta 1$  integrin can modulate the functionality and ligand specificity of the  $\alpha 7\beta 1$  receptor. We show that when coupled to the  $\beta 1D$  partner,  $\alpha 7X1$  splice variant shows a broader spectrum of specificity and can efficiently bind laminin-1. The formation  $\alpha 7X1/\beta 1D$  in vivo seems likely given that both subunits are present during the late embryonic stage of skeletal muscle development. This suggests that the simultaneous expression of  $\alpha 7X1$  along with  $\beta 1D$  could facilitate the organization and stability of the MTJ during the early stages of muscle fiber maturation.



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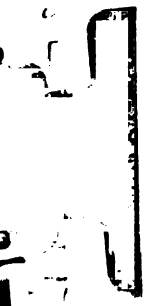
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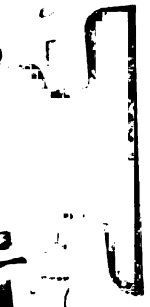
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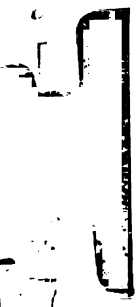
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**Appendix I.** Surface expression level of integrins in MCF-7 and transfectants

Cell line	Integrin pair*	Mean fluorescence intensity					
		$\alpha 2^a$	$\alpha 3^b$	$\alpha 6^c$	$\alpha 7^d$	Endogenous $\beta 1A^e$	Mouse $\beta 1A/D^f$
MCF-7	--/ $\beta 1A$	30	39	3	-	78	-
X1/A	$\alpha 7X1/\beta 1A$	25	27	6	43	86	-
X1/D	$\alpha 7X1/\beta 1D$	41	23	13	32	55	20
X2/A	$\alpha 7X2/\beta 1A$	56	64	7	40	58	-
X1/D-10	$\alpha 7X1/\beta 1D$	29	27	10	34	30	41
X1/D-12	$\alpha 7X1/\beta 1D$	18	8	9	25	14	15
X1/D-13	$\alpha 7X1/\beta 1D$	28	17	7	78	55	57
X1/D-17	$\alpha 7X1/\beta 1D$	21	17	12	28	31	15
X1/D-21	$\alpha 7X1/\beta 1D$	19	18	20	32	21	27
X1/D-27	$\alpha 7X1/\beta 1D$	28	18	5	49	47	31
X1/D-34	$\alpha 7X1/\beta 1D$	23	15	6	43	19	3
$\beta 1A$	--/ $\beta 1A$	52	31	2	--	27	49
$\beta 1D$	--/ $\beta 1D$	14	11	2	--	10	32

\*The parental and transfectants were processed for FACS using the indicated antibody as detailed in Material and Methods Section. The level of expression is expressed as the mean of 10,000 cells scanned excluding non-vital cells.

<sup>a</sup>mAb VM1, mouse anti- $\alpha 2$

<sup>b</sup>mAb VM2, mouse anti- $\alpha 3$

<sup>c</sup>mAb GoH3, rat anti- $\alpha 6$

<sup>d</sup>CY8, rat anti-mouse  $\alpha 7$ .

<sup>e</sup>mAb AIIB2, rat anti-human  $\beta 1$ .

<sup>f</sup>mAb MB1.2, rat anti-mouse  $\beta 1$ .





**Appendix II.** Surface expression level of integrins in MCF-7 Tet-Off cells and transfectants

Cell line	Integrin pair*	Mean fluorescence intensity					
		$\alpha 2^a$	$\alpha 3^b$	$\alpha 6^c$	$\alpha 7^d$	Endogenous $\beta 1A^e$	Mouse $\beta 1A/D^f$
Tet-Off	--/ $\beta 1A$	16	8	0	-	17	-
HuX1/A	$\alpha 7X1/\beta 1A$	4	3	0	294	24	291
HuX1/D	$\alpha 7X1/\beta 1D$	6	7	0	315	39	317
HuX2/A	$\alpha 7X2/\beta 1A$	10	5	0	374	25	299
HuX2/D	$\alpha 7X1/\beta 1D$	15	11	1	312	44	277

\*The parental and transfectants were processed for FACS using the indicated antibody as detailed in Material and Methods Section. The level of expression is expressed as the mean of 10,000 cells scanned excluding non-vital cells.

<sup>a</sup>mAb VM1, mouse anti- $\alpha 2$

<sup>b</sup>mAb VM2, mouse anti- $\alpha 3$

<sup>c</sup>mAb GoH3, rat anti- $\alpha 6$

<sup>d</sup>mAb 9.1, mouse anti-human  $\alpha 7$ .

<sup>e</sup>mAb AIIB2, rat anti-human  $\beta 1$ .

<sup>f</sup>mAb MB1.2, rat anti-mouse  $\beta 1$ .



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