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Differential expression of β 1, β 3 and β 4 integrins in sarcomas of the small, round, blue cell category

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Abstract Integrins are a large and complex family of membrane spanning αβ heterodimeric cell surface glycoproteins mediating cell/cell and cell/matrix interactions. Small, round, blue cell sarcomas (SRBCS) are a group of poorly differentiated tumours of various and in part uncertain histogenesis displaying similar cytomorphology. Among them are rhabdomyosarcomas (RMS), ganglioneuroblastomas [(G)NB], primitive peripheral neuroectodermal tumours (pPNET) and Ewing's sarcomas (ES). Thirty-two SRBCS were studied immunohistochemically for the distribution of β 1, β 3 and β 4 integrins in situ. We found complex and to some extent differential patterns of β1, β3 and β4 integrin subunit expression in different types of SRBCS: all of the sarcomas studied were consistently $\beta 1^+$, $\beta 4^-$, $\alpha 2^-$. Four of nine RMS were completely negative for all other integrin subunits studied while one RMS was α5+ throughout and three RMS were focally $\alpha 5^+$. Three RMS expressed the $\alpha 6$ and αv chains. In contrast to RMS, pPNET and ES, all of which were $\alpha 1^-$, $\alpha 3^-$, (G)NB were $\alpha 3^+$ and frequently co-expressed a1. The eight pPNET and seven ES studied showed a similarily restricted integrin profile that was limited to the expression of $\beta 1$ and $\alpha 5$ in nearly all cases. In summary, RMS were $\beta1^+$, $\alpha1^-$, $\alpha3^-$ and heterogeneously expressed $\alpha 5$ and $\alpha 6$. (G)NB were generally β 1+, α 1+, α 3+, α 5-, α 6-. pPNET and ES were β 1+, α 1-, $\alpha 3^-$, $\alpha 5^+$, $\alpha 6^-$. The data illustrate a complex expression pattern of various integrins in SRBCS, a differential expression pattern of some of the integrin subunits among different types of SRBCS and almost identical integrin profiles in pPNET and ES.

Key words Small round blue cell sarcomas · Integrins · Immunohistochemistry

This paper is dedicated to Prof. Dr. Dres. h.c. Wilhelm Doerr on the occasion of his 80^{th} birthday

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Introduction

Interactions between cells and extracellular matrix components are mediated, in part, via a large family of cell surface adhesion molecules known as integrins (recently reviewed in [23, 30]). Integrins are non-covalently linked membrane spanning αβ heterodimeric glycoproteins. Through an extracellular receptor domain which is built up by both subunits and the cytoplasmic portion of the β subunit which interacts with talin and α -actinin, integrins connect the extracellular milieu with the cytoskeleton [8, 27]. Currently, at least 15 different α chains and eight different β chains are known [22, 28, 30]. The integrin family is actually subdivided into different subgroups based on types of β chains which, in turn, can dimerize with different α chains. The largest subgroup are the β 1 integrins, formerly designated very late antigens since they were first described on T cells during late phases of activation in vitro [18]. The \(\beta 1 \) integrins consist of the common \(\beta \) chain [CD29 according to the cluster of differentiation (CD) nomenclature of cell surface molecules] that combines with at least six α chains (CD49a-f) to form six different αβ heterodimeric cell surface receptors. Furthermore, the \$1 chain has been shown to associate alternatively with the av chain (CD51) of the vitronectin receptor αvβ3 (CD51/CD61) [39]. Another example of the promiscuity within the different integrin subunits is the α6 chain which can also dimerize with the \(\beta \) chain (CD104) [20]. Receptors of the β 1, β 3 and β 4 integrin type were shown to bind to a variety of extracellular matrix proteins and to some cell adhesion molecules (Table 1). As cell adhesion molecules they are involved in different biological events such as embryogenesis, wound healing and immune response. Integrins are differentially expressed on a broad spectrum of normal cell types and their neoplastic equivalents [1, 2, 7, 24, 29, 45]. The descriptive term , small, round, blue cell sarcoma' (SRBCS) encompasses different tumours of neuroectodermal and mesenchymal derivation that, although similar in cytomorphology, have different biological properties [10]. Among SRBCS are

Table 1 Integrin subunits detected and monoclonal antibodies used in this study (CD cluster of differentiation, VCAM vascular cell adhesion molecule, ICAM intercellular adhesion molecule)

Antigen	Molecular weight (kDa)	CD number ^a	Clone	Isotype	Receptor for function	Reference
β1	130	CD29	K20	IgG2a	Common β chain of $\alpha 1$ to $\alpha 6$	[4]
β1 β3	110	CD61	SZ.21	IgG1	Common β chain of αv and αIIb (CD41)	[31]
β4	205	CD104	3E1	IgG1	Alternative β chain of α 6/laminin	[21]
ά1	210	CD49a	TS2/7	IgG1	Collagen/laminin	[19]
α 2	170	CD49b	CLB-	IgG1	Collagen/(laminin)/(fibronectin)	[16]
			thromb/	4		_
$\alpha 3$	130	CD49c	J143	?	Collagen/laminin/fibronectin/epiligrin/entactin	[14]
α4	150	CD49d	HP2/1	IgG1	Fibronectin/VCAM-1(CD106)/ICAM-2(CD102)	[32]
α5	135	CD49e	SAM1	IgG2b	Fibronectin	[37]
α6	120	CD49f	GOH3	IgG2a	Laminin	[35]
αν	120	CD51	AMF7	IgG1	Vitronectin/fibronectin/fibrinogen/von Willebrand's factor/collagen/osteopontin/thrombospondin	[38]

^a According to the fixing of the Nomenclature Committee of the 5th International Workshop and Conference on Leukocyte Differentiation Antigens, Boston, November 1993 [33]

rhabdomyosarcomas (RMS), ganglioneuroblastomas (GNB), neuroblastomas (NB), primitive peripheral neuroectodermal tumours (pPNET) and Ewing's sarcomas (ES). Histogenetic assignment of pPNET and ES is a matter of ongoing controversy [3, 5, 36, 41].

Most data on integrin expression in SRBCS are based on in vitro studies of various cell lines, such as NB, ES and RMS cell lines [6, 9, 11, 43]. The analysis in situ is still very incomplete. We therefore analysed the distribution pattern of β 1, β 3 and β 4 integrin subunits immunohistochemically in a series of 32 unselected SRBCS.

Materials and methods

Fresh frozen tumour tissue from 32 SRBCS was collected at our institute in a period of 5 years. The tumour series comprised nine RMS, five GNB, three NB, eight pPNET and seven ES. The diagnosis of each tumour case was based on routine paraffin sections (haematoxylin and eosin, periodic acid-Schiff, Masson-Goldner and Gomori silver stains) according to standard histopathological criteria as described by Enzinger and Weiss [12], combined with immunohistochemical analysis using monoclonal antibodies and antisera against structural antigens of mesenchymal cells (for review see [12, 26, 42]). All tissue samples were taken from incisional biopsies or surgical specimens. The tissue was snap-frozen in liquid nitrogen within 1 h after removal and stored at –70° C. Serial frozen sections of about 1 cm² and 4–6 µm thickness were airdried, acetone-fixed at room temperature for 10 min and immunostained immediately or stored at –20° C for a short period.

The primary monoclonal antibodies (mAb) used in this study are listed in Table 1. The mAb K20 (anti-β1), Gi9 (anti-α2), HP2/1 (anti-α4), SAM1 (anti-α5), GOH3 (anti-α6), SZ.21 (anti-β3) and AMF7 (anti-αν) were obtained from Dianova (Hamburg, Germany); mAb TS2/7 (anti-α1) was purchased from T Cell Sciences (Cambridge, Mass., USA); mAb P1B5 (anti-α3) and 3E1 (anti-β4) were obtained from Telios Pharmaceuticals (San Diego, Calif., USA). A polyclonal biotinylated sheep antibody to mouse immunoglobulin (Ig; reactive with all mouse isotypes), a polyclonal biotinylated sheep antibody to rat Ig for detection of rat-derived mAb GOH3, and a streptavidin-biotinylated peroxidase complex, all obtained from Amersham (Buckinghamshire, UK), served as a detection system for the primary antibodies. 3-Amino-9-ethyl-carbazole (AEC) and N'N-dimethylformamide were obtained from Sigma Chemical Company (St. Louis, Mo., USA).

After rehydration with phosphate-buffered saline solution (PBS; pH 7.4), the frozen sections were incubated for 1 h with primary mAb. The mAb as culture supernatants were applied undilut-

ed, ascites preparations were diluted 1:2000 in PBS, purified reagents were used in a protein concentration of about 10 µg/ml PBS. All incubation steps were carried out in a humid chamber at room temperature and were followed by double rinsing with PBS. AEC was used as chromogen [0.4 mg/ml in 0.1 mol/l (molar) acetate buffer, pH 5.0, with 5% N'N-dimethylformamide and 0.01% hydrogen peroxide for about 20 min]. The peroxidase reaction caused an intense red precipitate at the binding site of the primary antibody. The sections were then rinsed in tap water, counterstained with Harris' haematoxylin and mounted with glycerol gelatin.

Negative controls were performed in each case by substituting the primary antibody with PBS. No staining was observed except for scattered granulocytes due to endogenous peroxidase. This reaction was not blocked for the benefit of optimal antigenicity. Strongly stained stromal cells and/or endothelial cells and/or lymphocytes and histiocytes, always present in combinations characteristic of the respective antigen under study, served as positive intrinsic controls. The staining intensity corresponding to the integrin antigenic density in neoplastic cells was evaluated against the background of these cells and was assessed in a semiqualitative fashion as follows: +, strong staining; (+), weak staining; - no staining. In the presence of positive and negative neoplastic cells a semiquantitative evaluation was carried out: +/- indicates positive and negative tumour cells in about equal amounts, +>- indicates a clear prevalence of positive tumour cells; ->+ indicates a clear prevalence of negative tumour cells.

Results

The distribution patterns of $\beta 1$, $\beta 3$, $\beta 4$, $\alpha 1$ - $\alpha 6$ and αv integrin subunits within the tumour cells of 32 SRBCS studied are shown in detail in Table 2 and are summarized in Table 3. Generally, the cells displayed a granular staining that was located on both the cell membrane and in the cytoplasm.

RMS

The RMS studied comprised primary and metastatic lesions in about equal numbers and included six embryonal and three alveolar RMS. All RMS were consistently β 1+, β 3-, β 4- (Fig. 1). The expression of the α chains was heterogeneous. Two embryonal and two alveolar RMS lacked any detectable α 1-6 and α v chains. The un-

Table 2 Detailed expression patterns of β1, β3 and β4 integrin subunits in neoplastic cells of small round blue cell sarcomas (RMS a alveolar rhabdomyosarcoma, RMS e embryonal rhabdomyosarcoma, UC undifferentiated tumour cells, RB rhabdomyoblasts, GNB ganglioneuroblastoma, Nbl neuroblasts, GC immature ganglion cells, NB neuroblastoma, pPNET primitive peripheral neuroectodermal tumour, ES Ewing's sarcoma, + strong positivity for all tumour cells, (+) weak positivity for all tumour cells, - negative for all tumour cells, +/- positive and negative tumour cells in about equal amounts, +>- positive tumour cells clearly dominate negative tumour cells, ->+ negative tumour cells clearly dominate positive tumour cells

Type		β1	β3	β4	α1	α2	α3	α4	α5	α6	αν
RMS a		+	_	_		_	_	_		_	_
RMS a		+	-	_	-	_	_		_	_	_
RMS a		+	_	_	-		_	+	_	+	_
RMS e		+	_	_	_		_			_	_
RMS e		+	_		-	_	_	_	->(+)		-
RMS e		+		_	_	_	_	_	+		(+)
RMS e		(+)	-	_	_	_	_	_	->+	+>-	(+)
RMS e		+	_	_	_	_	_				_
RMS e	UC	+	_	_	_	-	-	+/-	+>-	+/-	_
	RB	+	_	_	+	-	+/-	+	+	+	+
GNB	Nbl	+	-	_	_	_	+	-	_	(+)/-	
	GC	+	-	_		-	+	_			
GNB	Nbl	+	+/-	_	(+)>	_	+>	_	_	(+)/-	+
	GC	+	_	_	(+)>	_	+/-	_	_	(+)/-	_
GNB	Nbl	+	_		_		+	-		_	_
	GC	+	_	_	_	_	+			_	_
GNB	Nbl	+	_	_	+	_	+	_		_	_
	GC	+	_	_	_	_	+	_		_	_
GNB	Nbl	(+)	_	_	_	_	+	_	_	-	_
	GC	(+)	-	_	_	_	+	_	_		_ 、
NB		+	->(+)	_	+		+		_	(+)>-	(+)
NB		+	-	-	+		+		_	****	
NB		+	_	_	+	_	+	_		_	_
pPNET		(+)			_		_	_	(+)	_	_
pPNET		+	_	-	_	_	_	_	+	_	_
pPNET		+	_	-	_	_	_	_	(+)	_	+
pPNET		(+)	_	_	_	_	_	-	(+)>-	_	
pPNET		+	_	-	_	_	_	_	(+)>-	-	_
pPNET		+	_		_	_	_	_	+	_	_
pPNET		(+)	_			_	_	_	-	_	_
pPNET		(+)		_	_	_	- ,	-	->(+)	_	
ES		(+)	_			_	+/	_	+	_	_
ES		+	-	_	_	_	_	_	(+)	_	_
ES		+	_	_	_	_	_		(+)	*******	
ES		+	_	_	_	_	_		(+)	-	
ES		+	_	_	_	_	_	_	+	-	>+
ES		(+)			_	_	_	(+)	(+)	_	(+)
ES		(+)>-	_	_	_	_	_	_	+/(+)	_	(+)

Table 3 Summarized expression of β 1, β 3 and β 4 integrin subunits in neoplastic cells of small, round, blue cell sarcomas (Figures without parentheses indicate the number of cases in which all tumour cells expressed the corresponding integrin subunit, those with parentheses include the number of cases in which the respective antigen was detectable in at least a minor neoplastic population)

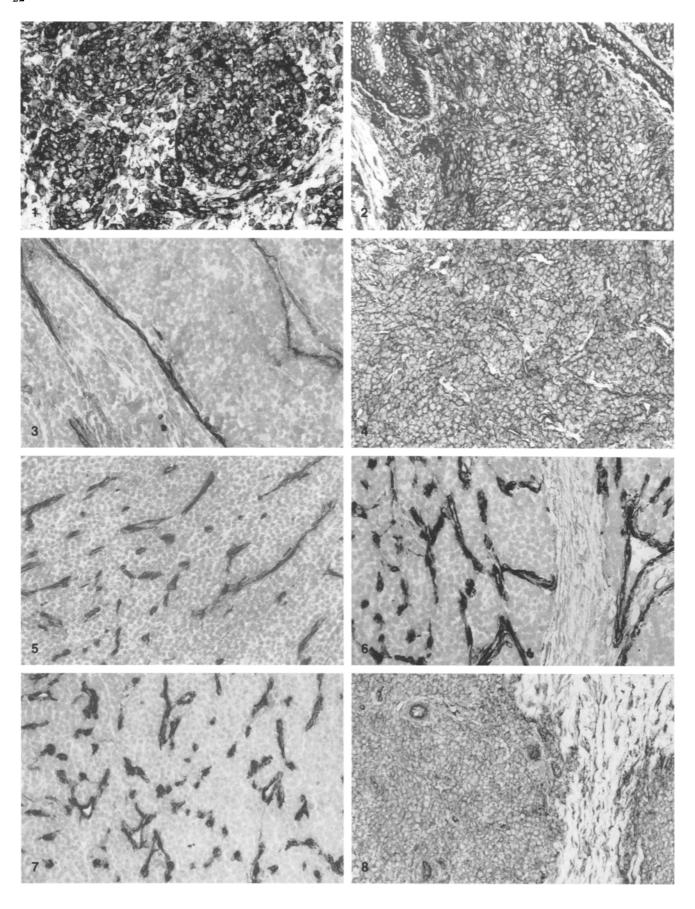
	RMS (<i>n</i> =9)	GNB (<i>n</i> =5)	NB (<i>n</i> =3)	pPNET (n=8)	ES (n=7)
β1	9	5	3	8	7
β3	_	(1)	(1)	-	_
β4	_	_		-	_
α1	(1)	(2)	3		_
$\alpha 2$	_	_	_	_	
$\alpha 3$	(1)	4(1)	3	~	(1)
α 4	1(1)		_	~	1
α 5	1(3)	_	_	4(3)	7
α6	1(2)	(2)	(1)		_
αν	2(1)	(1)	<u>i</u>	1	2(1)

differentiated tumour cell component of all RMS was $\alpha 1^-$, $\alpha 3^-$. One embryonal RMS contained a rhabdomyoblastic subpopulation which was $\alpha 1^+$, $\alpha 3^+$. One alveolar RMS was completely $\alpha 4^+$. In one embryonal RMS the rhabdomyoblastic tumour cells and about half of the un-

differentiated tumour cells were $\alpha 4^+$. The $\alpha 5$ subunit was expressed in the entire tumour cell population of one embryonal RMS. Three further embryonal RMS expressed $\alpha 5$ in a neoplastic subset. The $\alpha 6$ chain was consistently present in one alveolar RMS (Fig. 2) and heterogeneously expressed in two embryonal RMS. Two embryonal RMS were entirely αv^+ while in one embryonal RMS expression of αv was limited to a tumour cell subset.

(G)NB

The GNB and NB studied showed a strong expression of the $\beta1$ subunit. $\beta3$ was present in about half of the neuroblastic tumour cells of one GNB. One GNB and one NB expressed $\beta3$ in a neoplastic subset. All (G)NB were $\beta4^-$. Again, expression of the α chains was heterogeneous. All (G)NB were $\alpha2^-$, $\alpha4^-$ and $\alpha5^-$ (Fig. 3). $\alpha1$ was present in the neuroblastic cells of one GNB. Another GNB expressed $\alpha1$ in subsets of both the neuroblasts and ganglion cells. All NB were evenly $\alpha1^+$ (Fig. 4). Four of five GNB were entirely $\alpha3^+$, one case comprised $\alpha3^+$ and $\alpha3^-$ neoplastic cells in about equal amounts. All NB



were $\alpha 3^+$ throughout. The $\alpha 6$ subunit was focally expressed in ganglion cells and neuroblasts of one GNB and in the neuroblastic tumour cell population of one further GNB. One NB expressed $\alpha 6$ in the majority of tumour cells. Expression of the αv subunit was restricted to the neoplastic population of one NB and to neuroblastic tumour cells of a single GNB.

pPNET and ES

pPNET and ES showed a similar pattern of integrin subunit expression: all pPNET and ES were essentially $\beta1^+$, $\beta3^-$, $\beta4^-$, $\alpha1^-$ to $\alpha4^-$, $\alpha6^-$ (Figs. 5, 6, 7). As exceptions, one ES contained a $\beta1^-$ neoplastic subset, another case was focally $\alpha3^+$ and one further ES was evenly $\alpha4^+$. Some degree of heterogeneity emerged concerning the expression of the $\alpha5$ subunit. Four pPNET were entirely $\alpha5^+$ and three further pPNET were partly $\alpha5^+$. All seven ES were evenly $\alpha5^+$ (Fig. 8). The αv subunit was expressed in one of eight pPNET and in two of seven ES; one further ES showed an αv^+ neoplastic subset.

Against the background of tumour type three integrin subunit phenotypes emerged: RMS were essentially $\beta1^+$, $\alpha1^-$, $\alpha3^-$ and expressed $\alpha5$ and $\alpha6$ heterogeneously, (G)NB were generally $\beta1^+$, $\alpha1^+$, $\alpha3^+$, $\alpha5^-$, $\alpha6^-$ and pPNET and ES had similar integrin profiles being $\beta1^+$, $\alpha1^-$, $\alpha3^-$, $\alpha5^+$, $\alpha6^-$. Thus, the different types of SRBCS exhibited characteristic differences in their integrin subunit equipment.

Discussion

This is the first comparative immunohistochemical analysis of $\beta 1$, $\beta 3$, $\beta 4$ integrin subunit expression in different types of SRBCS in situ. Apart from homologies in integrin subunit expression like the consistent expression of

- Fig. 1 Embryonal rhabdomyosarcoma. All rhabdomyoblastic and undifferentiated tumour cells are β 1+ (×178)
- Fig. 2 Alveolar rhabdomyosarcoma. All tumour cells are $\alpha 6^+$. Duct remnants of infiltrated mammary gland are also $\alpha 6^+$ (×142)
- Fig. 3 Neuroblastoma. The tumour cell population is $\alpha 5^-$ contrasting with the $\alpha 5^+$ endothelial cells (×142)
- **Fig. 4** Neuroblastoma. The entire neoplastic population is $\alpha 1^+$ ($\times 214$)
- Fig. 5 Primitive peripheral neuroectodermal tumour. The tumour cell population is evenly $\beta 4^-$. $\beta 4^+$ endothelial cells serve as a positive intrinsic control of the immune reaction (×129)
- Fig. 6 Primitive peripheral neuroectodermal tumour. All tumour cells are $\alpha 1^-$. Endothelial cells are strongly $\alpha 1^+$ (×129)
- Fig. 7 Primitive peripheral neuroectodermal tumour. Expression of $\alpha 3$ is restricted to endothelial cells while being absent in the tumour cell population (×129)
- **Fig. 8** Ewing's sarcoma. The entire neoplastic population is $\alpha 5^+$. Endothelial cells and stromal fibrocytes/-blasts are $\alpha 5^+$ as well (×142)

the $\beta1$ chain and the absence of the $\alpha2$ and $\beta4$ chains in all SRBCS studied, we found three immunophenotypic patterns associated with the different types of SRBCS: RMS were $\alpha1^-$, $\alpha3^-$ and showed a heterogeneous expression of $\alpha5$ and $\alpha6$, (G)NB were $\alpha1^+$, $\alpha3^+$, $\alpha5^-$, $\alpha6^-$ and pPNET and ES were close to identical in their integrin subunit profile in being $\alpha1^-$, $\alpha3^-$, $\alpha5^+$, $\alpha6^-$. Thus, the three different modes of $\alpha1$, $\alpha3$, $\alpha5$, $\alpha6$ expression can be regarded as novel characteristics of RMS, (G)NB and pPNET/ES.

Our results are in most aspects in accordance with the rare data on integrin subunit expression in SRBCS in situ but contrast in some instances to findings observed in SRBCS cell lines in vitro.

Concerning RMS, current knowledge on the expression of integrins is limited to a few studies in vitro. Thus, the RMS cell line RD was shown to express the β 1, α 1, $\alpha 4$, $\alpha 5$ and $\alpha 6$ subunits while lacking the $\alpha 2$ and $\alpha 3$ chains [9, 40]. Expression of the α 4 and α 5 subunits was confirmed in various other RMS cell lines [6, 43]. Comparable to these data in vitro, some of our RMS expressed the $\alpha 4$, $\alpha 5$ and $\alpha 6$ subunits in parallel to the $\beta 1$ chain. However, in some cases the β chain was expressed in the absence of any detectable β 1-associated α subunit. It is conceivable that in RMS cells the β1 chain associated with another, as yet unknown α subunit to form a functionally active $\alpha\beta$ heterodimer. One such candidate, called α' or H36- α 7, has been described in chicken and murine myoblasts [25, 34]. Experimental evidence suggests that the presence of $\alpha 2$ in a RMS context might augment the metastatic potential of this tumour. Thus, Chan et al. [9] showed that the constitutively $\alpha 2^-$ and poorly tumourigenic RMS cell line RD had an enhanced growth rate and a higher capacity to form metastases in nude mice once successfully transfected with α 2 cDNA. In our study, although limited to a small number of specimens, all RMS were $\alpha 2^-$, both at the primary site and within the metastatic lesions. This does not exclude, however, that the disseminating RMS cells might have been $\alpha 2^+$ during the metastatic cascade.

NB and GNB were characterized by the presence of β 1, α 1 and α 3 chains while the other integrin subunits studied were absent in the clear majority of (G)NB studied. These data are in agreement with in situ studies performed by other investigators [13, 17]. In vitro, however, Yoshihara et al. [43] described differences in integrin subunit expression between the three different morphological types [neuroblastic (N-type), intermediate and substrate adherent (S-type)] NB cells. In line with our findings in situ N-type NB cells were characterized by very low levels of α5 expression when compared to Stype NB. Unexpectedly, however, expression of α 3 was more prominent in S-type NB cell lines. Moreover, in contrast with our data in situ and those presented by others [13, 17], expression of $\alpha 4$ was found in a variety of NB cell lines, irrespective of the morphological type [43]. These discrepancies might be due either to a clonal selection during the establishment of the cell lines or to culture conditions. Such effects have been proposed as an explanation for conflicting in situ/in vitro data in the melanoma system [2].

The pPNET and ES studied showed close to identical integrin profiles. In addition to the $\beta1$ subunit, pPNET and ES expressed the $\alpha5$ subunit in nearly all cases while most other $\beta1$ -associated integrin subunits, especially the $\alpha1$ and $\alpha3$ chains, were undetectable or only extremely rarely found in these tumours. This is in accordance with the phenotype of a PNE cell line that was $\alpha1^-$ but $\alpha5^+$ in vitro [11]. Furthermore, expression of $\alpha5$ was a consistent feature in three ES cell lines examined in vitro [43].

The almost identical integrin profile of pPNET and ES adds further evidence to the close nosological relationship of these two tumours that up to date is based on the common reciprocal translocation t(11,22)(q24,q12) [41, 44], the common expression of the MIC-2 antigen [3], and other immunophenotypic correspondences such as the consistent expression of HLA-A,B,C and the absence of CD9 and CD56/neural cell adhesion molecule [15, 26].

SRBCS are currently regarded as a morphologically rather uniform group of highly malignant and aggressive sarcomas. The phenotypic differences among the different types of SRBCS we have described herein most likely reflect differences in the mode of tumour growth and spread, since the molecules involved are fundamentally important regulators of the social behaviour of cells in tissues. However, one should refrain from speculating too much on the net effects that these different integrin receptor profiles might have in vivo. At this point we do not know whether the integrins expressed on SRBCS are assembled in a functional fashion. Furthermore, their ligand specificity will have to be re-defined for the cell types we have been dealing with. We cannot exclude the possibility that integrin functions and ligands not yet known might play a significant role. Nevertheless, it may well be that the adhesion receptor profile contributes to the fact that SRBCS typing is of clinical importance.

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