

Heparan Sulfate Heterogeneity in Skeletal Muscle Basal Lamina: Demonstration by Phage Display-Derived Antibodies

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The basal lamina (BL) enveloping skeletal muscle fibers contains different glycoproteins, including proteoglycans. To obtain more information on the glycosaminoglycan moiety of proteoglycans, we have selected a panel of anti-heparan sulfate (HS) antibodies from a semisynthetic antibody phage display library by panning against glycosaminoglycan preparations derived from skeletal muscle. Epitope recognition by the antibodies is strongly dependent on *O*- and *N*-sulfation of the heparan sulfate. Immunostaining with these antibodies showed a distinct distribution of heparan sulfate epitopes in muscle basal lamina of various species. Clear differences in staining intensity were observed between neural, synaptic, and extrasynaptic basal

laminae. Moreover, temporal and regional changes in abundance of heparan sulfate epitopes were observed during muscle development both *in vitro* and *in vivo*. Taken together, these data suggest a role for specific heparan sulfate domains/species in myogenesis and synaptogenesis. Detailed analysis of the functions of heparan sulfate epitopes in muscle morphogenesis has now become feasible with the isolation of antibodies specific for distinct heparan sulfate epitopes.

Key words: heparan sulfate proteoglycan; glycosaminoglycan; basal lamina; neuromuscular junction; myogenesis; synaptogenesis

The basal lamina (BL) enveloping skeletal muscle fibers plays various roles in muscle development and regeneration (Sanes, 1986; Wright et al., 1991). Several BL molecules have been identified as specifically synaptic, extrasynaptic, or common (Sanes, 1982; Hall and Sanes, 1993). On the protein level, these include isoforms of laminin, collagen, and entactin (Sanes et al., 1990; Chiu and Ko, 1994; Patton et al., 1997). Lectin staining (Sanes and Cheney, 1982; Iglesias et al., 1992) and a recent report on synapse-specific carbohydrates (Martin et al., 1999) indicate a spatial heterogeneity for carbohydrates as well.

Heparan sulfate proteoglycans (HSPGs), consisting of a core protein and a carbohydrate moiety [heparan sulfate (HS)], are main components of muscle BLs. So far, three BL HSPGs have been identified: perlecan, agrin, and type XVIII collagen (Noonan et al., 1991; Tsen et al., 1995; Halfter et al., 1998). HSPGs are implicated in developmental processes underlying myogenesis and synaptogenesis (Anderson and Fambrough, 1983; Anderson et al., 1984; Bayne et al., 1984; Dmytrenko et al., 1990). Perlecan may effect neuromuscular junction (NMJ) formation by binding growth factors (Peng et al., 1998). Agrin, a major HSPG of the synaptic BL (sBL), orchestrates acetylcholine receptor (AChR) clustering (Campanelli et al., 1994; Ruegg and Bixby, 1998). Heparin and HS are involved in AChR clustering induced by nerve (Hirano and Kidokoro, 1989) and agrin (Wallace, 1990). In muscle cell lines defective in glycosaminoglycan

(GAG) synthesis, a causal relationship between GAGs and AChR clustering is demonstrated (Ferns et al., 1993; Gordon et al., 1993; Mook-Jung and Gordon, 1996). HS binding may also mediate the synapse-specific anchoring of BL-resident proteins such as acetylcholine esterase and possibly certain laminin isoforms (Brandan et al., 1985; San Antonio et al., 1993; Patton et al., 1997).

Another major characteristic of HS is the binding of growth factors such as neuregulin (Fischbach and Rosen, 1997), midkine (Zhou et al., 1997), heparin-binding growth-associated molecule (Peng et al., 1995; Szabat and Rauvala, 1996), heparin-binding epidermal growth factor-like growth factor (Chen et al., 1995), and basic fibroblast growth factor. The latter protein is involved in postsynaptic differentiation (Peng et al., 1991) and maintenance of the proliferative state of satellite cells (Rapraeger et al., 1991; Olwin and Rapraeger, 1992; Olwin et al., 1994; Crisona et al., 1998). Considering the diversity of proteins that bind HS, HS molecules may contain unique domains (epitopes) that are specific for these interactions.

Studies of HSPGs have mainly been focused on the protein core. The structure and the role of the HS moiety are difficult to investigate because of a lack of appropriate tools. Only a few antibodies that recognize HS epitopes have been described (David et al., 1992; van den Born et al., 1992). Recently, we adapted phage display technology to obtain epitope-specific antibodies against HS (van Kuppevelt et al., 1998). Here we report on the isolation, characterization, and application of antibodies selected against HS-containing GAG preparations from skeletal muscle. We provide evidence for the existence of several specific, differentially distributed HS epitopes in (synaptic and extrasynaptic) muscle and nerve BLs. Moreover, we found a shift in abundance of these epitopes in BLs of developing muscle both *in vitro* and *in vivo*. These data suggest an involvement of specific HS epitopes in myogenesis and synaptogenesis.

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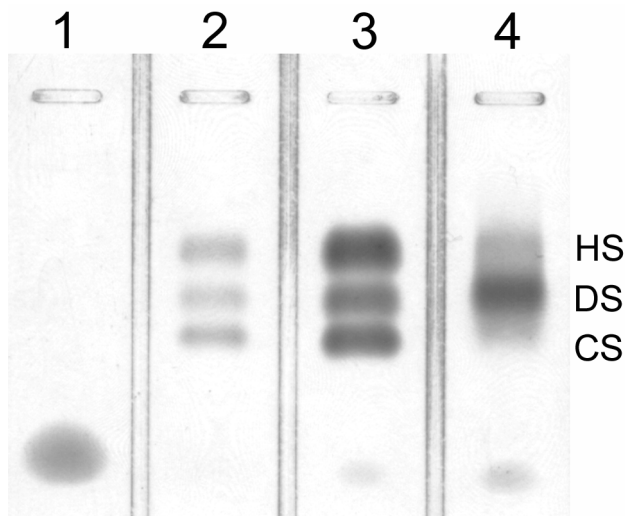


Figure 1. Silver-stained 1% agarose gel of a muscle glycosaminoglycan preparation. *Lane 1*, Sample buffer (control); *lane 2*, 5 ng GAG standard; *lane 3*, 20 ng GAG standard; *lane 4*, typical glycosaminoglycan preparation of mouse skeletal muscle. Dermatan sulfate (DS) is present at approximately fourfold higher concentration compared with chondroitin sulfate (CS) and heparan sulfate (HS).

MATERIALS AND METHODS

Materials. Synthetic single-chain variable fragment (scFv) library #1 (Nissim et al., 1994) was generously provided by Dr. G. Winter (Cambridge University, Cambridge, United Kingdom). Human skeletal muscle samples were generously provided by Prof. Dr. D. Ruiter (Department of Pathology, University of Nijmegen, Nijmegen, The Netherlands). *Torpedo marmorata* electric organ was generously provided by Dr. M. H. De Baets (Department of Immunology, University of Maastricht, Maastricht, The Netherlands). Mice (C3H, male, 70 d), rats (Wistar, male, six weeks), and rat embryos (Wistar, 10, 13, 16, and 19 d after conception) were obtained from the University of Nijmegen Central Animal Laboratory. C3H mouse-derived skeletal muscle (C₂C₁₂) cell line was purchased from American Type Culture Collection (Rockville, MD). Glycosaminoglycan-deficient myoblast (S₂₇) cell line was a generous gift of Dr. Z. Hall (Department of Physiology, University of California, San Francisco, CA); mutant Chinese hamster ovary (CHO) cell lines were kindly provided by Dr. J. Esko (Department of Biochemistry, University of Alabama, Birmingham, AL).

For phage display, two *Escherichia coli* strains were used: suppressor strain TG1 [K12, *supE*, *hsdΔ5*, *thi Δ(lac-proAB)*, F'(traD36, *proAB*⁺, *lacI*^q, *lac ZΔM15*)] and nonsuppressor strain HB2151 [K12, *ara*, *thi Δ(lac-pro)*, F'(proAB⁺, *lacI*^q, ZΔM15)]. Helper phage VCS-M13 was from Stratagene (La Jolla, CA).

All chemicals used were purchased from Merck (Darmstadt, Germany), unless stated otherwise. Bacterial media (2xTY and LB) and cell culture media were from Life Technologies (Paisley, Scotland). Chondroitinase ABC (from *Proteus vulgaris*, EC 4.2.2.4), chemically modified heparin kit, anti-chondroitin sulfate (CS) "stub" antibody (2B6), and anti-heparan sulfate stub antibody (3G10) were from Seikagaku Kogyo Co. (Tokyo, Japan). Heparinase III (from *Flavobacterium heparinum*, EC 4.2.2.8), heparan sulfate from bovine kidney and porcine intestinal mucosa, heparin from porcine intestinal mucosa, chondroitin 4-sulfate and chondroitin 6-sulfate from bovine trachea, dermatan sulfate (DS) from porcine intestinal mucosa, keratan sulfate from bovine cornea, hyaluronic acid from human umbilical cord, DNA from calf thymus, phenylmethylsulfonyl fluoride (PMSF), *N*-ethylmaleimide, aprotinin, sodium azide, pepstatin A, and bovine serum albumin (fraction V) were from Sigma (St. Louis, MO). Microtiter 96-well microtiter plates and immunotubes were from Greiner (Frickenhausen, Germany). Anti-c-Myc tag mouse monoclonal IgG (clone 9E10) was from Boehringer Mannheim (Mannheim, Germany), Anti-c-Myc tag rabbit polyclonal IgG (A-14) was from Santa Cruz Biotechnology (Santa Cruz, CA). Alkaline phosphatase-conjugated rabbit anti-mouse IgG was from Dakopatts (Glostrup, Denmark). Alexa 488-conjugated goat anti-rabbit IgG and tetramethylrhodamine isothiocyanate (TRITC)-conjugated α -bungarotoxin were from

Table 1. CDR3 sequences and germline V_H gene segments of anti-HS scFv antibodies

Clone	CDR3 sequence	V _H family	DP segment
AO4B05	LKQQGIS	3	53
AO4B08	SLRMNGWRAHQ	3	47
AO4F12	AMTQKKPRKLSL	1	7
RB4CB9	HAPLRNTRTNT	3	38
RB4CD12	GMRPRL	3	38
RB4EA12	RRYALDY	3	32
RB4EG12	SGRKYFRARDMN	1	4

Antibodies with the prefix AO were obtained after panning against GAGs from mouse skeletal muscle, whereas antibodies with the prefix RB were selected by panning against GAGs isolated from human skeletal muscle. CDR3 sequences are shown in single-letter amino acid code. V_H families and DP segments were deduced from the Sanger Centre Germline Query (http://www.sanger.ac.uk/DataSearch/gq_search.shtml) by applying the full-length V_H sequences of the anti-HS scFv antibody clones (nomenclature according to Tomlinson et al., 1992).

Molecular Probes (Eugene, OR). Mowiol (4–88) was from Calbiochem (La Jolla, CA). PCR chemicals and *Taq* polymerase (DNA polymerase from *Thermus aquaticus*) were from Promega (Madison, WI), PCR primers were from Biologio (Malden, The Netherlands), and restriction enzyme *Bst*NI was from New England Biolabs (Beverly, MA). ABI Prism Big Dye Terminator Cycle Sequencing Ready Reaction Kit was from PE Applied Biosystems (Norwalk, CT).

All experiments were performed at ambient temperature (21°C), unless stated otherwise.

Isolation of glycosaminoglycans from skeletal muscle. Mouse and human skeletal muscle specimens were homogenized, defatted in 20 vol of acetone at –20°C for 16 hr, and dried in a desiccator. Per gram of muscle tissue, 4 ml 50 mM sodium phosphate buffer, pH 6.5, containing 2 mM EDTA, 2 mM cysteine, and 10 U papain were added. Papain digestion was performed for 16 hr at 65°C, and the remaining debris was pelleted. Residual protein fragments were removed from the glycosaminoglycans by mild alkaline borohydride digestion in 0.5 M NaOH/0.05 M NaBH₄ at 4°C. After overnight digestion, the mixture was neutralized by addition of 6 M HCl. Residual protein fragments were precipitated by addition of 100% (w/v) trichloroacetic acid to a final concentration of 6% and precipitation at 0°C for 1 hr. Precipitated proteins were removed by centrifugation (10,000 × g for 20 min at 4°C), and glycosaminoglycans were isolated by addition of 5 vol of 100% ethanol to the supernatant and overnight precipitation at –20°C. After centrifugation (10,000 × g for 30 min at 4°C), the pelleted glycosaminoglycans were washed with 70% ethanol, dried, and dissolved in 10 mM Tris-HCl, pH 6.8. This crude glycosaminoglycan preparation was further deprived of protein contamination by DEAE Sepharose column chromatography, eluting glycosaminoglycans at 0.5 M and 1.0 M NaCl in 10 mM Tris-HCl, pH 6.8. GAG-containing eluates were pooled, and after ethanol precipitation the residual salt was removed by a 70% (v/v) ethanol wash. The resulting glycosaminoglycan preparations were dissolved in MilliQ water and stored at 4°C.

Phage display. Phage display was essentially performed as described (Van Kuppevelt et al., 1998). Synthetic scFv library #1 was subjected to four rounds of panning against mouse or human skeletal muscle glycosaminoglycan preparations. The library contains approximately 10⁸ different scFv antibody clones, composed of 50 different heavy (V_H) chain V segments with synthetic (randomly synthesized) complementarity-determining region 3 (CDR3) fragments and one light (V_L) segment. This library was *in vitro*-synthesized from V gene segments, derived from human lymphocytes, using PCR (Tomlinson et al., 1992; Nissim et al., 1994). After the last round of selection, single colonies were picked, and the antibodies expressed by these clones were evaluated for reactivity by ELISA. Clones displaying reactive antibodies were further analyzed by colony-PCR amplification of the antibody coding region and restriction digestion of the full-length PCR products with *Bst*NI (CC*^Δ₁/GG). Unique clones were grown at a larger scale, and individual plasmid DNAs were sequenced using PeIB-seq (5'-CCGCTGGATTGTTACTC-3') as a primer (located within the PeIB leader sequence).

Large scale preparation of antibodies. To produce large quantities of scFv antibodies, plasmid DNA from selected clones was used to transform nonsuppressor *E. coli* strain HB2151. Five hundred milliliters of

Table 2. Evaluation of anti-HS antibody specificity by ELISA

GAG preparation	AO4B05	AO4B08	AO4F12	RB4CB9	RB4CD12	RB4EA12	RB4EG12
K5 capsular polysaccharide (<i>E. coli</i>) ^a	–	–	–	–	–	–	–
HS (bovine kidney)	++	+++	+	+/-	++	+/-	+/-
HS (porcine intestinal mucosa)	++	++	+/-	+	+	+	+/-
HS (human lung, 0.5 M NaCl fraction) ^b	+	+/-	–	–	–	+	+/-
HS (human lung, 1.0 M NaCl fraction) ^b	++	+++	++	+	++	++	++
Heparin (porcine intestinal mucosa)	++	+++	+++	+	+++	++	+++
Heparin, desulfated and <i>N</i> -acetylated ^c	–	–	–	–	–	–	–
Heparin, desulfated and <i>N</i> -sulfated ^c	–	–	–	–	–	–	–
Heparin, <i>N</i> -desulfated and <i>N</i> -acetylated ^c	–	–	+/-	–	–	–	–

Antibody-containing supernatants of IPTG-induced *E. coli* HB2151 cultures were applied to various GAG preparations immobilized on microtiter plates. Bound antibodies were detected using anti-*c-Myc* mouse monoclonal antibody 9E10, followed by alkaline phosphatase-conjugated rabbit anti-mouse IgG, after which enzymatic activity was measured using *p*-nitrophenyl phosphate as a substrate. Substrate affinity: +++, very strong; ++, strong; +, moderate; +/-, weak; –, absent ($n = 5$).

^aSimilar to the HS precursor polysaccharide.

^bHS fraction eluting from anion exchange column at the NaCl concentration indicated.

^cInhibition-ELISA.

prewarmed 2xTY medium containing 0.1% (w/v) glucose and 100 μ g/ml ampicillin were inoculated with an overnight culture of transformed HB2151 and grown with vigorous shaking at 37°C until an OD₆₀₀ of 0.3 was reached. Induction was effectuated by addition of isopropyl- β -D-thiogalactopyranoside (IPTG) to a final concentration of 1 mM. After 3 hr incubation at 30°C the culture was cooled on ice for 20 min, and cells were pelleted (3000 \times *g* for 10 min at 4°C). One-tenth volume of 10 \times protease inhibitor mix [0.1 M EDTA, 250 mM iodoacetamide, 1 M *n*-ethylmaleimide, 1% (w/v) NaN₃, 1.5 mTIU/ml aprotinin, 0.1% (w/v) pepstatin A, 1 mM PMSF] was added to the supernatant, which was subsequently divided into aliquots and stored at 4°C. The cells were resuspended by vigorous vortexing in 5 ml ice-cold 200 mM sodium borate buffer, pH 8.0, containing 160 mM NaCl, 1 mM EDTA, and protease inhibitors. After centrifugation (5000 \times *g* for 30 min at 4°C), the supernatant (the periplasmic fraction containing the scFv antibodies) was filtered through a 0.45 μ m filter, dialyzed overnight at 4°C against PBS, divided into aliquots, and stored at –20°C.

Evaluation of antibody specificity by ELISA. Unless stated otherwise, supernatants of IPTG-induced HB2151 cultures were used for ELISA. Affinity of the antibodies to various molecules was evaluated by ELISA in two ways: scFv antibodies were applied to wells of Microton microtiter plates, coated with the molecule concerned (10 μ g/ml coating solution), and allowed to bind for 90 min. Alternatively, scFv antibodies were preincubated overnight with the test molecule (10 μ g/ml) in PBS/0.1% (w/v) Marvel, followed by transfer to and 90 min incubation in wells previously coated with heparin. Test molecules included glycosaminoglycan preparations from mouse and human skeletal muscle, HS preparations from bovine kidney and human lung, prepared as described above, commercially available heparan sulfate from bovine kidney and from porcine intestinal mucosa, heparin, chemically and enzymatically modified heparin, chondroitin 4-sulfate, chondroitin 6-sulfate, dermatan sulfate, keratan sulfate, hyaluronic acid, DNA, Marvel, and bovine serum albumin (fraction V). Bound scFv antibodies were detected using anti-*c-Myc* mouse monoclonal antibody 9E10, followed by alkaline phosphatase-conjugated rabbit anti-mouse IgG (60 min each). Plates were washed three times with PBS containing 0.1% (v/v) Tween-20 (PBST) after each incubation. Enzyme activity was detected using 1 mg/ml *p*-nitrophenyl phosphate in 1 M diethanolamine/0.5 mM MgCl₂, pH 9.8, and absorbance was read at 405 nm.

Immunohistochemistry. Periplasmic fractions of IPTG-induced HB2151 cultures were used for immunohistochemistry, unless stated otherwise. Location of the epitopes recognized by the antibodies in several tissues was assessed both on cryosections of tissue specimens and on monolayers of cultured cell lines. Tissues included human skeletal muscle and diaphragm, rat and C3H mouse skeletal muscle, rat denervated skeletal muscle, rat embryos, and *T. marmorata* electric organ. Tissue specimens were snap-frozen in liquid nitrogen-cooled isopentane and stored at –80°C. Wild-type and glycosylation-deficient CHO cell lines were studied at confluency, whereas myoblast (C₂C₁₂ and S₂₇) cell lines were analyzed at various stages of differentiation. Cells were grown as previously described (Portiér et al., 1999), differentiated in Ultrosar brain extract medium, washed three times with PBS, dried overnight, and

stored at –80°C. Cryosections (3 or 5 μ m) or tissue cultures were rehydrated for 10 min in PBS, blocked with PBS containing 0.1% (w/v) BSA for 20 min, and incubated with scFv antibodies for 90 min. Bound antibodies were detected using anti-*c-Myc* rabbit polyclonal antibody A-14 and Alexa 488-conjugated goat anti-rabbit IgG (60 min each). For visualization of AChR clusters, TRITC-conjugated α -bungarotoxin was included in the final incubation. Cryosections or tissue cultures were washed three times with PBS after each incubation. Finally, the cryosections or tissue cultures were fixed in 100% methanol, dried, and embedded in Mowiol [10% (w/v) in 0.1 M Tris-HCl, pH 8.5/25% (v/v) glycerol/2.5% (w/v) NaN₃]. As a control, primary, secondary, or conjugated antibodies were omitted.

Evaluation of antibody specificity by immunohistochemistry. To assess the heparan sulfate specificity of the scFv antibodies, cryosections or tissue cultures were preincubated with heparinase III to digest heparan sulfate [0.02 U/ml in 50 mM NaAc/50 mM Ca(Ac)₂, pH 7.0] overnight at 37°C, or with chondroitinase ABC, which digests chondroitin and dermatan sulfate (1 U/ml in 25 mM Tris-HCl, pH 8.0) for 30 min at 37°C. As a control, cryosections or tissue cultures were incubated in the reaction buffer without enzyme. After washing three times with PBS and blocking with PBS/0.1% (w/v) BSA, cryosections and tissue cultures were incubated with antibodies and processed for immunofluorescence as described above. The efficiency of chondroitinase ABC treatment was evaluated by incubation of cryosections with an antibody (2B6) against chondroitin sulfate “stubs,” generated by chondroitinase. Heparan sulfate stubs were visualized using anti- Δ -heparan sulfate antibody (3G10).

Denervation of rat skeletal muscle. The musculus gastrocnemius and the musculus soleus of the left legs of young adult rats were denervated by cutting the efferent motor nerves innervating these muscles. The ends of these nerves were fastened to the musculus biceps femoris to prevent reinnervation (Degens et al., 1992). After 11 d, rats were killed, and the calves of both the left (denervated) and right (control) legs were isolated and processed as described in immunohistochemistry.

RESULTS

Selection of antibodies against skeletal muscle GAGs

To select scFv antibodies against skeletal muscle GAG epitopes, GAGs were isolated from human and C3H mouse skeletal muscle. Typically, 10 μ g GAG could be purified from 1 gm muscle tissue (wet weight). All GAG preparations contained approximately equal amounts of CS and HS and were approximately fourfold richer in DS (Fig. 1).

Four rounds of panning were performed against mouse skeletal muscle-derived GAG preparations, resulting in antibodies that bear the prefix AO. Antibodies with the prefix RB were obtained after panning against human skeletal muscle-derived GAGs. This approach yielded a set of unique anti-HS antibodies, based on the

amino acid sequence of their heavy chain CDR3, a major determinant in antigen specificity (Table 1).

Characterization of antibodies

All antibodies showed a high reactivity in ELISA for the GAG preparation against which they were selected, whereas the reactivity for various GAG species derived from other tissues varied significantly. Despite the fact that the antibodies were selected against a GAG mixture that consisted predominantly of DS, antibodies showed affinity only for HS and heparin. No reactivity was observed with chondroitin 4-sulfate, chondroitin 6-sulfate, dermatan sulfate, keratan sulfate, hyaluronic acid, DNA, Marvel (blocking agent), and Microlon (data not shown). Antibodies reacted to various extents with a highly sulfated HS fraction (eluting at 1.0 M NaCl in ion exchange chromatography) and with a low-sulfated fraction (eluting at 0.5 M NaCl) of human lung (Table 2). All antibodies showed a major cross-reactivity with heparin, which is highly sulfated. Antibodies AO4B05, AO4B08, and (to a somewhat lesser extent) RB4CD12 showed a high reactivity for HS from bovine kidney and porcine intestinal mucosa, whereas all other antibodies interacted only moderately or weakly. K5 capsular polysaccharide from *E. coli*, which is similar to the HS precursor, was not bound by any of the antibodies.

To investigate which chemical groups are recognized by the different antibodies, we determined the reactivity of the antibodies toward modified heparin preparations (Table 2). Completely desulfated and *N*-acetylated heparin as well as completely desulfated and *N*-sulfated heparin were not recognized by any of the antibodies. Heparin that was *N*-desulfated and *N*-acetylated also was not recognized by the antibodies, except for AO4F12, which showed a weak binding.

To ascertain the HS specificity of the antibodies, immunofluorescence studies were performed on cryosections of skeletal muscle tissue that were treated with heparinase III before incubation. Heparinase treatment of cryosections resulted in a total loss of staining for all antibodies (Fig. 2), whereas treatment with chondroitinase ABC did not (data not shown). Staining of heparinase-treated cryosections with anti-HS stub antibody 3G10 (which reveals all HS that is present) showed HS to be equally distributed in synaptic and extrasynaptic BL (Fig. 2c).

Cell lines that are defective in GAG synthesis are not surface-stained by anti-HS antibodies

To further establish the anti-HS nature of the scFv antibodies, we investigated cell lines that are defective in GAG synthesis. Developmental stages from half-confluent to 8 d of differentiation of the S₂₇ cell line (Gordon and Hall, 1989) and confluent cultures of different CHO cell lines [wild type, *N*-acetylglucosaminyl- and glucuronosyltransferase-deficient; *pgsD*-677 (Lidholt et al., 1992), heparan sulfate uronic acid 2-*O*-sulfotransferase deficient; *pgsF*-17 (Dr. J. Esko, personal communication), and xylosyl transferase deficient; *pgsA*-745 (Esko et al., 1985)] were analyzed by immunofluorescence.

In contrast to wild-type myoblast cell line C₂C₁₂ (see below), the surface of S₂₇ myoblasts was not immunoreactive for any of the antibodies, nor were places of cell–cell contact. On alignment and fusion, and at day 8 of differentiation, myotubes were not stained either, indicating that the BL of this mutant cell line does not contain any of the HS epitopes recognized by any of the antibodies (Table 3, Fig. 3). A noteworthy observation was the distinct staining of perinuclear and cytosolic granules by some antibodies (Table 3).

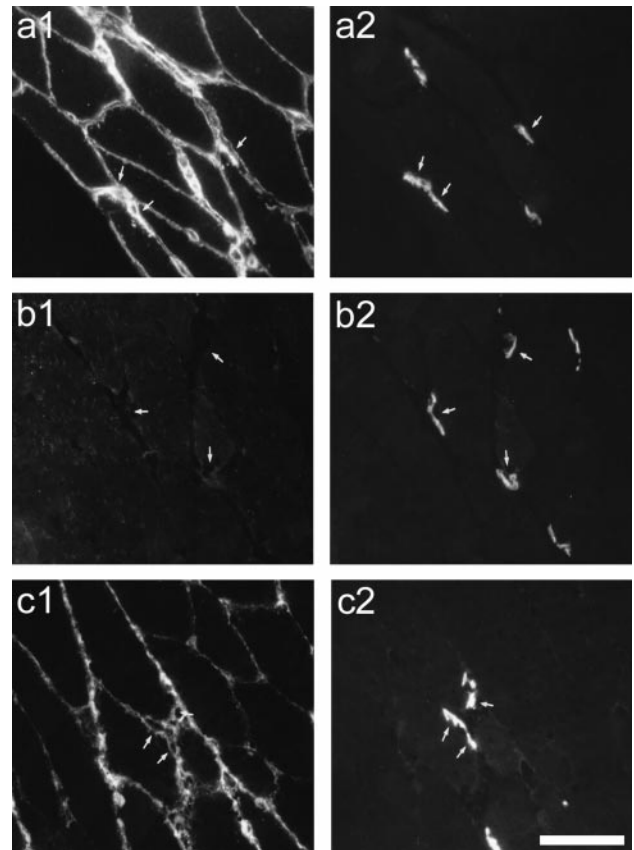


Figure 2. Staining of heparinase III-treated skeletal muscle cryosections with anti-HS scFv and anti-HS stub antibodies. Nontreated (*a*) and heparinase III-treated (*b*, *c*) cryosections of mouse skeletal muscle tissue were incubated with periplasmic fraction of anti-HS antibody AO4F12 (*a*, *b*) or anti-heparan sulfate stub antibody (3G10) (*c*). Bound scFv antibodies were visualized by incubation with rabbit polyclonal anti-c-Myc IgG (*a1*, *b1*), followed by Alexa 488-conjugated goat anti-rabbit or anti-mouse IgG (*a* and *b*, or *c*, respectively). AChR clusters present in the neuromuscular junction were visualized using TRITC-conjugated α -bungarotoxin (*a2*–*c2*). Although in untreated tissue the AO4F12 epitope is clearly present in the muscle BL (*a1*), staining disappeared during heparinase treatment (*b1*), indicating the HS nature of the epitope. Staining of heparan sulfate stubs in heparinase-treated tissue showed HS to be present throughout the muscle BL (*c1*). Note the higher staining intensity of AO4F12 at NMJs (*a1*, *a2*, arrows), regardless of the overall quantity of HS in the NMJ (*c1*, *c2*, arrows). Scale bar, 50 μ m.

Wild-type CHO cells showed a clear surface staining at sites of cell–cell contact when incubated with antibodies AO4B05, AO4B08, AO4F12, RB4CB9, and RB4CD12 (Table 3, Fig. 4*a,b*), whereas incubation with RB4EA12 and RB4EG12 did not (Table 3, Fig. 4*c*). None of the glycosylation-defective CHO mutant cell lines showed any surface staining (Fig. 4). As in the S₂₇ cell line, some antibodies showed a distinct staining of perinuclear and cytosolic granules (Table 3, Fig. 4).

Anti-HS antibodies bind distinct HS epitopes in skeletal muscle basal lamina

Incubation of cryosections of human, rat, and mouse skeletal muscle with each of the anti-HS antibodies yielded a clear staining of the muscle BL, which was similar in the species examined (Table 3, Fig. 5). Staining patterns of the antibodies on muscle BL were mutually distinct, ranging from a strong staining of the entire BL (AO4B05, AO4B08, AO4F12, RB4CB9, and

Table 3. Immunostaining patterns of anti-HS antibodies

Tissue	AO4B05	AO4B08	AO4F12	RB4CB9	RB4CD12	RB4EA12	RB4EG12
Mature human, rat, and mouse skeletal muscle							
Extrasyntaptic basal lamina	++	++	++	++	++	+/-	-
Synaptic basal lamina	+++	++	+++	++	+++	+++	+
Endoneurium and perineurium	++	+	+++	+	+++	+++	+
Capillary basal lamina	++	++	++	+	+++	-	-
Arterial basal lamina	++	+	++	++	++	-	-
Electric organ (<i>T. marmorata</i>)							
Electrocyte, noninnervated face	++	+	+	+/-	++	+/-	+
Electrocyte, innervated face	++	+	+	++	++	+/-	+
Endoneurium	++	+	+	-	+	+/-	+
Perineurium	+/-	+	++	++	+	-	+/-
Rat embryo skeletal muscle (19 d <i>in utero</i>)							
Extrasyntaptic basal lamina	++	++	++	+	+++	-	+/-
Synaptic basal lamina	++	++	+++	++	++	-	++
Nerve basal lamina	++	++	+++	++	+++	+/-	+
Smooth muscle basal lamina (artery)	++	++	++	++	++	-	-
<i>C</i> ₂ <i>C</i> ₁₂ skeletal muscle cell line							
Cell surface (on contact places)	+++	++	++	++	+++	- <i>d</i>	- <i>d</i>
Basal lamina (during differentiation)	+++ ^a	++ ^a	+++ ^a	+	+++ ^b	- <i>d</i>	- <i>d</i>
AChR clusters	+++ ^a	++	+ ^a	++	+++	-	-
<i>S</i> ₂₇ cell line (defective in proteoglycan synthesis)							
Cell surface (on contact places)	-	- <i>d</i>	- <i>d</i>	- <i>d</i>	- <i>c</i>	- <i>d</i>	- <i>d</i>
Basal lamina (during differentiation)	-	- <i>d</i>	- <i>d</i>	- <i>d</i>	- <i>c</i>	- <i>d</i>	-
CHO cell line (wild type)							
Cell surface	++	++	++	+	+	- <i>c</i>	- <i>d</i>
CHO cell line (677 mutant)							
Cell surface	-	-	- <i>d</i>	- <i>c</i>	- <i>c</i>	- <i>c</i>	- <i>d</i>
CHO cell line (F17 mutant)							
Cell surface	-	- <i>d</i>	- <i>d</i>	- <i>c</i>	-	- <i>d</i>	- <i>d</i>
CHO cell line (745 mutant)							
Cell surface	-	-	-	- <i>c</i>	- <i>c</i>	- <i>d</i>	- <i>d</i>

Cryosections of human, rat, and mouse skeletal muscle, *T. marmorata* electric organ, and rat embryo, as well as fixed monolayers of CHO (wild-type and glycosylation-deficient mutants, grown to confluency), and *C*₂*C*₁₂ and *S*₂₇ cells (differentiated for 0–15 d) were incubated with periplasmic fractions containing anti-HS scFv antibodies. Bound antibodies were visualized by incubation with rabbit polyclonal anti-c-Myc IgG followed by Alexa 488-conjugated goat anti-rabbit IgG. AChR clusters present in muscle neuromuscular junction, *T. marmorata* electrocytes, and *C*₂*C*₁₂ myotubes were visualized using TRITC-conjugated α -bungarotoxin. Staining intensity: +++, very strong; ++, strong; +, moderate; +/-, weak; -, absent [*n* = 3 (cryosections), *n* = 2 (tissue cultures)].

^aStaining intensity decreased with differentiation level.

^bStaining intensity increased with differentiation level.

^cSmall granules within the cytosol: ++.

^dSmall granules around the nucleus: ++.

RB4CD12), to a staining concentrated in (RB4EA12), or almost exclusive for (RB4EG12) the sBL. Antibodies AO4B05, AO4F12, RB4CD12, RB4EA12, and RB4EG12 stained the sBL more intensely than the extrasynaptic BL. The BL of neural tissues showed very strong (AO4F12, RB4CD12, and RB4EA12), strong (AO4B05), or moderate (AO4B08, RB4CB9, and RB4EG12) staining. BLs of blood vessels showed strong to moderate (AO4B05, AO4B08, AO4F12, RB4CB9, and RB4CD12) or no (RB4EA12 and RB4EG12) staining. The latter two antibodies hardly stain muscle BL extrasynaptically and appear to be neuron- and synapse-specific. Most antibodies that stain blood vessels showed differences in staining intensity between arteries, large blood vessels, and capillaries.

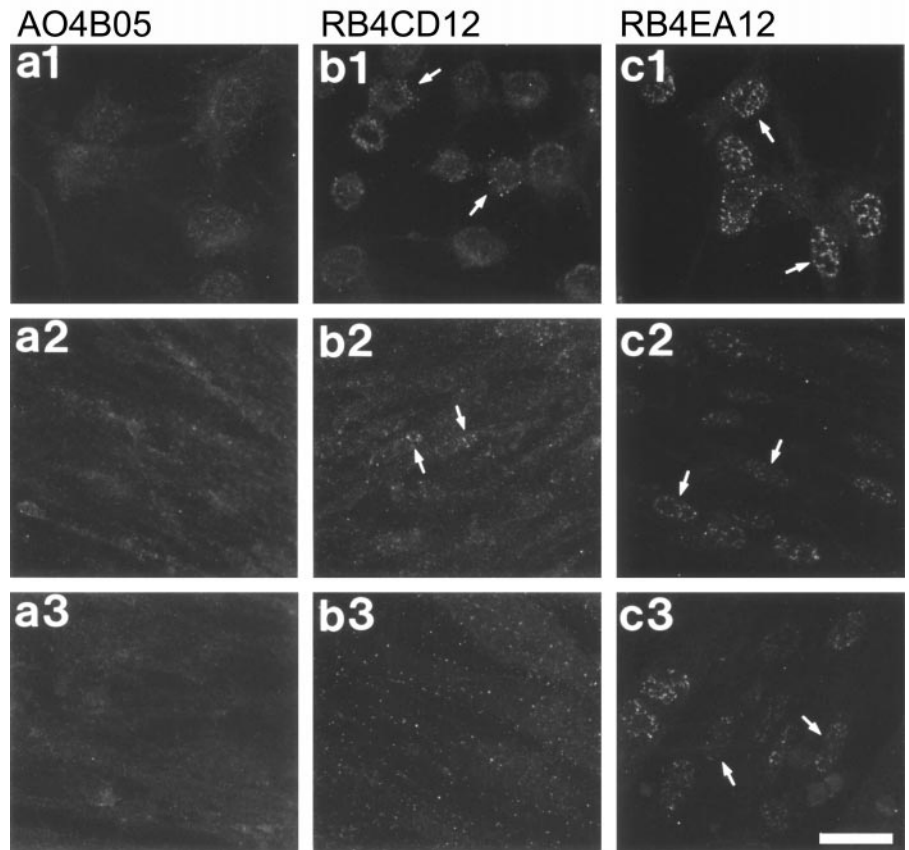
The staining patterns of anti-HS antibodies provided convincing evidence for the existence and unique distribution of multiple HS epitopes within the skeletal muscle BL. To investigate the distribution of these HS epitopes with regard to the sBL, cryosections containing NMJs were incubated both with

the antibodies and with TRITC-conjugated α -bungarotoxin. α -Bungarotoxin exclusively binds AChRs, thus allowing identification of NMJs. The AO4F12 epitope does not fully colocalize with AChR clusters, yet there is considerable overlap between the distribution of the AO4F12 epitope in the sBL and the presence of dense patches of AChR on the postsynaptic membrane (Fig. 5*a1–a3*). RB4CD12, on the other hand, showed an almost complete colocalization with AChR clusters (Fig. 5*b1–b3*). RB4EA12 showed a strong preference for neural and synaptic BL, thus completely colocalizing with AChR clusters in NMJs (Fig. 5*c1–c3*). Finally, the RB4EG12 epitope showed a moderate staining that was limited to neural and synaptic BLs only (Fig. 5*d1–d3*).

HS epitopes recognized by anti-HS antibodies abundant in *T. marmorata* electric organ

Because the anti-HS antibodies showed differential staining patterns with regard to nerve- and muscle-derived (extrasyn-

Figure 3. Staining of S_{27} muscle cell line with anti-HS scFv antibodies. S_{27} cultures were grown to confluency and subsequently differentiated up to 8 d. Cultures of different developmental stages [half confluent (*a1–c1*), 1 d (*a2–c2*), and 8 d (*a3–c3*) of differentiation] were fixed and subsequently incubated with the periplasmic fraction of scFv antibodies AO4B05 (*a*), RB4CD12 (*b*), and RB4EA12 (*c*), respectively. Bound antibodies were visualized by incubation with rabbit polyclonal anti-c-Myc IgG followed by Alexa 488-conjugated goat anti-rabbit IgG. None of the epitopes recognized by any of the antibodies can be visualized at the surface of myoblasts (*a1–c1*). For AO4B05, staining is not visible in aligning myoblasts (*a2*) or during myotube formation (*a3*). The epitope recognized by RB4CD12 is present in perinuclear granules in myoblasts (*b1*, arrows). During the alignment of myoblasts, the granular staining around the nucleus persists (*b2*, arrows) to change into a predominant cytosolic granular staining during myotube formation (*b3*). ScFv antibody RB4EA12 strongly stains perinuclear granules in myoblasts (*c1*, arrows). In aligning myoblasts and during myotube formation, the granular staining around the nucleus persists (*c2*, *c3*, arrows). Scale bar, 25 μ m.



aptic and synaptic) BLs, and to investigate whether the HS epitopes are present in nonmammalian species as well, we tested the antibodies for BL staining in the electric organ of the electric ray (*T. marmorata*). The electric organ is evolutionary derived from muscle tissue and shows dense patches of AChR clustering on the innervated face of the electrocytes. The various anti-HS antibodies showed a distinct staining of the electric organ (Table 3, Fig. 6), ranging from a strong staining of both electrocyte and endoneurial BLs [AO4B05 (Fig. 6*a1–a4*), AO4B08, AO4F12 (Fig. 6*b1–b4*), and RB4CD12], to a predominantly strong [RB4CB9 (Fig. 6*c2–c4*)], moderate (RB4EG12) or weak (RB4EA12) staining of the electrocyte BL. AO4B08, AO4F12 (Fig. 6*b2–b4*), and RB4CD12 stained specific regions of the electrocyte with a higher intensity. AO4F12 (Fig. 6*b1*), RB4CB9 (Fig. 6*c1*), and, to a lesser extent, AO4B08 and RB4CD12 stained the perineurial BL. The endoneurial BL was strongly stained by AO4B05 (Fig. 6*a1*) and to a lesser extent by AO4B08, AO4F12 (Fig. 6*b1*), RB4CD12, and RB4EG12.

Double-label micrographs of the epitope recognized by AO4B05 with α -bungarotoxin showed that the AO4B05 epitope partially colocalizes with the dense patches of AChR clusters on the innervated face of the electrocytes (Fig. 6*a4*). Double labeling of the AO4F12 epitope showed a slightly lower staining intensity of the noninnervated membrane, as compared with the innervated membrane, except for some brightly stained regions (Fig. 6*b4*). The RB4CB9 epitope was almost exclusively located at the electrocyte-innervated face, virtually completely colocalizing with the AChR clusters (Fig. 6*c4*).

Anti-HS antibodies show a developmental occurrence of HS epitopes in skeletal muscle basal lamina

The diversity of staining patterns obtained with the antibodies in mature skeletal muscle prompted us to investigate the occurrence of HS epitopes during muscle development. Special attention was paid to changes in the occurrence of specific HS epitopes within the endomysial, neural, and synaptic BL. This study was performed in three ways. First, cryosections of rat embryos at various developmental stages (days 10, 13, 16, and 19 *in utero*) were studied. In this way, the occurrence of and possible changes in BL–HS epitopes during muscular development and synaptogenesis could be studied in the presence of both muscular and neural tissue. Second, cultures of the mouse skeletal muscle cell line C_2C_{12} at developmental stages ranging from half-confluent to 15 d of differentiation were analyzed. In doing so, we could monitor the presence of and changes in HS epitopes during myogenesis, as well as during the clustering of AChRs in the presence of muscular tissue only. Third, cryosections of denervated skeletal muscle of rat were studied. In denervated muscle cells, we looked at a possible upregulation or downregulation of HS epitopes as a result of the regeneration process.

In early embryonic stages of the rat (days 10–16), strong staining of the endomysial as well as a distinct interaction with neural BL was observed on immunostaining with AO4B05, AO4B08, AO4F12, RB4CB9, and RB4CD12. Antibody RB4EA12 predominantly stained neural tissue, whereas RB4EG12 showed an amorphous staining in developing muscle regions (data not shown). Rat embryos at day 19 *in utero* showed a more defined organ texture in cryosections, which enabled us to examine the presence

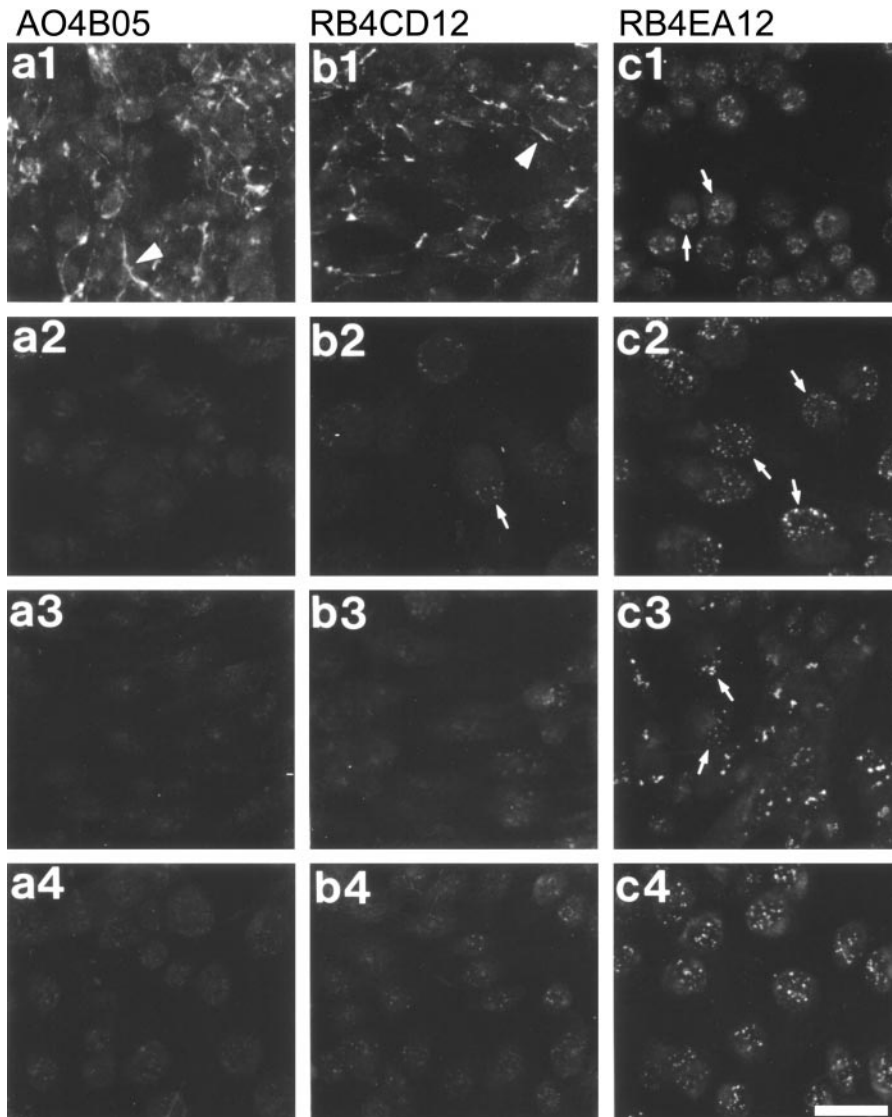


Figure 4. Staining of wild-type and glycosylation-deficient CHO cell lines with anti-HS scFv antibodies. CHO cultures [wild type (*a1–c1*), *N*-acetylglucosaminyl- and glucuronosyltransferase-deficient *pgsD-677* (*a2–c2*), heparan sulfate uronic acid 2-*O*-sulfotransferase-deficient *pgsF-17* (*a3–c3*), and xylosyl transferase-deficient *pgsA-745* (*a4–c4*)] were grown to confluency and subsequently fixed and incubated with the periplasmic fraction of scFv antibodies AO4B05 (*a*), RB4CD12 (*b*), and RB4EA12 (*c*), respectively. Bound antibodies were visualized as in Figure 3. The AO4B05 epitope is present to a high degree at the surface of wild-type CHO cells where cell–cell contacts are made (*a1*, arrowhead). Staining is not visible in any of the CHO mutant cell lines (*a2–a4*). Wild-type CHO cells are moderately stained at the surface by antibody RB4CD12 at places of cell–cell contact (*b1*, arrowhead). In CHO mutant cell line *pgsD-677* a faint granular perinuclear staining is visible (*b2*, arrow), whereas cell lines *pgsF-17* and *pgsA-745* show a slightly elevated background staining (*b3* and *b4*, respectively). The epitope recognized by RB4EA12 does not appear at the surface in any of the CHO cell lines but shows a distinct, perinuclear, and granular staining in all CHO cell lines used. In wild-type, *pgsD-677*, and *pgsF-17* cells, these granules are predominantly located at the perinuclear region on one side of the cell (*c1–c3*, arrows). In *pgsA-745* cells, the granular staining is present around the entire nucleus (*c4*). Scale bar, 25 μ m.

of the HS epitopes in greater detail (Table 3, Fig. 7). Although RB4CD12 showed a strong staining of the entire neural and endomysial BL, the staining intensity was markedly lower in the sBL (Fig. 7*a*). RB4CB9, on the other hand, stained the sBL considerably stronger than the extrasynaptic BL (Fig. 7*b*). RB4EG12, binding HS epitopes present in neural and synaptic BL in fully developed skeletal muscle, strongly interacted with HS epitopes within the sBL and showed a faint, although definite staining of the extrasynaptic BL (Fig. 7*c*). The epitope recognized by RB4EA12, preferentially staining neural tissue and sBL in mature muscle, could hardly be visualized in BL of skeletal muscle tissue at day 19 of rat embryogenesis. However, this antibody did stain large cytosolic granules (Fig. 7*d*). Staining of BL in tissues other than skeletal muscle was also observed (data not shown).

Cultures of mouse skeletal muscle cell line C_2C_{12} were incubated with antibodies at stages ranging from half-confluent to 15 d of differentiation (Table 3, Figs. 8–10). Immunostaining with AO4B05, AO4B08, AO4F12, RB4CB9, and RB4CD12 resulted in a strong staining of the myoblast surface. An intense staining was observed at places where myoblasts made contact. However, on alignment and fusion (processes that trigger BL formation), the

entire myotube surface was stained. AChR clusters, which develop on the surface of multinucleated myotubes at approximately day 3 of differentiation, were also stained by these antibodies. AO4B05 (Fig. 8), RB4CB9, and RB4CD12 (Fig. 9), especially, showed an enhanced staining of the myotube BL at sites of AChR clustering. A striking feature was that the overall staining intensity decreased strongly during differentiation for antibodies AO4B05 (Fig. 8), AO4B08, and AO4F12. Antibodies RB4EA12 and RB4EG12 were not able to stain the surface of C_2C_{12} cells at any stage of differentiation. Both antibodies stained small cytosolic granules that were predominantly present near the nuclei (Fig. 10).

Incubation of cryosections of rat skeletal muscle 11 d after denervation with various anti-HS antibodies did not result in staining patterns that were any different from control muscle. After heparinase treatment, no differences in staining intensity with anti-HS stub antibody 3G10 could be seen in BLs of denervated versus control muscle (data not shown).

DISCUSSION

In this paper, we report the selection of a set of unique anti-HS scFv antibodies. The HS epitopes recognized by these antibodies

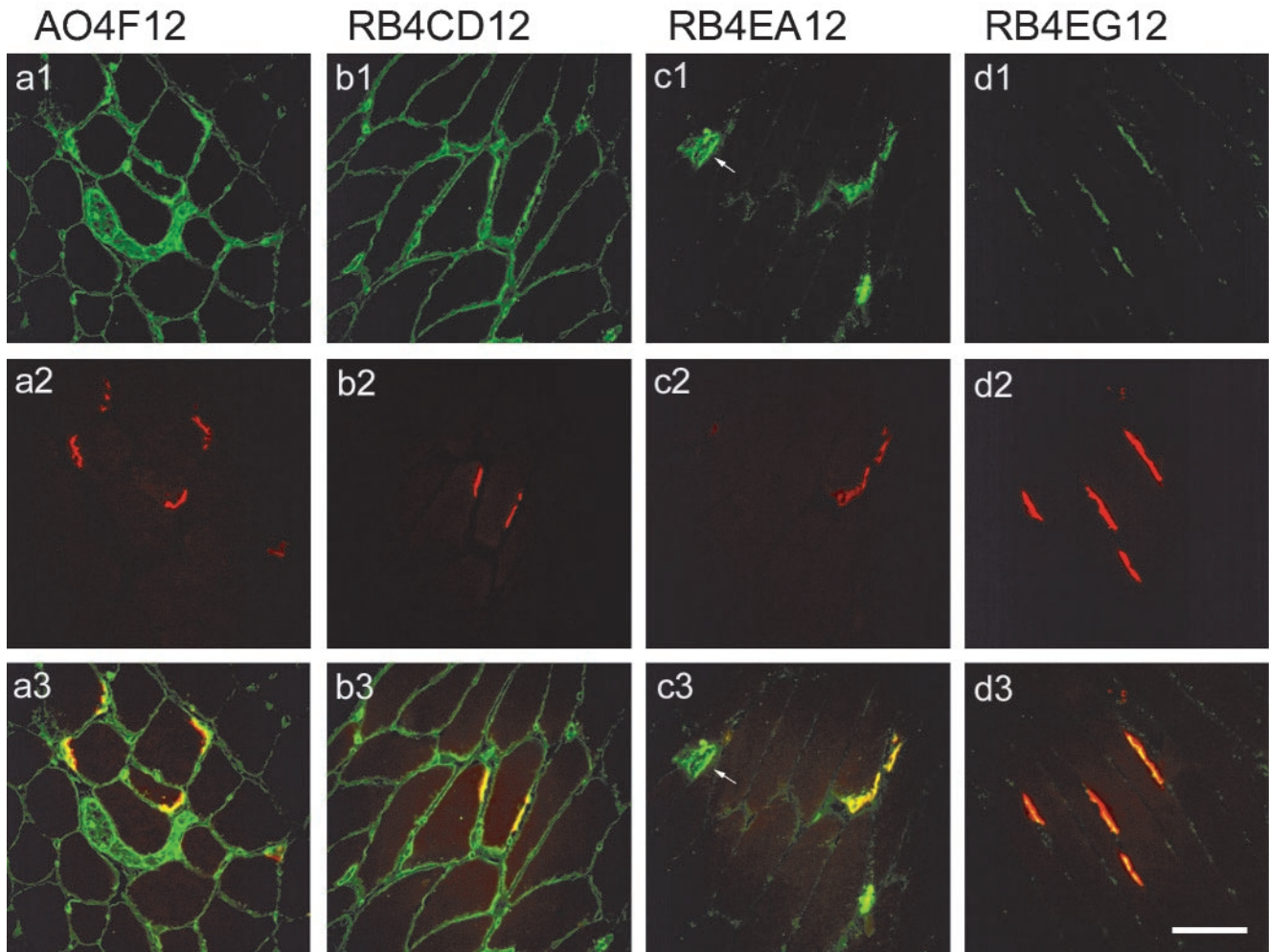


Figure 5. Staining of mouse skeletal muscle basal lamina with anti-HS scFv antibodies. Cryosections of C3H skeletal muscle were incubated with periplasmic fractions of anti-HS antibodies AO4F12 (*a*), RB4CD12 (*b*), RB4EA12 (*c*), and RB4EG12 (*d*), respectively. Bound antibodies were visualized by incubation with rabbit polyclonal anti-c-Myc IgG followed by Alexa 488-conjugated goat anti-rabbit IgG (*a1–d1*). AChR clusters present in the neuromuscular junction were visualized using TRITC-conjugated α -bungarotoxin (*a2–d2*). Double-label micrographs (*a3–d3*) show in yellow the colocalization of the HS epitopes bound by the scFv and AChR clusters. The epitope recognized by AO4F12 is present in endoneurial and perineurial as well as in endomysial BLs, but is clearly more abundant in synaptic versus extrasynaptic BL (*a1*). Note that this epitope does not fully colocalize with AChR clusters; there is a clear overlap from the BL epitope recognition (green) via a zone in which both epitopes are present (yellow) to the dense patches of AChR (red) (*a3*). The RB4CD12 epitope is also present throughout neural and endomysial BLs and is slightly more abundant at NMJs (*b1*) but covers the entire region of AChR clustering (*b3*). Antibody RB4EA12 stains epitopes present in neural BL to a larger extent than those present in endomysial BL (*c1, c3, arrows*), shows a high abundance in sBL (*c1*), and covers areas of AChR clustering entirely (*c3*). The epitope recognized by RB4EG12 hardly stains endomysial BL but resides in neural BL and at NMJs (*d1*), where it does not completely cover areas of AChR clustering (*d3*). Scale bar, 50 μ m.

are shown to be differentially distributed in BLs of both developing and mature skeletal muscle. GAG preparations isolated from mouse and human skeletal muscle specimens were used to select a series of anti-HS antibodies by phage display. Despite the enrichment of the muscle GAG preparations for DS, only anti-HS antibodies were selected. To our knowledge, no anti-DS antibodies have been described so far.

In ELISA, all anti-HS scFv antibodies showed a differential reactivity with several HS preparations and with heparin, reflecting the epitope specificity of each antibody. The requirement of both *N*- and *O*-sulfate groups for proper epitope recognition was shown by desulfation of heparin, which is known for its high number of disaccharide units and high levels of *N*-sulfation. Desulfation completely abolished recognition by all antibodies, and *N*-resulfation could not restore the heparin–antibody interaction. Because CS and DS are not bound by any of the antibod-

ies, sulfation patterns specific for HS are likely to be important in the structure of the epitopes involved.

In our experiments, CHO cells showed a distinct HS staining for most antibodies, which was less intense than the staining of C₂C₁₂ cells. This is probably because HS from CHO cells is relatively poorly sulfated [40–45% *N*-sulfation and ~0.8 sulfate/disaccharide (Bame et al., 1991)]. None of the cell-surface HS epitopes recognized by any of the antibodies described here could be detected in cell lines that are defective in GAG synthesis. This was the case with the S₂₇ cell line, a genetic variant of the C₂ mouse skeletal muscle cell line, which is severely hampered in GAG synthesis but does align and fuse to form myotubes during differentiation (Gordon and Hall, 1989). Several CHO mutants defective in GAG synthesis caused by the loss or impaired functioning of enzymes involved in glycosylation (Esko et al., 1985; Lidholt et al., 1992) failed to show any cell surface staining, which

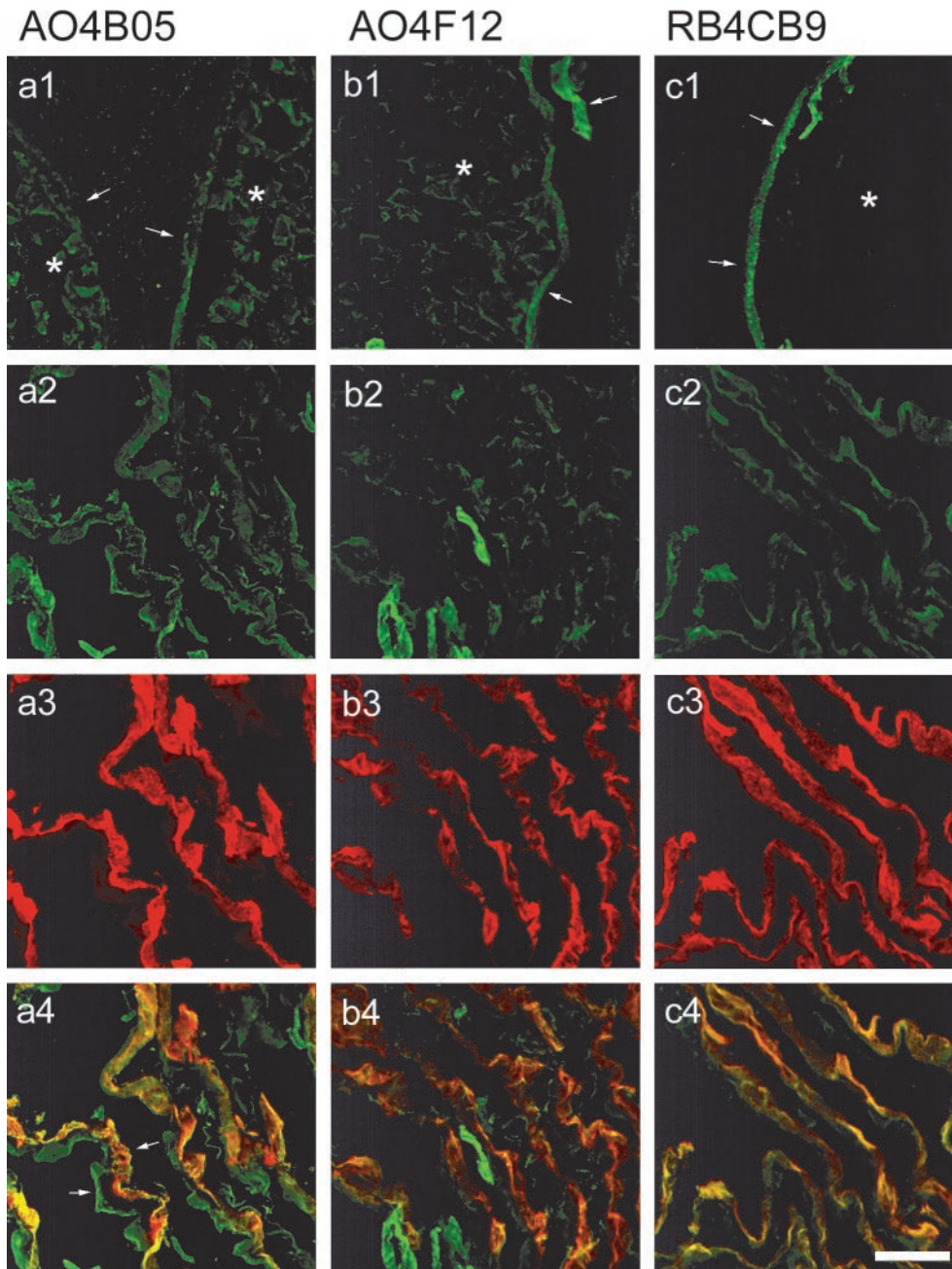


Figure 6. Staining of *Torpedo* electric organ with anti-HS scFv antibodies. Cryosections of *Torpedo* electric organ were incubated with periplasmic fractions of anti-HS scFv antibodies AO4B05 (*a*), AO4F12 (*b*), and RB4CB9 (*c*), respectively. Bound antibodies (*a1–c1* and *a2–c2*) and AChR clusters (*a3–c3*) were visualized as in Figure 5. Double-label micrographs (*a4–c4*) show in yellow the colocalization of AChR clusters and the HS epitopes bound by the scFvs. The AO4B05 epitope is present in large quantities in the endoneurium (*a1*, asterisks) and in electrocyte BLs (*a2*) but hardly at all in the perineurium (*a1*, arrows). Double staining of electrocytes reveals that this epitope is present not only on the innervated face but throughout the electrocyte BL (*a4*, arrows). The epitope recognized by AO4F12 is present in the endoneurium (*b1*, asterisk) but is more abundant in the perineural BLs (*b1*, arrows). In electrocytes, the presence of this epitope on the innervated face is less pronounced, whereas specific regions of the noninnervated membrane accommodate the epitope in large amounts (*b2*). Double staining shows this epitope to be partially colocalized with AChR clusters on the electrocyte-innervated membrane (*b4*). The endoneurium is not stained by antibody RB4CB9 (*c1*, asterisk), whereas this epitope is very abundant in the perineurium (*c1*, arrows). In electrocytes, its presence is restricted to the innervated membrane (*c2–c4*). Scale bar, 50 μ m.

indicates their inability to properly synthesize the HS epitopes involved. The granular staining seen with some antibodies in many cells may reflect the staining of certain cellular compartments such as the Golgi apparatus or lysosomes. Because of the defective cellular machinery for the correct synthesis of GAGs, immature HS epitopes or degradation products of HS molecules may be confined to these organelles.

All anti-HS scFv antibodies showed distinct reactivity in immunofluorescence with the BL of mature skeletal muscle. Staining patterns of the antibodies on human and rat muscle were consistent with those obtained on mice, reflecting an interspecies conservation of the epitopes involved. Most antibodies stained the entire muscular BL, but some antibodies showed a more intense staining in synaptic regions. Because of the presence of junctional folds in the postsynaptic membrane, BL is two- to

threefold more abundant at NMJs than extrasynaptically (Sanes and Chiu, 1983). This local concentration of BL might explain the higher staining intensity of some antibodies at the NMJ, but we did not observe a higher abundance of HS in the synaptic cleft by heparitinase III digestion and anti-stub staining. A more appealing explanation is the possibility that certain HS epitopes are specifically concentrated in the sBL. The incomplete overlap of the AO4F12 epitope with AChR clusters, in contrast with e.g., RB4CD12 and RB4EA12 (Fig. 5), suggests differences in location of these epitopes within the sBL. Antibodies that predominantly recognize epitopes present in neural and synaptic BL, such as RB4EA12 and RB4EG12, may indicate the neural origin of the epitopes involved. Results obtained on aneurally cultured skeletal muscle cells support this view, because these antibodies did not stain BLs at sites of AChR clustering (see further). The synapse-

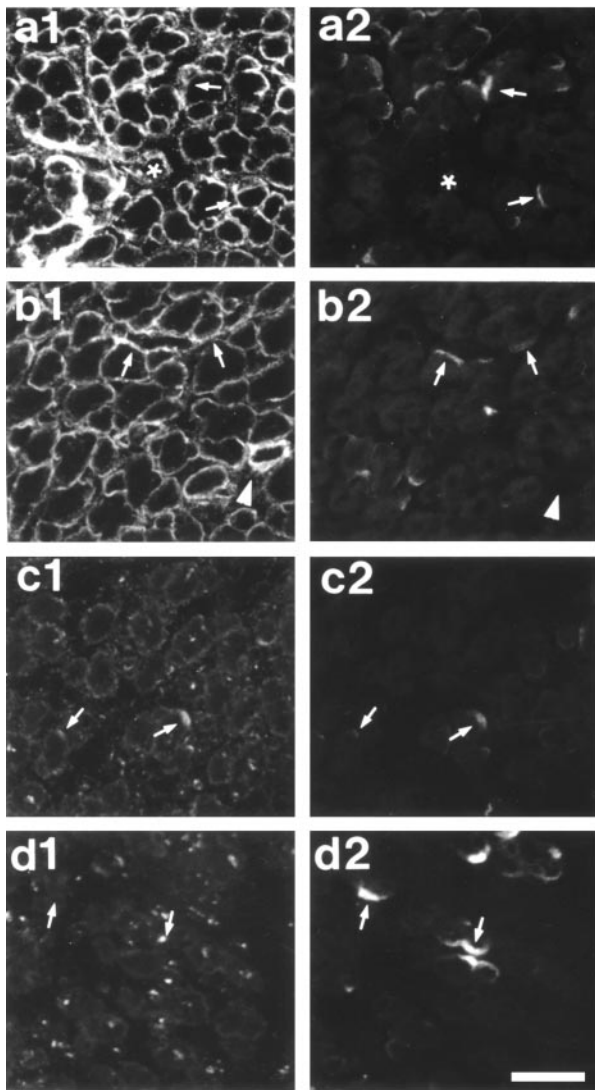


Figure 7. Staining of rat embryo skeletal muscle with anti-HS scFv antibodies. Cryosections of rat embryos (day 19 *in utero*) were incubated with periplasmic fractions of anti-HS scFv antibodies RB4CD12 (*a*), RB4CB9 (*b*), RB4EG12 (*c*), and RB4EA12 (*d*), respectively. Bound antibodies (*a1–d1*) and AChR clusters (*a2–d2*) were visualized as in Figure 5. The epitope recognized by antibody RB4CD12 is present throughout neural (asterisk) and endomysial BLs (*a1*) and is slightly less abundant in the sBL of developing NMJs (*a1, a2, arrow*). The epitope recognized by RB4CB9 is present in smooth muscle BL (arrowheads point at an artery) throughout the endomysial BL (*b1*) but is more abundant at NMJs (*b1, b2, arrows*). RB4EG12 stains, at a low level, throughout the endomysial BL (granular staining in *c1*), but the epitope involved is more abundant at developing NMJs (*c1, c2, arrows*). The epitope recognized by RB4EA12 is only slightly present in the endomysial BL (*d1*) and absent in sBL (*d1, d2, arrows*). Scale bar, 25 μ m.

specific occurrence of distinct HS epitopes may prove to be causal for the restricted location of NMJ-resident, HS-binding proteins such as agrin, acetylcholine esterase, growth factors, and certain laminin isoforms.

Most HS epitopes recognized by the antibodies proved to be located close to AChR clusters, present on the innervated face of electrocytes, in the electric organ of the electric ray (*T. marmarota*). Anti-HS antibodies recognized their epitopes, which were embedded in mutually distinct patterns and quantities within neural BLs and in BLs on both the innervated and non-

AO4B05

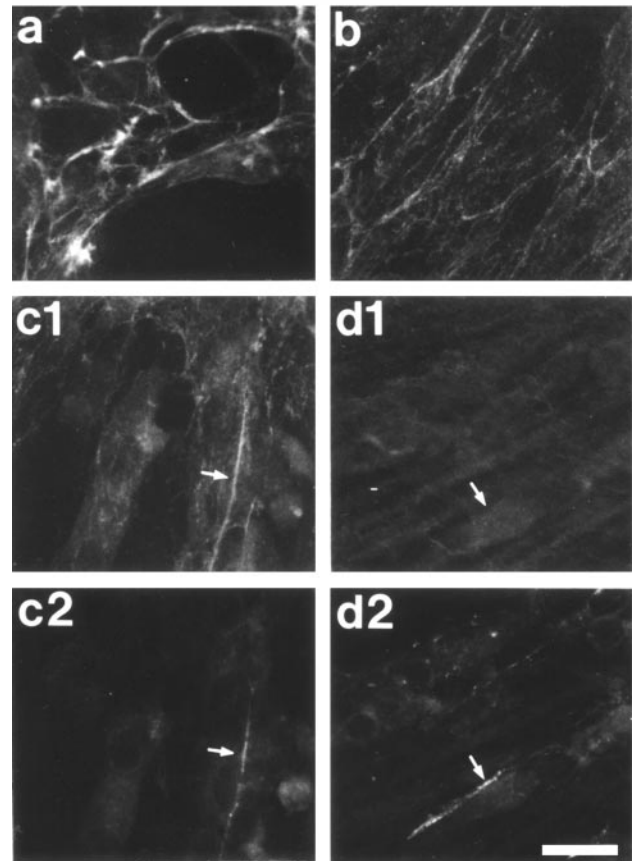


Figure 8. Staining of C_2C_{12} muscle cell line with scFv antibody AO4B05 during differentiation. C_2C_{12} cultures were grown to confluency and subsequently differentiated up to 15 d. Cultures of different developmental stages [half confluent (*a*), 1 d (*b*), 8 d (*c1, c2*), and 15 d (*d1, d2*) of differentiation] were fixed and subsequently incubated with the periplasmic fraction of scFv antibody AO4B05. Bound scFv antibodies (*a, b, c1, d1*) and AChR clusters present on the surface of multinucleated myotubes (*c2, d2*) were visualized as in Figure 5. The epitope recognized by AO4B05 is present to a high degree at the surface of myoblasts in regions where cell–cell contacts are made (*a*). In aligning myoblasts (*b*), a clear surface staining is visible that becomes more pronounced after myotube formation at sites where AChR clusters develop (*c1, c2, arrows*). As differentiation proceeds, both the staining of the BL and the staining at AChR clusters decrease (*d1, d2, arrows*). Scale bar, 25 μ m.

innervated side of the electrocytes. Despite the conserved distribution of the epitopes with regard to neural, synaptic, and extra-synaptic BL among the mammals tested, the distribution within the elasmobranch electric organ appeared to differ. Extracellular matrix isolated from *Torpedo* electric organ can induce AChR clustering in fibroblasts (Hartman et al., 1991). The heavily glycosylated HSPG agrin appears to be involved in the clustering of AChR in *Torpedo* electrocytes (Cartaud et al., 1996). The staining patterns of our anti-HS antibodies on cryosections of the electric organ add proof to the mutually distinct HS epitopes involved and raise curiosity about their function in organ morphogenesis.

During myogenesis in developing rat embryos, some of the HS epitopes were present in endomysial and synaptic BL in a pattern different from that seen in mature muscular tissue. Because NMJs appear between day 14 and 16 of embryonic life (Engel, 1994), the occurrence of HS epitopes during synaptogenesis was investi-

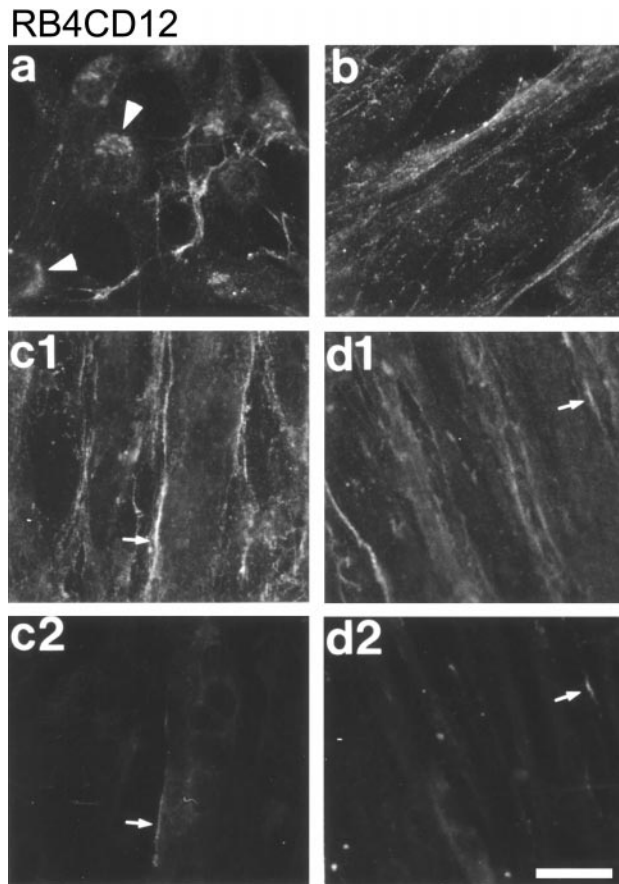


Figure 9. Staining of C_2C_{12} muscle cell line with scFv antibody RB4CD12 during differentiation. C_2C_{12} cultures were grown to confluency and subsequently differentiated up to 15 d. Cultures of different developmental stages [half confluent (*a*), 1 d (*b*), 8 d (*c1*, *c2*), and 15 d (*d1*, *d2*) of differentiation] were fixed and subsequently incubated with the periplasmic fraction of scFv antibody RB4CD12. Bound scFv antibodies (*a*, *b*, *c1*, *d1*) and AChR clusters present on the surface of multinucleated myotubes (*c2*, *d2*) were visualized as in Figure 5. The RB4CD12 epitope can be visualized at the myoblast surface at sites where cells have made mutual contacts (*a*). Note the perinuclear staining of the myoblasts at half-confluent stage (*a*, arrowheads). After alignment, a strong surface staining is visible (*b*). During myotube formation, this staining intensifies, especially at sites of AChR clustering (*c1*, *c2*, arrows). Ongoing differentiation does not lead to reduced AChR cluster staining, whereas the overall staining of the BL decreases slightly (*d1*, *d2*, arrows). Scale bar, 25 μ m.

gated on cryosections of embryos at days 10–19 *in utero*. Most antibodies stained endomyisial as well as neural BLs during embryonic muscular development, as may be expected on the basis of their staining patterns in mature skeletal muscle tissue. However, clear differences in developmental appearance could be distinguished for epitopes recognized by some antibodies (RB4CB9, RB4CD12, RB4EA12, and RB4EG12), especially at sites of synaptogenesis. Local binding of growth factors and cytokines to specific HS sequences, as reviewed recently by Lyon and Gallagher (1998), may prove to be elemental in tissue morphogenesis. The distinct distribution in both time and space of these HS epitopes argues for such a regulatory mechanism.

The HS epitopes recognized by our antibodies were present in C_2C_{12} skeletal muscle cells at various stages of differentiation. Aneurally grown C_2C_{12} myoblasts start aligning when they reach confluency. When culture medium is changed to differentiation medium containing 10% rat brain extract, AChR clusters appear

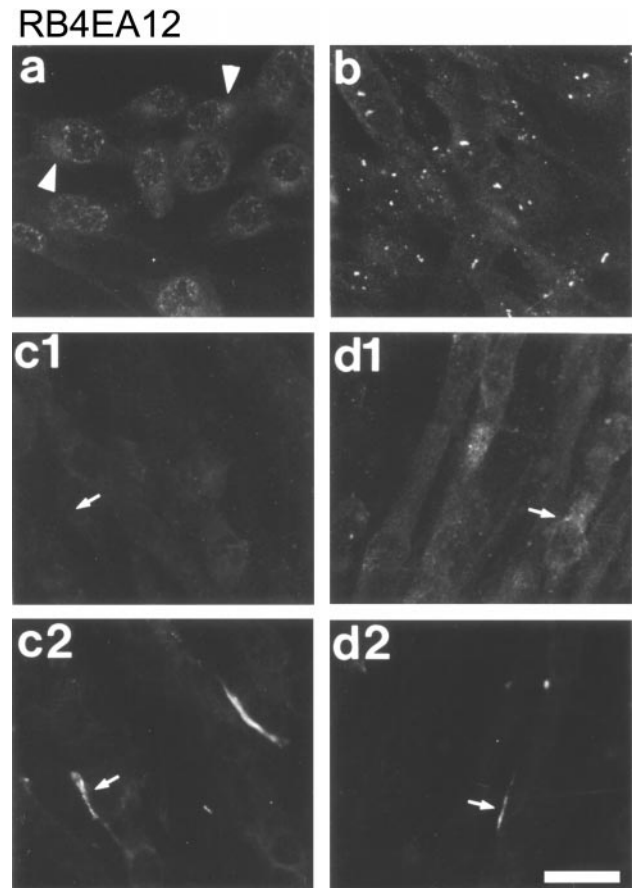


Figure 10. Staining of C_2C_{12} muscle cell line with scFv antibody RB4EA12 during differentiation. C_2C_{12} cultures were grown to confluency and subsequently differentiated up to 15 d. Cultures of different developmental stages [half confluent (*a*), 1 d (*b*), 8 d (*c1*, *c2*), and 15 d (*d1*, *d2*) of differentiation] were fixed and subsequently incubated with the periplasmic fraction of scFv antibody RB4EA12. Bound scFv antibodies (*a*, *b*, *c1*, *d1*) and AChR clusters present on the surface of multinucleated myotubes (*c2*, *d2*) were visualized as in Figure 5. The epitope recognized by RB4EA12 is not present at the surface of myoblasts in regions where cell–cell contacts have been made (*a*). Note the faint staining in the perinuclear region of myoblasts at the half-confluent stage (*a*, arrowheads). A granular staining within the cytosol is visible in early stages of alignment (*b*) but disappears during myotube formation (*c1*). During further differentiation, no BL staining is seen at AChR clusters (*d1*, *d2*, arrows). Nevertheless, a faint perinuclear staining occurs in some myotubes displaying AChR clusters (*d1*, *d2*, arrows). Scale bar, 25 μ m.

at approximately day 3 of differentiation (Portier et al., 1999). On mutual contact, C_2C_{12} myoblasts expressed most of the HS epitopes described in this paper in large quantities on their surface. Alignment and fusion resulted in a complete staining of the newly formed BL by corresponding antibodies. These observations are in accordance with the threefold increase in HS synthesis in myotube cultures, compared with proliferating or aligning cultures (Noonan et al., 1986). Some antibodies showed steady levels or even a marked increase in overall staining intensity of the BL during further differentiation, consistent with the upregulation of the HSPG glypican during C_2C_{12} differentiation (Brandan et al., 1996). Overall BL staining intensity of other antibodies decreased during later stages of differentiation. These results may be related to observations of Larraín et al. (1997a,b) on downregulation of the HSPGs perlecan and syndecan-1 during C_2C_{12} cell differentiation. AChR cluster formation was accompa-

nied by a strong local increase of certain HS epitopes, arguing for a possible role of these epitopes in the clustering of this ion channel. Antibodies RB4EA12 and RB4EG12 were not capable of BL staining at any stage of C₂C₁₂ cell differentiation, in accordance with their supposed neural origin.

Attempts to detect possible changes in the abundance of HS epitopes in denervated skeletal muscle proved to be elusive. Endomysial and neural BLs persist after damage or degeneration of either muscle or nerve cells, or both (Hall and Sanes, 1993). Synaptic and extrasynaptic proteoglycan deposits are conserved in both size and morphology in denervated skeletal muscle (Anderson et al., 1984), serving as scaffolds for the regeneration of both muscle and nerve tissue, thus causing NMJs to develop at sites where they were present before the degeneration. Moreover, Fadic and coworkers (1990) reported proteoglycan synthesis to be upregulated after denervation. Recently, GAGs were shown to be potent stimulants of insulin-like growth factor-1-mediated muscle reinnervation (Gorio et al., 1998). Because HS binds several growth factors involved in tissue morphogenesis and because of the unique distribution of certain HS epitopes, we suspect certain roles for HS epitopes in this regeneration process.

In conclusion, we show that it is possible to select for highly specific anti-HS antibodies against GAG preparations from skeletal muscle. The antibody-defined HS epitopes have distinct distribution characteristics in skeletal muscle BL and are similarly distributed in humans, rats, and mice. Obvious differences in extrasynaptic and synaptic BL staining were observed in mature versus developing skeletal muscle. The unique distribution patterns in skeletal muscle of the HS epitopes recognized by the scFv antibodies described in this article, both in time and in space, raise questions as to the biological roles of these HS epitopes. Of special interest are their roles in myogenesis, more specifically in synaptogenesis and the accompanying postsynaptic specializations such as the clustering of AChRs and other ion channels. The occurrence of these HS epitopes in HSPGs that have already been implicated in developmental processes awaits further investigation. Tools are now available to study more accurately the role of HS epitopes separate from their core protein.

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