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(DAPC) present in the nucleus of HeLa cells: Members of the nuclear DAPC associate with the nuclear matrix

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ABSTRACT: Dystrophin is an essential component in the assembly and maintenance of the dystrophinassociated protein complex (DAPC), which includes members of the dystroglycan, syntrophin, sarcoglycan and dystrobrevin protein families. Distinctive complexes have been described in the cell membrane of different tissues and cultured cells. In this work, we report the identification and characterization of a novel DAPC present in the nuclei of HeLa cells, which contains dystrophin Dp71 as a key component. Using confocal microscopy and cell fractionation analyses, we found the presence of Dp71, β -sarcoglycan, β -dystroglycan, α -and β -syntrophin, α 1-and β -dystrobrevin and nNOS in the nuclei of HeLa cells. Furthermore, we demonstrated by coimmunoprecipitation experiments that most of these proteins form a complex in the nuclear compartment. Next, we analyze the possible association of the nuclear DAPC with the nuclear matrix. We found the presence of Dp71, β dystroglycan, nNOS, β -sarcoglycan, α/β syntrophin, α 1-dystrobrevin and β dystrobrevin in the nuclear matrix protein fractions and in situ nuclear matrix preparations from HeLa cells. Moreover, we found that Dp71, β-dystroglycan and β -dystrobrevin co-immunoprecipitated with the nuclear matrix proteins lamin B1 and actin. The association of members of the nuclear DAPC with the nuclear matrix indicates that they may work as scaffolding proteins involved in nuclear architecture.

Introduction

Dystrophin, the product of the Duchenne muscular dystrophy gene (DMD) [1], is an essential component of a multi-protein complex collectively termed the dystrophin-associated protein complex (DAPC) [2]. Proteins that comprise the DAPC are structurally organized into three distinct subcomplexes: the cytoskeletal proteins dystrophin, dystrobrevins (α and β subunits) and syntrophins (α , β and γ subunits); the dystroglycans (α and β subunits); and the sarcoglycans (α , β , γ , δ and ε subunits). In skeletal muscle, the DAPC is assembled around dystrophin; this scaffold links the actin cytoskeleton to the basement membrane via the transmembrane protein β dystroglycan and anchors the syntrophins and dystrobrevins to the muscle membrane [2,3]. Loss of dystrophin leads to disassembly of the complex and muscle degeneration [3]. It has been shown that neuronal nitric oxide synthase (nNOS) binds to skeletal muscle syntrophin through PDZ domain interactions [4]; thus, the DAPC may also regulate signal transduction.

Many of the DAPC components found in skeletal muscle, including β -and ϵ -sarcoglycans, dystroglycans and syntrophins, are also expressed in other tissues [5–8]. Likewise, several dystrophin isoforms are ubiquitously expressed [9,10]. Therefore, distinct DAPC may form in different tissues implying that they may have important roles outside skeletal muscle. In fact, the characterization of different DAPCs in non-muscle tissues as well as cultured cells has been recently documented [11–13].

Dystrophin Dp71 is a DMD gene isoform produced from an internal promoter located on intron 62 [14]. As dystrophin Dp71 is the most widely expressed DMD gene product, it is expected that a number of non-muscle DAPC have this protein as a key component. In fact, the association of Dp71 with proteins of the DAPC such as β -dystroglycan, dystrobrevins and syntrophins has been reported [15–17]. Moreover, it has been evidenced that Dp71 is necessary for the anchorage and/ or organization of the brain DAPC [18].

An additional complication in understanding the diversity of DAPC functions has emerged from the evidence showing that several DAPC components are present in the nuclear compartment, such as α -and β -dystrobrevins [17,19] and α -, β -and γ 1-syntrophins [20]. Moreover, by using green fluorescence protein-Dp71 protein fusions (GFP-Dp71), we have previously shown that Dp71 splicing variants containing amino acids encoded by exon 71 and/or 78 distribute predominantly in the nuclei of HeLa cells [21].

Although the evidence mentioned above raises the possibility that a Dp71containing DAPC (Dp71-DAPC) exists in the nucleus, the biochemical characterization of such protein complex remains to be approached. To address this issue, we analyzed the expression and subcellular distribution of Dp71 splicing isoforms and DAPC components in HeLa cells, a human cell line widely used to characterize the molecular mechanisms underlying the function of nuclear proteins. Using confocal microscopy and cell fractionation analyses we showed the presence of Dp71, β -sarcoglycan, β dystroglycan, α -and β -syntrophins, α 1-and β -dystrobrevins and nNOS in the nuclei of HeLa cells. Moreover, we demonstrated by co-immunoprecipitation experiments that most of these proteins form a complex in the nuclear compartment. Nuclear fractionation analyses together with in situ nuclear matrix preparations showed that Dp71, β dystroglycan, α 1-dystrobrevin, β -dystrobrevin, α/β syntrophin, β -sarcoglycan and nNOS, components of the nuclear DAPC, are present in the HeLa nuclear matrix. Furthermore, we found that Dp71, β -dystroglycan and β -dystrobrevin copurified with the nuclear matrix proteins lamin B1 and actin. Our findings suggest the presence of a DAPC in the nuclei of HeLa cells, and the association of several members of this complex with the nuclear matrix may indicate their participation as nuclear scaffolding proteins.

Materials and methods

Cell culture

HeLa cells were grown in Dulbecco's modified Eagle medium (DMEM), (Invitrogen, Carlsbad, CA, USA) supplemented with 10% (v/v) neonatal serum, 25 U/ml penicillin and 25 μ g/ml streptomycin (Invitrogen), plated on 100-mm-diameter dishes and maintained at 37°C in a humidified incubator with CO2 atmosphere.

RNA extraction and RT-PCR

Total RNA was isolated from confluent cell cultures using a High Pure RNA Isolation Kit (Roche Applied Science, Indianapolis, IN, USA). RNA concentration was determined spectrophotometrically at OD 260 nm. For cDNA synthesis, 1 µgof total RNA was primed with random hexadeoxynucleotides and reverse transcribed using 2 U of MuLV reverse transcriptase (Invitrogen), according to manufacturer's instructions. PCR reactions, containing 2.5 µl of cDNA/20 mM Tris–HCl (pH 8.4)/50 mM KCl/1.5 mM MgCl2/2.5 µM of each dNTP/200 ng of each oligonucleotide, and 1U of Taq DNA polymerase (Invitrogen) were performed in a Gene Amp PCR System 2400 Thermal Cycler (Perkin Elmer, Boston, MA, USA). The sequence of primers and the expected size for the amplified products are given in Table 1.

Immunofluorescence and confocal microscopy analysis

Cells, plated on coverslips, were washed with cytoskeleton buffer (10mM MES/150mM NaCl/5mM EGTA/5mM MgCl2/ 5mM glucose), fixed in 2% (w/v) paraformaldehyde in PBS for 15min at room temperature and permeabilized with 0.1% (v/v) Triton X-100 in PBS for 5min. Fixed cells were blocked 20min with 0.5% (w/v) gelatin in PBS and incubated overnight at 4°C with primary antibodies (Table 2). The day after, cells were washed with PBS and incubated for 1h with a Cy3-conjugated secondary anti-rabbit antibody, a fluoresceinconjugated (FITC) secondary anti-mouse IgG antibody (Amersham Biosciences, UK, England) or FITC secondary anti-rabbit IgG antibody (Amersham Biosciences). For counterstaining, cells were incubated for 20min at room temperature with Texas Red-phalloidin (Molecular Probes) or during 5 min with propidium iodide (PI) (Sigma, St. Louis, MO, USA). After washing, coverslips were mounted on microscope slides with VectaShield (Vector Laboratories Inc. Burlingame, CA, USA) and viewed throughout a confocal laser scanning microscope (TCP-SP2, Leica, Heidelberg Germany) using 63× and 100× oilimmersion plan apochromat objectives (NA 1.32 and 1.4, respectively). Sixtyeight consecutive single sections were obtained simultaneously for one or two channels throughout the Z axis of the sample. The resulting stack of images were

projected and analyzed onto the two-dimensional plane using a pseudocolor display green (FITC) and red (Texas Red and Cy3) for double labeling experiments. Fluorochromes were excited at 488nm (for FITC) wavelength for single labeling, and additionally at 560nm (for Texas Red and Cy3) wavelength in double labeling experiments. Single optical sections of the stacks were selected to analyze the colocalization patterns between two markers.

Table 1 Expression of DAPC members by RT-PCR analysis				
mRNA	Sequence (5'–3')	Product size	Expression	
β-Dystroglycan	F-gtggtggaatggaccaacaa R-agcaatgattccagcaatga	348 pb	Positive	
α-Sarcoglycan	F-gcctacaatcgagacagctt R-agcccaccttgatgtatacc	283 pb	Negative	
β-Sarcoglycan	F-gtttttcagcaagggacaac R-atcggcacacatgcagag	444 pb	Positive	
γ-Sarcoglycan	F-tgctcatcatccttgttgtg R-gcagacaggtacagcttcc	700 pb	Negative	
δ-Sarcoglycan	F-acgccaaagaaatccagtcc R-ggtctgctctgacattaggt	291 pb	Negative	
α-Syntrophin	F-caggacatcaagcagattgg R-aagtttctcaagggctgc	498 pb	Positive	
β2-Syntrophin	F-atgccgatcctcatctccaa R-atggcagccctgaacaagta	902 pb	Positive	
y2-Syntrophin	F-ttagggtccccagggccatc R-tggacccggttctctgaact	774 pb	Negative	
α1-Dystrobrevin	F-gggtacattcacagtggagc R-tgggtttcagcctgtgtt	569 pb	Positive	
β-Dystrobrevin	F-catctcctccatctactatc R-ccagcatttttccaccacac	185 pb	Positive	
nNOS	F-tcggtgttcagcaaatccac R-cctcctgcccatcatcgt	491 pb	Positive	

Table 2 Antibodies used in this study					
Antibody	Position of antigen	Nature	Specificity	Reference/resource	
Dys2	Peptide 3669–3685	Monoclonal	Dp71d isoform	Novocastra	
2166	C-terminal 17 amino acids	Polyclonal	Dp71d isoform	[17]	
5F3	Alternative C-terminal 31aa	Monoclonal	Dp71f isoform	[45]	
11.11	(rounder sequence)	D 1 1 1	- D. (1	[LC]	
JAF	C-terminal 7 amino acids	Polyclonal	B-Dystroglycan	[46]	
β-521	Peptide 521–738	Polyclonal	β-Dystrobrevin	[47]	
β-CTFP	Peptide 459–465	Polyclonal	β-Dystrobrevin,	[47]	
			α-dystrobrevin		
			1 and 2		
α-CTFP	C-terminal of α1-dystrobrevin	Polyclonal	α1-Dystrobrevin	[47]	
mDp71	First 7 amino acids	Polyclonal	Dp71	Blake D.	
nNOS	Peptide 1095–1289	Monoclonal	Neuronal nitric oxide	Transduction	
			synthase	laboratories	
NCL-b-DG	Peptide PKNMTPYRSPPPYVP-P	Monoclonal	β-Dystroglycan	Novocastra	
Pan-syntrophins	Peptide VTRLGLLA	Polyclonal	α/β-Syntrophin	Mornet D.	
β-sarcoglycan	Peptide AGYIPIDEDRL	Polyclonal	β-Sarcoglycan	[48]	
Actin	Central domain of actin	Monoclonal	All isoforms of actin	[49]	
Lamin B1	Non-specified	Monoclonal	Lamin B1 isoform	Zymed	
NCL-b-SARC	N-terminal of sarcoglycan	Monoclonal	β-Sarcoglycan	Novocastra	
Sp3	C-terminal of Sp3	Polyclonal	Sp3 isoforms	Santa Cruz	
			••••	Biotechnology	
Vinculin	N-terminal of vinculin	Polyclonal	Vinculin	Santa Cruz	
				Biotechnology	
Calnexin	Amino acids 1–70	Polyclonal	Calnexin	Santa Cruz	
				Biotechnology	
α-Syntrophin	N-terminal of α -syntrophin	Polyclonal	α-Syntrophin	Santa Cruz	
				Biotechnology	
β2-Syntrophin	Internal region of β2-syntrophin	Polyclonal	β 2-Syntrophin	Santa Cruz	
				Biotechnology	

Isolation of total, cytosolic and nuclear protein extracts

To obtain total extracts, we washed cell dishes with cold PBS and suspend in complete protease inhibitor mixture (Roche Applied Sciences) containing 50mM Tris–HCl (pH 8.0) and 0.1% (w/v) DNase I. Homogenates were sonicated for 30s and protein concentrations were determined by the Bradford protein dye binding method (Bio-Rad, Hercules, CA, USA) using bovine serum albumin as standard. To obtain nuclear and cytosolic protein extracts, we harvested cells in cold buffer I [0.32M sucrose/10mM Tris–HCl (pH 8.0)/3mM CaCl2/ 2mM magnesium acetate/0.1mM EDTA/1mM DTT/0.5mM PMSF/0.5% (v/v) NP-40/complete protease inhibitor mixture], Dounce homogenized and spun at 200×g for 10min at 4°C. The supernatant was saved as a cytosolic fraction, and the pellet was suspended in buffer I, mixed with buffer II [2M sucrose/ 10mM Tris–HCl (pH 8.0)/5mM magnesium acetate/0.1mM EDTA/1mM EDTA/1mM DTT/0.5mM PMSF/complete

Gel electrophoresis and Western blot

Equal amounts of protein extracts (80µg) were mixed with an equalvolumeof 2×samplebuffer[50mMTris-HCl (pH 6.8)/15% (w/v)SDS/5% (v/v)Bmercaptoethanol/20% (v/v) glycerol/0.1% (w/v) bromophenol blue] and boiled for 3min. Proteins were resolved by 10% (v/v) sodium dodecyl sulfatepolyacrylamide gel electrophoresis (SDS-PAGE) and electrotransferred onto nitrocellulose membrane. Membranes were blocked for 1h in TBS-T [150mM NaCl/100mM Tris-HCl (pH 8.0)/0.5% (v/v) Tween-20] with 6% (w/v) low-fat dried milk and then incubated overnight at 4°C with the respective primary antibody (Table 2). When anti-rabbit antibodies were employed, a further incubation step in TBS [150mM NaCl/ 100mM Tris-HCl (pH 8.0)] with 2.5% (v/v) β -mercaptoethanol was carried out. Following three washes in TBS-T, membranes were incubated with the appropriate horseradish peroxidaseconjugated secondary antibody (Amersham Biosciences) and developed using the ECL Western blotting analysis system (Amersham Biosciences).

Immunoprecipitation

Nuclear extracts obtained in a final volume of 250µl were incubated for 1h at 4°C with 5µg of the appropriate polyclonal antibody, previously bound to protein A-agarose. As control for non-specific interactions, parallel incubations with no polyclonal antibodies were performed. The immune complexes were precipitated by centrifuging for 2min at 500 × g and washed twice in NET-BSA [150mM NaCl/50mM Tris–HCl (pH 7.5)/1mM EDTA/0.1% (v/v) NP-40/0.5mM PMSF/0.25% (w/v) BSA]. Precipitated proteins were eluted by boiling in sample buffer for 10min, separated on a 10% (v/v) SDS–PAGE and immunodetected as described above.

Nuclear matrix preparations

High salt isolation of nuclear matrix was carried out essentially as previously reported, with some modifications [22,23]. Nuclei, isolated by centrifugation in a 2-M sucrose cushion, as described above, were stripped of membranes by resuspension in 150µl of Triton X-100 extraction buffer [0.5% (v/v) Triton X-100/10mM Tris–HCl (pH 7.4)/2.5mM MgCl2], followed by incubation on ice for 15min. Membrane-depleted nuclei were collected by centrifugation at 5000 × g for 10min at 4°C. Chromatin and soluble proteins were removed by suspension in 300µl of DNase I digestion buffer [10mM Tris–HCl (pH 7.4)/ 0.1mM DTT/10mM MgCl2] containing 200U/ml RNase-free DNase I and incubated 2h at

37°C, addition of 100µl stop buffer [10mM Tris–HCl(pH7.5)/200mMEDTA(pH 8.0)] and400µlof2× high salt buffer [20mM Tris–HCl (pH 7.4)/1.3M (NH4)2SO4] and further 5min incubation at 4°C. Sample was centrifuged at 10,000 × g for 10min at 4°C and the nuclear matrix pellet was resuspended in 150µl of RIPA lysis buffer [50mM Tris–HCl (pH 7.4)/5mM DTT/150mM NaCl/1% (v/v) NP-40/0.5% (w/v) sodium deoxycholate/0.1% (w/v) SDS/0.5mM PMSF]. Nuclear matrix proteins were separated on a 10% (v/v) SDS–PAGE and immunodetected as described above.

In situ nuclear matrix was prepared essentially as previously reported [24,25]. Briefly, HeLa cells grown on coverslips were washed twice with ice-cold PBS followed by incubation with CSK buffer [10mM Pipes (pH 6.8)/100mM NaCl/30mM sucrose/3mM MgCl2/1mM EGTA/0.5% (v/v) Triton X-100], supplemented with complete protease inhibitor mixture to remove cytosolic soluble proteins. After 10min at 4°C, the chromatin was eliminated by incubating with DNase I digestion buffer [CSK buffer containing 200U/ml of RNase-free DNase I] for 2h at 37°C. Next, to remove the remaining DNA and associated proteins, the nuclei were incubated with extraction buffer [CSK buffer containing 650mM (NH4)2SO4] for 20min at 4°C. Finally, samples were washed with the CSK buffer and then, fixed with CSK and 4% (w/v) paraformaldehyde for 40min in CSK. No detectable staining of nuclei with propidium iodide was observed, demonstrating effective removal of DNA by this procedure. the Immunofluorescence analysis on in situ nuclear matrix was performed as described above for whole cells.

Results

Expression and subcellular localization of endogenous Dp71 protein isoforms in HeLa cells

Using GFP-Dp71 gene fusions, we have previously demonstrated that alternative splicing determines the nuclear or cytoplasmic localization of exogenously expressed Dp71 protein isoforms in HeLa cells [21]. Hence, to corroborate these results, we decided to analyze the expression and distribution patterns of endogenous Dp71 protein isoforms in this cell line. HeLa cells are commonly used to study the function of nuclear proteins. To differentiate between Dp71 protein variants, we utilized different antibodies raised against the Dp71 C-terminal domain (Table 2): Dys2 and 2166, which recognize specifically the protein isoform containing the domain encoded by exon 78 (Dp71d), and 5F3, which detects exclusively the protein variant bearing 31 new amino acids that are incorporated into the molecule due to the alternative splicing of exon 78 (Dp71f). A typical immunoblot for the detection of Dp71 isoforms is shown in Fig. 1A; a band with an apparent molecular mass of 71kDa was observed when total protein extracts from HeLa cells were probed with either Dys2 or 5F3 antibodies. These results indicate that both Dp71d and Dp71f are expressed in HeLa cells. We did not observed any immunoreactive band above the 70-kDa protein marker, which indicates that besides Dp71, HeLa cells do not express any other DMD gene product at detectable levels.

Next, to study the subcellular distribution of each Dp71 variant, immunofluorescence and confocal microscopy analyses were carried out.

Immunostaining of Dp71f revealed a scattered signal distributed throughout the cytoplasm and excluded from nuclei. In contrast, although Dp71d staining was also present in the cytoplasm, it accumulated mainly in the nuclei, leaving the nucleoli clear of signal (Fig. 1B). To confirm the distinctive subcellular distribution of each Dp71 isoform, we carried out immunoblotting analyses of nuclear and cytosolic protein extracts. To verify the fidelity of our fractionation procedure, we employed vinculin, a focal adhesion protein [26], and calnexin, a membrane protein from the endoplasmic reticulum, as cytoplasmic markers. On



Fig. 1 – Expression and subcellular localization of endogenous Dp71 protein isoforms in HeLa cells. (A) Total protein extracts from HeLa cells were separated on 10% (v/v) SDS–PAGE, transferred to nitrocellulose membranes and probed with 5F3 and Dys2 monoclonal antibodies for the detection of Dp71f and Dp71d isoforms, respectively. Migration of Dp71 protein isoforms is indicated on the right by an arrow. Positions of molecular weight standards are shown on the left. (B) Cells, cultured on glass coverslips for confocal microscopy analysis, were immunostained with the primary monoclonal antibodies described above, and the specific signal was developed using a fluorescein-conjugated secondary antibody. Representative optical Z sections were selected to show the subcellular distribution of Dp71 isoforms. Scale bar = 16 µm. (C) Cellular fractionation was performed as described in Materials and methods. Nuclear (n) and cytosolic (c) protein extracts were subjected to immunoblot analysis using vinculin and calnexin antibodies as cytoplasmic markers and lamin B1 antibody as nuclear marker. (D) Subcellular fractions were analyzed with 5F3 and 2166 antibodies for the detection of Dp71f and Dp71d protein isoforms, respectively. Dp71f and Dp71f protein isoforms are expressed in HeLa cells; Dp71f distributes

the other hand, lamin B1, a nuclear matrix-associated protein [27], was used as nuclear marker. As expected, the vinculin (135kDa) and calnexin (90kDa) bands were found exclusively in the cytosolic fraction whereas the lamin B1 band (68kDa) was absent in the cytosolic fraction but present in the nuclear extract (Fig. 1C). These results indicate that nuclear preparations were free of cytosolic proteins and membrane proteins from cytoplasm. Regarding the distribution of Dp71 isoforms, the Dp71f immunoreactive band was undetectable in the nuclear fraction but prominent in the cytosolic preparation; in contrast, the Dp71d immunoreactive band, clearly observed in the nuclear extract, was absent in the cytosolic fraction (Fig. 1D).

Expression of DAPC members in HeLa cells

To examine the expression profile of the dystrophin-associated protein complex (DAPC) components in HeLa cells, we performed RT-PCR and immunoblot analyses. We designed different pairs of primers to amplify specifically the cDNA of each DAPC member. Positive controls were carried out on human muscle total RNA or rat brain total RNA and the identity of the RT-PCR products was determined by DNA sequencing (data not shown). In summary, we determined β -sarcoglycan, α -syntrophin, that β -dystroglycan, β 2-syntrophin, **α**1dystrobrevin, β-dystrobrevin and nNOS are expressed at mRNA level in HeLa cells (Table 1). On the basis of these results, we further investigated the presence of the DAPC proteins in total extracts of HeLa cells using antibodies directed specifically to each member (Table 2). Representative immunoblots of DAPC members are shown in Fig. 2 (see lanes corresponding to total extracts). By using a monoclonal antibody anti-β-dystroglycan (NCL-b-DG), a robust band migrating at 43kDa was observed, which corresponds to the expected molecular weight of β-dystroglycan; in addition, a faint band migrating between the 43-and 59-kDa markers was observed. The expected band for β -sarcoglycan positioned above the 43-kDa protein marker was obtained after using a monoclonal antibody anti-βsarcoglycan (NCL-b-SARC). By using the pan-specific antibody α-CTFP, two bands with apparent molecular weights of 70 and 94kDa were revealed, which must correspond to β -and α 1-dystrobrevin, respectively. A band of approximately 60kDa, which must correspond to α -and/or β -syntrophin, was revealed after using a pan-specific antibody for α -and β -syntrophins. Finally, a band of 150kDa that corresponds to nNOS expression was obtained after using a commercial antinNOS antibody.

Subcellular distribution of DAPC components in HeLa cells

To examine the subcellular localization of DAPC members, we subjected HeLa cells, grown on coverslips, to immunofluorescence and confocal microscopy analysis using the appropriate fluorescein-conjugated secondary antibodies. As

shown in Fig. 3, β -dystroglycan displayed a predominant cytoplasmic staining with a perinuclear distribution that could correspond to endoplasmic reticulum or nuclear membrane labeling; in addition, we observed a scarce diffuse punctuate labeling inside nuclei (viewed specifically with propidium iodide), which was excluded from nucleoli. On the other hand, α 1dystrobrevin and β -sarcoglycan immunostaining spanned the whole cell body but accumulated mainly in the nuclei (Fig. 3). Finally, as indicated by the phalloidin labeling of cytoplasmic actin, the immunostaining of β -dystrobrevin, nNOS and α/β syntrophin appeared to be mostly nuclear but, excluded from nucleoli. To distinguish between α -and β -syntrophin, we employed specific antibodies for α -or β 2-syntrophin (Table 2) in the immunofluorescence assays. We observed that β 2syntrophin displayed a predominant nuclear localization whereas α -syntrophin accumulated evenly between cytoplasm and nucleus (data not shown).

To support the immunofluorescence results with biochemical evidence, we carried out subcellular fractionation of HeLa cells into cytoplasmic and nuclear extracts and we subjected



Fig. 2 – Subcellular distribution of DAPC components in HeLa cells. Total (t), nuclear (n) and cytoplasmic (c) extracts, obtained as indicated in Materials and methods, were separated on 10% (v/v) SDS–PAGE, transferred to nitrocellulose membranes and probed with antibodies directed specifically against β -dystroglycan (NCL-b-DG), β -sarcoglycan (NCL-b-SARC), α 1-dystrobrevin and β -dystrobrevin (β -CTFP), α/β -syntrophin, nNOS and actin (Table 2). For comparison, equal amounts of cytoplasmic and nuclear proteins were loaded in the corresponding lanes. Immunodetected DAPC members are indicated at the top of the panels. Position of protein markers is shown at the right. Arrows at the left denote the migration of the expected protein bands. All of the DAPC components analyzed by cellular fractionation were recovered in the nuclear fraction.



Fig. 3 – Immunofluorescence analysis of DAPC components in HeLa cells. Cells immunostained with the following antibodies: anti- β -dystroglycan (JAF), anti- α 1-dystrobrevin (α -CTFP) and anti- β -sarcoglycan (NCL-b-SARC) (green color) were counterstained with propidium iodide (PI, red color) to visualize nuclei. Cells immunolabeled with anti- β -dystrobrevin antibody (β -521), anti-nNOS and anti- α/β -syntrophin (Pan-syntrophins) (green color) were counterstained with Texas red phalloidin (red color) to visualize cytoplasmic actin. After double staining, cell preparations were subjected to confocal microscopy analysis and single optical Z sections were selected in each case to show the distribution of DAPC components. The perinuclear localization of β -dystroglycan is denoted by arrows. All merged images are shown in the right panels. DAPC members displayed different nuclear distribution patterns. Scale bar = 20 µm.

the resulting protein extracts to SDS–PAGE and immunoblotting analyses. The β dystroglycan immunoreactive band (43kDa) was prominent in the cytosolic fraction but was practically absent in the nuclear extract; in contrast, a second slower migrating band was observed predominantly in the nuclear fraction showing a weak signal in the cytosolic extract (Fig. 2). The same pattern of immunoreactive bands for β -dystroglycan was obtained with two different antibodies (JAF and NCL-b-DG). It has been observed that phosphorylation of β dystroglycan causes an upward mobility shift in HeLa cells [28]; therefore, the retarded band might correspond to a phosphorylated isoform of this protein that accumulates in the nucleus of HeLa cells. The β -sarcoglycan immunoreactive band seemed to accumulate in the cytosolic extract showing a comparative weaker intensity in the nuclei. In contrast, the nNOS protein signal was observed mostly in the nuclear fraction (Fig. 2). In regard to the distribution of dystrobrevins, it was observed that both β -dystrobrevin and α 1-dystrobrevin protein levels are enriched in the nuclear preparation being higher the cytoplasmic protein levels of the latter protein (Fig. 2). Finally, the immunoreactive band corresponding to α/β -syntrophin was distributed evenly between the two cellular compartments (Fig. 2). Based on the immunofluorescence data obtained with antibodies directed specifically against α -syntrophin or β -syntrophin, it could be proposed that the immunoreactive band revealed in the nucleus with the pansyntrophin antibody corresponds mainly to β -syntrophin. In addition, because actin represents a nuclear/cytoplasmic protein that has been found associated with Dp71 in cultured cells [29], we decided to analyze its subcellular distribution as well. We observed that cytoplasmic protein levels of actin were similar to those found in the nuclear fraction (Fig. 2).

Overall, the immunofluorescence and cellular fractionation studies evidenced the presence of DAPC components in the HeLa cell nuclei; however, certain differences were observed in the nuclear abundance of the DAPC members by comparing these two methodologies: For instance, immunostaining of α/β -syntrophin appeared to be mainly nuclear but the fractionation analysis indicated that protein levels of α/β syntrophin are portioned similarly between cytoplasm and nucleus. These inconsistencies could be due to the principle of each methodology; in the confocal images, mid-optical Z-sections were selected to emphasize the nuclear presence of the DAPC members whereas in the



Fig. 4 – Dp71 forms a DAPC in the HeLa nuclei. Detergent-soluble nuclear extracts from HeLa cells were immunoprecipitated with antibodies specific to Dp71 (mDp71), β -dystrobrevin (β -521), α 1-dystrobrevin (α -CTFP) and β -dystroglycan (JAF). Co-immunoprecipitated proteins were analyzed by immunoblotting with antibodies directed against Dp71 (mDp71), β -dystroglycan (NCL-b-DG), nNOS and α 1-and β -dystrobrevin (β -CTFP). As negative control, an antibody directed against the transcription factor Sp3 was employed. Representative results from three independent immunoprecipitation assays are shown. β -DG, β -dystroglycan; α 1-DB, α 1-dystrobrevin; β -DB, β -dystrobrevin; IP, immunoprecipitation; WB, western blotting. Input, soluble nuclear extracts subjected to immunoblotting. The unspecific binding lane refers to immunoprecipitation experiments where the respective primary antibody was omitted. Dp71, β -dystroglycan, nNOS, α 1-dystrobrevin and β -dystrobrevin, copurified in the nuclear fraction.

fractionation studies a more accurate determination of the cytoplasm-nucleus ratio of these proteins was provided.

Dystrophin Dp71 forms a DAPC in the nuclei of HeLa cells

To explore the possibility that Dp71 interacts with DAPC components within HeLa nuclei to form a DAPC complex in this organelle, we immunoprecipitated nuclear protein extracts with an anti-Dp71 antibody raised against the N-terminus domain of Dp71 (mDp71) and analyzed by Western blotting with a panel of antibodies directed against different DAPC components. It was observed that βdystroglycan, nNOS, α 1-dystrobrevin and β -dystrobrevin immunoprecipitated with Dp71. In reciprocal experiments, Dp71 was immunoprecipitated by antibodies directed against β -dystroglycan (JAF), α 1-dystrobrevin (α -CTFP) and β -dystrobrevin (β -521), confirming the association of Dp71 with these proteins (Fig. 4). Moreover, Dp71, nNOS, α 1-dystrobrevin and β -dystrobrevin were found in the immunoprecipitates of β -dystroglycan; reciprocal experiments showed that β -dystroglycan immunoprecipitated with α 1-and β -dystrobrevin (Fig. 4). In addition to Dp71 and β -dystroglycan, immunoprecipitates of α 1-and β dystrobrevin contained nNOS. Interestingly, α 1-and β -dystrobrevin copurified, which suggest that they are part of the same protein complex within HeLa nuclei. We were unable to detect β -sarcoglycan in the Dp71, β -dystroglycan and α 1-and Bdystrobrevin immunoprecipitates (data not shown). Specificity of the

immunoprecipitation assays was established by the following negative controls: (a) Assays where the immunoprecipitating antibodies were omitted showed that none of the proteins examined precipitated in a unspecific way (Fig. 4, lane "unspecific binding"); and (b) Western blotting experiments with an antibody directed against the transcription factor Sp3 showed that the DAPC immunoprecipitates excluded an unrelated protein (Fig. 4). Overall, our data indicate that Dp71 forms a DAPC in the nuclei of HeLa cells, which contains, at least, β -dystroglycan, α 1-/ β -dystrobrevin and nNOS.

Components of the nuclear DAPC associate with the nuclear matrix of HeLa cells

Because nuclear matrix has been proposed to be the underlying structural support for subnuclear organization of diverse protein complexes [30,31], we examined whether or not the members of the nuclear Dp71-containing DAPC complex bind to the HeLa nuclear matrix. We obtained nuclear and nuclear matrix protein extracts and in situ nuclear matrix preparations by a sequential high salt extraction procedure previously described [22–25]. To verify the authenticity of the nuclear matrix preparations, we analyzed the distribution of the nuclear matrix proteins lamin B1 and actin as well as the distribution of the soluble transcription factor Sp3. Fig. 5A shows optical sections of nuclear matrix preparations stained with antibodies directed against Dp71 or the different DAPC components. We obtained the typical staining of lamin B1 localized at the nuclear periphery with a dotted organization in the inner nuclear matrix (Fig. 5A). As expected for a soluble nuclear protein, the Sp3 immunolabeling was excluded from the nuclear matrix preparation (Fig. 5A). Interestingly, the immunostaining of Dp71 as well as β dystroglycan, β dystrobrevin, α 1-dystrobrevin and nNOS was retained within the nucleoskeletal structure displaying specific labeling patterns. The staining of Dp71 and β -dystroglycan distributed mainly in the nuclear matrix periphery forming a ringlike arrangement; besides, the signal of the latter protein produced some diffuse labeling in the inner nucleoskeletal structure that may correspond to invaginations of the nuclear envelope in the nucleus (Fig. 5A). The α 1-dystrobrevin immunostaining displayed a punctate labeling throughout the nuclear matrix structure that left nucleoli clear of signal. On the other hand, β-dystrobrevin labeling distributed through the nucleoskeletal structure showing patches with labeling. Finally, nNOS immunodetection more intensive displayed а discontinuous staining throughout the nucleoskeletal structure that forms a netlike arrangement (Fig. 5A). It should be noted that fluorescence pattern of Dp71, Bdvstroglycan and lamin B1 colocalized at certain extent in the nuclear matrix periphery, which may indicate their physical interaction.



Fig. 5 – Association of Dp71 and DAPC components with the nuclear matrix of HeLa cells. (A) In situ nuclear matrix preparations were obtained as described in Materials and methods. Immunofluorescence confocal microscopy images of representative nuclear matrices stained with antibodies directed against Sp3, lamin B1, Dp71d (2166), β -dystroglycan (JAF), β -dystrobrevin (β -521), α 1-dystrobrevin (α -CTFP) and nNOS are shown. As control, a phase contrast micrograph of the in situ nuclear matrix preparation used for Sp3 immunostaining is also shown. Immunostaining of Dp71, β -dystroglycan, β -dystrobrevin, α 1-dystrobrevin and nNOS was retained within the nuclear matrix preparation whereas that of Sp3 was not. Immunostaining of lamin B1, Dp71d and β -dystroglycan in the nuclear matrix periphery is denoted by arrows. (B) Equal amounts of nuclear (N) and nuclear matrix (NM) protein extracts, obtained as described in Materials and methods, were subjected to immunoblotting analysis. Proteins were visualized by using specific antibodies against Sp3 (negative control of the nuclear matrix

extracts), lamin B1, actin, Dp71 (mDp71), β -dystroglycan (NCL-b-DG), α 1-and β -dystrobrevin (β -CTFP), nNOS, β -sarcoglycan (NCL-b-SARC) and α/β -syntrophin. Representative results from three independent assays are shown. The position of the protein markers is shown at the left. Arrows point to the specific bands of Sp3. The slower-migrating band of β -dystroglycan is denoted by an arrow. The expected protein bands for lamin B1, actin, Dp71, β -dystroglycan, α 1-and β -dystrobrevin, nNOS, β -sarcoglycan and α/β -syntrophin were recovered in the nuclear matrix fraction.

To obtain protein extracts from the nuclear matrices of HeLa cells, previously isolated nuclei were digested with DNase I and extracted with 1.3M (NH4)2SO4, and the nuclear material remaining after treatment was electrophoresed and immunoblotted with antibodies directed against Dp71 or to the different DAPC components. As expected, the immunoreactive bands for lamin B1 (68kDa) and actin (43kDa) were present in both the nuclear and the nuclear matrix fractions whereas those of Sp3 protein (115, 80 and 70kDa) were present in the nuclear extract but excluded from the nuclear matrix fraction (Fig. 5B). Fig. 5B also shows that the Dp71, β -dystroglycan, α 1-and β dystrobrevin, nNOS, β sarcoglycan and α/β -syntrophin immunoreactive bands were recovered in both the nuclear and nuclear matrix fractions after using the antibodies mDp71, NCLb-DG, β-CTFP, anti-nNOS, NCL-b-SARC and pan-syntrophins, respectively. The additional slower-migrating band of β -dystroglycan, noted in the nuclear fraction, was also revealed in the nuclear matrix extract (indicated by black arrow, Fig. 5B). It was also observed that only α 1-dystrobrevin displayed enrichment in the nuclear matrix fraction, compared with the nuclear extract (Fig. 5B). Altogether, confocal microscopy and immunoblotting analyses of HeLa nuclear matrix preparations evidenced that some members of the nuclear DAPC are nuclear matrix-associated proteins.



Fig. 6 – Interaction of Dp71 and DAPC members with nuclear matrix proteins. Detergent-soluble nuclear extracts from HeLa cells were immunoprecipitated with antibodies specific to Dp71 (mDp71), β -dystroglycan (JAF), α 1-dystrobrevin (α -CTFP) and β -dystrobrevin (β -521). Co-immunoprecipitated proteins were analyzed by Western blotting with antibodies directed against actin and lamin B1. Representative results from three independent immunoprecipitation assays are shown. IP, immunoprecipitation. WB, Western blotting. Input, soluble nuclear extracts subjected to immunoblotting analysis. The unspecific binding lane refers to immunoprecipitation experiments where the respective primary antibody was omitted. β -DG, β -dystroglycan; α 1-DB, α 1-dystrobrevin; β -DB, β -dystrobrevin.

Interaction of nuclear DAPC components with the nuclear matrix proteins actin and lamin B1

To verify the experimental evidence concerning the association of nuclear DAPC components with the nuclear matrix, we examined whether any of the DAPC members establishes physical interactions with the nuclear matrix proteins lamin B1 and actin. Therefore, nuclear protein extracts prepared from HeLa cells were immunoprecipitated with specific antibodies for Dp71, α 1-dystrobrevin, β dystrobrevin or ßdystroglycan and the resulting immunoprecipitates were immunoblotted with antibodies directed against lamin B1 and actin. Fig. 6 shows that the amount of actin and lamin B1 recovered in the immunoprecipitates of Dp71, β -dystroglycan and β -dystrobrevin was clearly higher than that obtained in the negative control experiment. In contrast, the intensity of the actin and lamin B1 immunoreactive bands was undistinguishable between the α 1-dystrobrevin immunoprecipitate and the negative control experiment. Western blotting assays using an anti-Sp3 antibody demonstrated that immunoprecipitates of DAPC members excluded an unrelated protein (see lower panel of Fig. 4). Altogether, these results indicate that actin and lamin B1 copurified specifically with Dp71, β -dystroglycan and β -dystrobrevin, but not with α 1-dystrobrevin.

Discussion

The characterization of different dystrophin-associated protein complexes (DAPCs), such as those localized in the skeletal muscle, neuromuscular junction, brain, lung, retina and kidney, has been reported [3,11–13,17,32]. In addition, an increasing number of cellular functions associated with these multiple protein complexes have been proposed, such as membrane stability, cellular signaling and force transduction [7,33,34]. The specific function of each DAPC seems to be determined by the cellular environment where it resides as well as by the unique combination of proteins that conforms the complex.

In this work, we report for the first time the existence of a DAPC complex in a cellular compartment different from cell membrane, the nucleus. We identified and characterized a DAPC present in the nuclei of HeLa cells, which contains dystrophin Dp71 as a key component. Besides, Dp71, the complex is composed, at least, of β -dystroglycan, α 1-and β dystrobrevin and nNOS. Previously, the presence of Dp71 and several DAPC components in the nuclei of different cell types has been described [17,19,20]. The nuclear distribution of Dp71 has been observed in cultured hippocampal neurons

[35] and the PC12 cell line [36,37]. Additionally, α -dystrobrevin was found in the nuclei of the human cell line HL-60

[19] whereas β -dystrobrevin was observed in the nuclear compartment of

hippocampal pyramidal neurons [17]. Furthermore, the nuclear localization of α -, β -and γ 1syntrophin has been described in HeLa cells [20]. Finally, the nuclear localization of nNOS in the rat brain and MDCK cells has been also reported [38]; the authors proposed that nNOS may traffic nuclear proteins to the cytosol via its PDZ domain.

Although the evidence mentioned above point to the existence of a nuclear DAPC complex, to our knowledge, this is the first report establishing the formation of a Dp71contaning DAPC complex within nuclei. Our statements are based on the following experimental evidence: We have demonstrated by immunofluorescence and cell fractionation experiments that Dp71, ßdystroglycan, nNOS, β -sarcoglycan, α/β -syntrophin and α 1-and β -dystrobrevin colocalize in the HeLa cell nuclei. In addition, we established by co-immunoprecipitation assays that Dp71, β -dystroglycan, nNOS, α 1 and β -dystrobrevin associate by themselves to form a protein complex within this organelle. Because two different dystrobrevins, $\alpha 1$ and β , colocated in the HeLa nuclei, it would be expected that at least two distinct Dp71-containing complexes are formed in this cellular compartment, each one including a specific dystrobrevin. However, we found that α 1 and β -dystrobrevin copurified together in the HeLa nuclear fraction suggesting that they are part of a single protein complex. This finding contrasts to the previously described cell membrane-associated complexes where a single dystrobrevin is included in each DAPC [11]. Further biochemical experiments are clearly needed to define the precise arrangement of dystrobrevins in the nuclear DAPC. We were unable to show any association of β -sarcoglycan with the nuclear DAPC; nevertheless, because this protein is distributed in the nuclear compartment of HeLa cells, the possibility that β -sarcoglycan binds to this novel protein complex still open.

The existence of a nuclear DAPC raises important questions about the molecular mechanism governing its nuclear transport, either as a protein complex or as independent components. Selective nuclear import occurs through the nuclear pore complex [39], which allows small molecules (under 45kDa) to diffuse freely into and out of the nucleus [39]. The targeting of larger proteins requires the presence of either a classical or a bipartite nuclear localization signal (NLS), both of which are composed of basic amino acids [40,41]. In this regard, a computational analysis of the entire amino acid sequence of each DAPC component showed that only the β dystroglycan contains a stretch of basic amino acids (RKKRKGK) that could correspond to an NLS (amino acids 776-782). Therefore, it is tempting to speculate that DAPC components gain access to the soluble transport machinery through its association with β -dystroglycan. Alternatively, it couldbepossiblethatDp71orany otherofthe DAPC components contains an atypical but functional NLS motif [42]. It has been recently reported that α -, β -and γ 1syntrophin are recruited into HeLa cell nuclei via their interaction with the enzyme diacylglycerol kinase- ζ [20]. Hence, it seems that several alternative molecular mechanisms could be involved in the transport of DAPC components to HeLa nuclei.

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The existence of a nuclear DAPC elicits interesting questions regarding its physiological function within this organelle; to start exploring this issue, we determined whether any of the nuclear DAPC members associates with the nuclear matrix. It is now becoming increasingly clear that the nuclear matrix provides the underlying support for nuclear compartmentalization; hence, nuclear factors localize to discrete subdomains within the nucleus to perform specialized functions, such as DNA organization and replication, gene transcription and RNA processing [43]. Rather than behave as soluble, easily extractable factors, we were surprised to find that Dp71 and various DAPC components, such as β dystroglycan, β -sarcoglycan, β dystrobrevin, α 1-dystrobrevin, nNOS and α/β syntrophins, were resistant to nuclei extraction with detergent, nucleases and salt. Such treatments are normally used to extract chromatin components including transcription factors or histones. Thus, we have demonstrated that members of the Dp71-containing DAPC are not only localized in the nucleus; but also tightly associated with the nuclear matrix. Interestingly, the immunofluorescencelabeling pattern of Dp71 and β -dystroglycan colocalized at certain extent with that of lamin B1, especially at the nuclear matrix periphery. These results indicate that these three proteins might colocalize spatially. Supporting these data, we found that Dp71, β dystroglycan and β -dystrobrevin co-immunoprecipitated with the nuclear matrix proteins lamin B1 and actin. Because lamin B1 is a structural component of the nuclear matrix that facilitates the interaction between chromatin and the inner nuclear membrane [27] and actin is a component of several mammal chromatin-remodeling complexes [44], the association of Dp71 and some DAPC members with those proteins might indicate that they work as scaffolding proteins involved in nuclear architecture. Additional biochemical studies are necessary to establish whether the interaction between DAPC components and actin and lamin B1 is direct or due to an association within the context of a larger nuclear matrix complex.

Based on the results of this work and on the well-known protein-protein interactions occurring among DAPC members at the cell membrane, it is tempting to speculate that the arrangement of the nuclear DAPC is similar to that displayed by the DAPC at the cell membrane. We hypothesize that β dystroglycan and β sarcoglycan are inserted in the inner nuclear membrane via their respective transmembrane domains. Therefore, the C-terminal domain of β-dystroglycan would interact with the WW domain of Dp71, which in turns might bind to β dystrobrevin by their coiled-coil domains in the C-terminal region of each protein. Likewise, we propose that α 1-and β -dystrobrevin bind each other via a coiled-coil interaction forming a dystrobrevin heterodimer. On the other hand, nNOS may be recruited to the nuclear DAPC by interacting with α/β -syntrophin [20] or with a PDZ-containing nuclear protein. We propose that the physical contact between the DAPC and the nuclear matrix might be mediated by the interaction of β dystroglycan with lamin B1. In addition, Dp71 might contact nuclear actin via its N-terminal actin-binding domain [29].

In summary, we have identified and characterized a novel Dp71-containing

DAPC present in the nuclei of HeLa cells. In addition, we have revealed that several members of this novel complex are associated with the nuclear matrix. Altogether, our results suggest that the nuclear DAPC may work as a nuclear scaffolding complex involved in nuclear organization.

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