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Keywords:	CD133, CNS neoplasms, stem cells



CD133 expression is associated with small round blue cell tumour morphology in human CNS neoplasms,

Jens Schittenhelm¹, Perikles Simon², <u>Patrick N. Harter⁶, Cornelia Zachskorn⁶</u>, Holger Schlaszus¹, Frauke Röttger³, Monika <u>Winkels³</u>, Michael Weller^{4,5}, Richard Meyermann¹, Michel Mittelbronn⁶

¹Institute of Brain Research, University of Tuebingen, Germany; ²Department of Sports Medicine, Rehabilitation and Disease Prevention, Johannes Gutenberg University, Mainz, Germany ³Miltenyi Biotec GmbH, Bergisch-Gladbach, Germany ; ⁴Department of Neurology, Hertie-Institute of Clinical Brain Research, University of Tuebingen, Germany; ⁵Department of Neurology, University Hospital of Zurich, Switzerland; ⁶Institute of Neurology (Edinger Institute), Goethe University, Frankfurt, Germany

Address for correspondence: Michel Mittelbronn, MD Institute of Neurology (Edinger Institute) Johann Wolfgang Goethe University D-60325 Frankfurt Germany Tel: +49(0)69-6301-84169 Fax: +49(0)69-6301-84150

Key words: CD133, CNS neoplasms, stem cells Running title: CD133 in CNS neoplasms Deleted: in primary and secondary human CNS neoplasms Deleted: (SRBCT) Deleted: s

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Abstract:
Aims: CD133 is considered as a marker for brain tumour initiating cells. However
most data about CD133, is derived from animal or in vitro studies. The aim of our
study was to characterize CD133 expression, distribution and morphological features
of CD133 ⁺ cells in primary and secondary human CNS neoplasms.

Methods and results: Tumours were analyzed by realtime RT-PCR, western blot, flow cytometry and immunohistochemistry. Our results show that only small round blue cell tumours (SRBCT) exhibit strong and consistent CD133 expression. Interestingly, glioblastomas, large cell carcinomas or sarcomas were negative for CD133. Only single glioblastomas with focal small cell component exhibited CD133 immunoreactivity in the SRBCT component. In addition, CD133 expression did not correlate with the expression of other markers associated with stem cell differentiation including CD15 or nestin.

Conclusions: This indicates that CD133 expression in human CNS neoplasms may be independent from the grade of malignancy but strongly correlates with SRBCT morphology. Together with recent findings showing that CD133 is regulated by hypoxia and CD133⁻ cells exhibiting also stem cell properties, our data strongly guestions the suitability of CD133 as a brain tumour stem cell marker, in vivo.

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Introduction:

Over 20 years ago, first evidence supported the hypothesis that neoplasms might arise from common stem cells capable to give rise to tumours of various differentiation.¹ Since human stem cells possess the capacity for self-renewal as well as unlimited growth and current treatment strategies mostly fail to eradicate tumour stem cell,² Therefore, the actual aim is not only to treat the tumour bulk, but also to focus on cancer stem cells.³ Cancer stem cells are <u>functionally</u> defined by assays of self-renewal, multilineage differentiation and sphere formation and it was claimed that these cells are <u>could be selected</u> by the five-transmembrane protein CD133 (prominin-1).⁴ <u>After stimulation with either epidermal growth factor (EGF) or basic</u> fibroblast growth factor (bFGF) multipotent, self-renewing neural stem cells proliferate <u>within the spheres</u> and are considered <u>as</u> candidate cells for autologous transplantation after neuronal loss.^{5,6}

The CD133 antigen was detected on germ cells and on progenitor cells of all germ layers, including ectoderm, mesoderm and entoderm as well as neuroectoderm.⁷⁻¹¹ <u>More recently</u>, different splice isoforms of the CD133 antigen were described.^{7,12} While CD133-1 was <u>stronger expressed in the</u> fetal brain and adult skeletal muscle, CD133-2 (the 9 amino acid shorter variant of CD133-1), showed higher expression levels on hematopoietic stem cells and was inversely correlated <u>with</u> the terminal differentiation marker involucrin pointing to a <u>spatio-temporal</u> down-regulation of CD133-2 with the grade of cell maturation.¹² The AC133/AC141 epitopes bound by the widely used monoclonal CD133 antibodies are glycosylated and regulated independently of the much more widespread expression of their mRNA<u>s</u>.¹³

While the number of CD133⁺ cells increased with the grade of malignancy in some tumours like melanocytic neoplasms, others such as prostate cancer failed to show a

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correlation of the amout of CD133⁺ cells and tumour grade.^{14,15} Very recently, CD133

expressing tumour cells <u>were shown to be responsible for radio- and</u> chemoresistance in gliomas, ^{16,17}

Another study revealed that CD133 positive progenitor cells considerably contribute to tumour neo-vascularization.¹⁸ One way to overcome, the long-term propagation of CD133 positive stem cells in neoplasms is the blockade of the Notch signaling pathway finally leading to cell cycle exit, apoptosis and cellular differentiation via its downstream target Hes1.¹⁹ The amount of circulating CD133 positive cells seems to be regulated by chemotactic factors such as interleukin-8 which on the other hand is significantly upregulated in various neoplasms.²⁰⁻²³ In an epithelial cell model, one feature of both CD133 isoforms seems to consist in organizing plasma membrane protrusions.²⁴ Since morphological features are potentially mediated by CD133 under normal conditions, the aim of our study was not to use functional assays which may alter number and expression level of CD133-positive cells but rather to analyze morphological properties and the distribution pattern of the so-called CD133- positive tumour cells within various human neoplasms under conditions that more closely resemble the *in vivo* conditions in the human brain. In the present study we analyze tumours not only in vitro but also fresh viable tumour tissue and paraffin embedded specimens, since many cell culture conditions may irrevocably alter the cells.²⁵ This might be important for further therapeutical approaches focussing on the detection of tumour stem cells. Herein, we demonstrate that CD133⁺ expression is strongly associated with a small round blue cell tumour (SRBCT) phenotype representing neoplasms with a very primitive morphological differentiation. In contrast, other than SRBCTs – even though being very malignant – were almost devoid of CD133⁺ cells.

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Materials and Methods:

Cell lines

CD34-positive stem cells, burkitt lymphoma cells, the malignant glioma cell lines A172, LNT-229, LN-308, TG98G, U373MG (kind gift of Dr. N. Tribolet, Lausanne), 4 different neuroblastoma cell lines (N-Myk single copy, N-Myk 25, N-Myk 30, N-Myk 100), the retinoblastoma cell line WERI-Rb1, the medulloblastoma cell line DAOY (kind gift of Dr. K. Schilbach, Tübingen) were maintained in DMEM supplemented with 2 mM L-glutamine (Gibco Life Technologies, Paisley, UK), 10% FCS (Biochrom KG) and penicillin (100 IU/ml) / streptomycin (100 µg/ml) (Gibco). CD34+ stem cell were mobilized in healthy donors. Regimen for mobilization of donor stem cells was 10µg/kg/g G-CSF s.c. 5 days prior to leucaphereses, performed on days 5 and 6. CD34+ progenitors were positively selected from leucapheresis products by magnetic-activated cell sorting using Clinimax CD34 reagent and Clinimacs device from Miltenyi Biotec (Bergisch Gladbach, Germany). Purity of CD34+ population was > 99.99%. Progenitor cells were suspended in DMSO and stored in liquid nitrogen until use. Informed consent was obtained from all CD34+ cell donors in accordance with the Declaration of Helsinki. In addition, several primary glioblastoma cells were cultured under different media conditions.

Patients and tissue samples

We investigated a total of 641 paraffin tumour samples obtained from the brain tumour bank of the Institute of Brain Research, University of Tuebingen. These consisted of 295 samples of 213 glioblastoma cases (including 13 cases with known small cell component), 173 samples of 132 astrocytoma cases (including 27 WHO grade I, 44 WHO grade II and 61 WHO grade III tumours), 12 primary cerebral B-cell

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lymphomas, 67 medulloblastomas (including 11 desmoplastic variants), 10 CNS-PNET, 19 small cell lung carcinoma metastases, 30 non-small cell carcinoma metastases, 10 melanomas, 10 sarcoma metastases, 6 neuroblastomas (2 peripheral abdominal, 1 central spinal, 3 esthesioneuroblastomas), 5 pineoblastomas and 4 retinoblastomas (Supplementary Table 1), In addition 79 samples of 24 autopsy cases with normal human brain belonging to an established brain bank (Brain Bank Tübingen) and 26 specimens of cases with biopsies of non-neoplastic CNS lesions (Supplementary Table 2) were also included. In all cases mentioned above full mount sections were stained with the exception of astrocytoma samples where additional samples were partially constructed as tissue microarray (Manual Tissue Mircroarrayer, Beecher Instruments, Sun Prairie, USA; two representative cores measuring 600µm for each tumour; n=78 WHO grade IV, n=75 WHO grade III, n=35 WHO grade II and n=24 WHO grade I). For western blot and RT-PCR frozen tissue of selected CD133-positive and negative cases (n=25) was studied. Utilization of the human specimens was in accordance with the University's ethics commission. Pathological diagnoses were made by at least two experienced neuropathologists according to the WHO criteria.²⁶

Preparation of RNA and reverse transcription to cDNA

Total RNA was isolated from the various cryo-preserved tissues and cell lines after homogenisation using Trizol Reagent (Invitrogen, Karlsruhe, Germany). All samples were subjected to treatment with RNase free DNase I (Roche Diagnostics, Mannheim, Germany). Consecutively, 500 µg of each RNA sample was reversetranscribed with 200 U of Moloney Murine Leukemia Virus Reverse Transcriptase (M-MLV-RT, Promega, Mannheim, Germany), using 0.5 µg of Random Primer

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(Promega). Glyceraldehyde-3-phosphate-dehydrogenase (gapdh) served as housekeeping-gene.

Quantitative real-time RT-PCR

Quantitative real-time RT-PCR including primer-design and control of the specificity and efficiency of the primer pairs was done as described previously. ^{27,28} The sequences of the PCR primers used in this study are listed in table 1. All primers were chosen to span an intron in order to control for and avoid priming of gDNA contaminations. The sense primers were constructed to contain an intron spanning internal to the respective sense primer, which was specific for the respective splice variant of CD133 (see Figure 1 for details). This procedure enabled us to avoid any priming of gDNA, since the sense primer are cDNA specific and to ensure maximum comparability of differential CD133_1 and CD133_2 expression by reducing technical artefacts using the same reverse primer and generating similar amplicon lengths. Quantitative real time RT-PCR was linear over the expression range. For the evaluation of all PCR runs, 'Cycle Threshold' values (CTs) obtained for the different amplicons were processed to Mean Normalized Expression (MNE) as described previously.²⁹ Replicates within one plate were used for statistical analysis. MNE values and the respective 99%-confidence intervals (CI) of the means are reported.

Immunoblotting

For immunoblotting, <u>12</u>, tumour samples, <u>5</u> established cell lines and <u>4</u> primary glioblastoma cell cultures were used. Tissue specimens were homogenized and placed overnight on ice in Tris-HCI-buffer pH 7.6 containing 125mM NaCI, 2.5% Brij 96/98 and a protease inhibitor cocktail (Complete[™], Roche). After centrifugation the

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remaining pellet was resuspended in 50mM Tris; 150mM NaCl, 1% Triton X100 and 1% NP-40 placed on ice for 6 h. Protein concentration was determined photometrically at OD₂₈₀ against lysis buffer. The protein lysates were stored at -80 $^{\circ}$ C pending immunoblotting. Whole-cell extracts for SDS-PAGE were prepared by solubilizing washed cells in hot SDS-sample buffer and incubating at 95℃ for 5 minutes. Proteins were separated by 10% SDS-PAGE and subsequently transferred onto nitrocellulose membranes. MBI Fermentas Prestained Protein Ladder was used as molecular weight standard. Following electrophoresis, the samples were transferred to a polyvibylidene difluoride (PVDF) membrane (Hybond, Amersham) and blocked overnight with 5% bovine serum albumin BSA in Tris-buffered-saline-Tween (TBST; 50 mM Tris-HCl pH 7.6, 150 mM NaCl, 0.1% (v/v) Tween-20). The membranes were incubated with primary antibodies in TBST (1µg/ml) against CD133 clones W6B3C1 (recognizing CD133-1) and 293C3 (recognizing CD133-2, both from Miltenyi, Bergisch-Gladbach, DE), Synaptophysin (Biogenex, San Ramon, USA), pan-cytokeratin (Dako, Glostrup, DK) and glial fibrillary acidic protein GFAP (Neomarkers, Freemont, USA). Exposition times were 3 to 5 mins. A subsequent

incubation with an antibody against beta-actin (Santa Cruz) was performed to determine equal protein loading of each lane.

Flow Cytometry

Immunocytofluorescence staining for CD133 monoclonal antibodies W6B3C1 (recognizing CD133-1) and 293C3 (recognizing CD133-2, both from Miltenyi) was performed on cultured cells. 1×10^6 cells were collected, washed in PBS, and incubated with 100µl 3.7% formaldehyde for 30mins. After addition of 1ml PBS with 2mM EDTA and 0.5% bovine serum albumin (PEB), cells were permeabilized in 100µl 0.5 saponin buffer (20mins) before incubation with the primary antibodies for

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15mins (5µg/ml in saponin buffer). After a second washing step, cells were incubated with isotype-matching rat anti-mouse IgG fluorochrome-conjugated secondary antibodies at optimal dilutions for another 15mins (Miltenyi). Finally cells were washed again with PEB. One color analysis was performed on a FACS (FACSCalibur, BD, NJ, USA) using scatter gate with a minimum of 5,000 events counted. Isotype controls were performed for each antibody.

Immunohistochemistry

Immunohistochemistry was performed as described previously.³⁰ Briefly, samples were put in 4% formalin (pH 7.4) after surgery and routinely embedded in paraffin. Four-micron-thick sections were stained routinely with hematoxylin and eosin and in adjacent serial sections, immunohistochemistry was performed with mouse immunoaffinity-purified IgG1 antibodies against CD133 clones W6B3C1 (dilution 1:50) and AC133 (1:10, both from Miltenyi). Deparaffinized and rehydrated sections were immunostained by using the Benchmark Immunohistochemistry System (Ventana, Tucson, AZ, USA). A standard protocol including a cell conditioning pretreatment for 60mins was used. I-View-Inhibitor (Ventana) was applied for 4min and endogenous peroxidase of the tissue sections was blocked with 3% H₂O₂ in methanol for 14min. CD133 antibodies were applied with an incubation time of 32 min. Incubation of amplifier A and B for 8min each were followed by avidin and biotin blockers for 4 mins respectively and I-View-Biotin-Ig for 8mins (all Ventana). Before diaminobenzidine/H₂O₂ visualization (8min), incubation with I-View SA-HRP (8min) was applied. Finally, secions were incubated with a copper enhancer (4min) andcounterstained with haematoxylin. Positive control (WERI-Rb1 cell culture slides)

and negative controls (omission of the primary antibody from the diluent) were run

with each batch of staining

Results:

Both CD133 splice variants are differentially regulated in vivo and in vitro.

CD133_1 as well CD133_2 showed the highest mean normalized expression levels in CD34-positive stem cells and in the retinoblastoma cell line WERI-Rb1 (Figure, 2). While both variants were detectable in four different neuroblastoma cell lines (N-Myk single copy, N-Myk 25, N-Myk 30, N-Myk 100) and in a medulloblastoma cell line (DAOY), the CD133_2 splice variant was not detectable in two malignant glioma cell lines (A172, LN308) or showed levels just above the treshold (LNT-229). In a burkitt lymphoma cell line only CD133_1 mRNA expression was detectable but not CD133_2. Similiar levels of both CD133 mRNA splice variants were observed in vivo in small round and blue cell tumours (medulloblastoma, PNET and pineoblastoma

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tumour samples). In contrast CD133_1 expression levels in astrocytic neoplasms were either not detectable or displayed levels just above treshold, while CD133_2 expression was always present in higher levels.

In vitro and in vivo expression of CD133 is consistent only in small round blue cell tumours.

Immunoblot analysis using two different CD133 antibodies (W6B3C1 and 293C3) allowed the detection of 120 kDa bands in WERI-Rb1 and DAOY cell-lines (Figure, 3a) but were lacking in U373, T98G and LNT-229 glioma cell lines (data not shown). Presence of 43 kDa beta-actin or 49 kDa GFAP bands in the latter served as control for protein integrity. Flow cytometry revealed membrane-associated CD133 (W6B3C1 and 293C3) in WERI-Rb1 (Figure 3b) and DAOY (Figure 3c) while only a small minority of glioma cell lines could be considered to express membranous CD133 (Figure 3d). Both lineage specific markers (e.g. CD56, Figure 3b) and CD133 were strongly expressed on SBRCT, while non SRBCT entities showed an absence or only very weak frequency of CD133 positive cells compared to isotype and lineage specific positive controls. The results of flow cytometry and immunoblotting nicely correlated with CD133 expression on cultured cells (Supplementary Figure 2). In vivo results were similar showing strongest <u>CD133</u> expression patterns in pineoblastomas and medulloblastomas, while all glioblastomas and most primary glioblastoma cell cultures showed no distinct bands for CD133 (Figure 3a and Supplementary Figure 1).

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CD133 immunoreactivity is mainly restricted to small round blue cell tumours in vivo. The CD133 antibody 293C3, which recognizes the CD133-2 antigen was not suitable for immunohistochemistry. Two different CD133 clones directed against CD133-1 (W6B3C1 and AC133) showed similar results in test rows, with slightly nicer staining results for AC133, Therefore, AC133 was used for further evaluation. We examined 79 different normal brain samples of 24 autopsy cases including the cerebrum, the cerebellum and the brainstem. No AC133-immunoreactive cells were found in normal control CNS tissue specimens (Supplementary Figure 3). Six out of 56 medulloblastomas (Fig.4a-c) and 2 of 11 desmoplastic medulloblastoma variants showed strong expression of AC133 on the majority of tumour cells. In addition 3 out of 5 pineoblastomas (Fig. 4d and e), 4 out of 4 retinoblastomas (Fig. 4f), 1 out of 6 neuroblastomas (an esthesioneuroblastoma) (Fig. 4g) were immunopositive for AC133. Further, formalin-fixed and paraffin embedded cell pellets of different cell lines exhibiting CD133-positive (W6B3C1 and 293C3) phenotype in both FACS and WB analysis such as DAOY or WERI-Rb1 (Fig. 4i) kept their immunoreactivity also in AC133 immunocytochemistry (Supplementary Figure 2). All AC133 positive tumour cells exibited the typical morphology of small round and blue cells with large nucleus and small cytoplasm. In addition all 295 samples of 200 classical glioblastomas (Fig. 4j, k), all 173 samples of 27 WHO grade I, 44 WHO grade II and 61 WHO grade III astrocytomas remained negative for AC133. We further examined nonneuroepithelial tumours. All 12 primary cerebral B-cell lymphomas, 10 malignant sarcomas and 10 melanomas were devoid of AC133 expression. 9 out of 19 small cell lung carcinomas were immunoreactive for AC133 (Fig. 4h) while in the group of 30 non-small cell carcinomas two metastases of breast cancer focally displayed granular immunoreactivity in the cytoplasm of few tumour cells for AC133. None of

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the 16 non-small cell lung carcinoma metastases included in this group <u>showed</u> <u>positivity</u> for AC133. <u>CD133 did not at all correlated with the expression of other</u> <u>putative stem cell markers such as CD15 or nestin (Supplementary Figure 4)</u>.

CD133 expression in glioblastomas are restricted to GBM with small cell component. Since all conventional glioblastomas examined were immunonegative and AC133 expression was restricted mainly to tumour cells with small round and blue phenotype we <u>further</u> examined 13 glioblastomas with <u>more</u> undifferentiated <u>tumour</u> areas - a so called small cell component. These samples only included cases with history of prior radiotherapy. no known In one case we observed immunohistochemical expression of AC133 in a minority of such cells (Fig. 4I). All its w other normal astrocytic differentiated tumour cells within this glioblastoma remained immunonegative for AC133.

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Discussion:

Depending on the tissue of origin, putative cancer stem cells (CSC) have been identified by an array of different markers including CD34 for acute myeloid leukaemia and non small cell lung cancer, CD44 for breast and prostrate cancer, while colorectal cancer and brain tumour CSC are considered to express the cell surface marker CD133 (Prominin-1).^{10,15,31-34} CD133 positive brain tumour-initiating cells are capable to proliferate in serum-restricted medium by forming tumourigenic neurospheres and differentiate into mature cells under normal serum-conditions.³⁵ These cells may exhibit increased radio- and chemoresistance through increased DNA repair mechanisms.^{16,17,36,37} CD133-positive brain tumour cells have also been reported in retinoblastomas and medulloblastomas - tumours known to exhibit a poorly differentiated morphology. Therefore, these tumours are also named small round blue cell tumours.^{34,38} In addition, it has been noted that CD133 positive cells isolated from glioblastomas show absence or very low expression of glial fibrillary acidic protein suggesting that they might not be fully differentiated astrocytic tumour cells.^{37,39}

In the current study we aimed to determine the expression levels of CD133 in various brain tumours with emphasis on immunohistochemistry to identify the morphology of CD133-positive tumour cells. Surprisingly, neither immunoblotting nor immunohistochemistry revealed a considerable expression of CD133 in commonly used glioma cell lines and in a large number of glioblastomas in contrast to high expression levels in many SRBCTs. These results were supported by absent or low-level CD133_1 expression in realtime quantitative PCR and CD133 expression data in flow cytometry being almost similar to isotype controls. This contrasts with data of

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other studies that have reported CD133 to be present in gliomas.^{34,40,41} These differences may be in part explained by the fact, that data in other studies were obtained on serum-free cultured neurospheres, while our cell lines were mainly grown under serum conditions. Even the minimal CD133-positive expression in astrocytic cell lines in our flow cytometry analysis could not be detected by immunohistochemistry although we screened a large number of glioblastoma tumour samples. Our observation, is, further corroborated by an independent flow cytometry, study reporting less than 1% viable CD133+ cells in astrocytomas and grades.42 oligodendrogliomas WHO independent of their CD133 immunohistochemistry is considered to be difficult, therefore many groups only use frozen tissue. The use of strong signal amplification techniques results in high numbers of CD133-positive cells also in astrocytomas. 43-45 Since these numbers are higher than <u>flow cytometry</u> results suggests, this indicates a possible overstaining. <u>Our</u> carefully adjusted immunohistochemical protocol permits a strong and unambiguously interpretable staining pattern without background staining on paraffin sections using the AC133 clone.

We were able to demonstrate that CD133 in medulloblastomas is not only expressed *in vitro*³⁶ but also *in vitro* and that other tumours with a small round <u>blue cell</u> phenotypes such as pineoblastoma, neuroblastoma, retinoblastoma and and small cell lung cancer specimens often consist of <u>completely</u> CD133-positive tumour cells. To our knowledge, CD133 expression in some of the entities in our study (e.g. pineoblastomas) has not been reported so far. The role of the the CD133_2 splice variant which was present in all tumours examined still needs further investigation, Interestingly in epithelial tumours both isoforms seem to be equally distributed at low levels.⁴⁶ CD133 expression was also observered in a significant number of cases of

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small cell lung cancer but <u>was not fully restricted to a SRBCT phenotype in epithelial</u> <u>neoplasms</u> in <u>contrast to</u> all neuroepithelial tumours. However, <u>both CD133-positive</u> cases_were metastases of breast cancer <u>in which</u> CD133 expression has been reported and for which CD44 (and not CD133) is considered <u>to be a putative stem</u> cell marker.^{33,47}

It is noteworthy that multipotent neural stem cells present in the adult brain (located within the subgranular zone of the hippocampal dentate gyrus and the subventricular zone of the lateral ventricles) morphologically resemble undifferentiated small round plue cells and not astrocytes.⁴⁸ Furthermore CD133⁺ cells in neuroepithelial SRBCTs frequently show a perivascular distribution pattern with signs of clonal expansion suggesting that there might be the possibility that these cells are integrated hematopoietic stem cells rather than primary tumour intrinsic cells. The recruitment of hematopoietic progenitor cells by VEGF expression of tumour cells has been already demonstrated.⁵⁴ A close relationship between CD133 positive cells and adjacent endothelial cells was noted in *in vitro* experiments showing that these cells are able to secrete VEGF. ^{16,49} CD133-positive cells not closely related to tumour vessels have been reported to reside in pseudopalisading areas of necroses. ⁴⁴ These areas are mainly subjected to low oxygen concentrations. From cell experiments it is known that cultured glioma cells are capable to upregulate CD133 on protein and mRNA level when kept unter hypoxic conditions.^{13,50,51}

Very recent studies demonstrated that primary glioma cell cultures analysed by flow cytometry, did, also consist of CD133-negative tumour cells, that fulfilled stem cell criteria. These cells were tumourigenic in nude mice. Furthermore, CD133-positive cells could also be obtained from CD133-negative tumourigenic cells.^{41,52,53} Even though the CD133 portion represents undifferentiated cells, only a minority of these

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glioblastomas. ⁵⁵ These data led to the notion that CD133 expression in freshly	
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In conclusion, our data shows that CD133 expression in human CNS neoplasms is	
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correlates with SRBCT morphology.

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Figure legends:

Figure 1: Position of sense and antisense primers on the different positions for the CD133_1 and CD133_2 splice variants generating similar amplicon lengths. The pairs were constructed spanning at least one intron to avoid priming possible genomic DNA contaminations.

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Figure 4: Immunohistochemical results (CD133, clone AC133): Strong membranous staining of small round blue cell tumours: a-c): medulloblastomas; d-e) pineoblastomas; e) ; f) retinoblastoma; g) neuroblastoma <u>(esthesioneuroblastoma)</u>; h) metastasis of small cell lung carcinoma; i) retinoblastoma cells (WERI-Rb1). Within the CD133-positive small round blue cell tumours, endothelial cells or perivascular non-neoplastic cells remain immunonegative. j-k) classical

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glioblastomas without small round blue cell features were completely immunonegative for CD133; I) In contrast only small round and blue cells of a glioblastoma partially exhibiting small cell differentiation show a rather diffuse pattern.

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Supplementary Figure 1: Immunoblotting results (CD133, clone W3B6C1: 120kDa band, control: beta-actin: 43 kDa band). WERI-Rb1 cells were used as positive control. In addition, fresh frozen tissue of 4 different primary glioblastomas and primary cell cultures of 3 different primary glioblastomas under different growth conditions were investigated.

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Supplementary Figure 3: CD133 expression in biopsies of non-neoplastic CNS diseases. In contrast to small round blue cell neoplasms such as medulloblastomas (a), non neoplastic CNS diseases including epilepsies (b: Wyler grade 3; c: Wyler grade 4) as well as surrounding tissue of arterio-venous malformations (d), epilepsy brains without gross morphological changes (e) or cerebellar bleeding (f) remained negative for CD133.

Supplementary Figure 4: Comparison of CD133, CD15 and nestin expression in CNS neoplasms (n=30). (A) WERI-Rb1 for CD133, (B) a brain infarction for CD15, (C) and a fetal brain of the for nestin were used as positive controls. Small round blue cell neoplasms including pineoblastoma (D-F) and medulloblastoma (G-I) were labelled for CD133 (D, G), CD15 (E, H) and nestin (F, I). CD133-negative normal brain (J) and astrocytoma ((K) pilocytic astrocytoma WHO grade I, (L) diffuse astrocytoma WHO grade II, (M) anaplastic astrocytoma WHO grade III and (N, O) glioblastoma WHO grade IV) cases were further investigated for nestin (J-N) and CD15 (N).

Tables:

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Table 1: Primer sequences used in this study for real-time RT-PCR. All primers were

chosen to span at least one intron to control for and exclude the possibility of

contamination with genomic DNA.

Product mRNA Sense Primer Antisense Primer Size 5'TTG ATT ATG ACA5'ACC CCA CCA GAG GCA 120 bp Cd133 1 AGC CAG AAA CTG3' TCA G3' 5'GAC AAG ATT GTC 5'ACC CCA CCA GAG GCA_{85 bp} Cd133 2 TAC TAT GAA GCA TCA G3' GG3' 5'GCC TGG AGA AAC5'TGG TCC TCA GTG TAG96 bp GAPDH CTG CCA AG3' CCC AAG3'

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28

Supplementary Table 1: Epidemiological data. For tumour specimens received from

other pathological institutes, no epidemiological data were obtained

Deleted: contamination with genomic DNA.

(e.g.

neuroblastomas, retinoblastomas).

Diagnosis Glioblastoma Astrocytoma WHO III	mean age	<u>sex (m/f)</u>	
	<u>58.4</u>	54/35	
ASTOCYTOTIA WHO III	46.0	32/20	
Astrocytoma WHO II	43.7	20/16	
Astrocytoma WHO I	21.1	16/9	
Medulloblastoma	19.7	41/25	
PNET	44.1	9/1	
Neuroblastoma	N/A	N/A	
Carcinoma	63.4	24/9	

Supplementary Table 2: Epidemiological data of cases with biopsies of non-

neoplastic CNS lesions.

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Diagnosis	Age	Sex
pineal cyst + adjacent CNS	40	М
pineal cyst + adjacent CNS	28	M
pineal gland + adjacent CNS	58	F
mesial temporal sclerosis Wyler 3	<u>35</u>	<u>F</u>
mesial temporal sclerosis Wyler 4	<u>28</u>	M
mesial temporal sclerosis Wyler 3	<u>37</u>	M
mesial temporal sclerosis Wyler 3	<u>35</u>	<u>F</u>
pineal cyst + adjacent CNS	<u>7</u>	Μ
pineal gland + adjacent CNS	<u>27</u>	M
CNS + Stroke	<u>54</u>	M
CNS + CAA	<u>81</u>	M
CNS + hematoma	<u>55</u>	M
Cerebellum	<u>84</u>	F
<u>CNS + AVM</u>	20	M
PML	<u>64</u>	<u>F</u>
<u>CNS + cavernoma</u>	<u>52</u>	M
<u>CNS + AVM</u>	<u>60</u>	M
<u>CNS + AVM</u>	27	M
normal CNS	55	E
Cerebellum + bleeding	<u>70</u>	F
<u>CNS + cavernoma</u>	21	M
<u>CNS + cavernoma</u>	<u>19</u>	M
CAA	<u>83</u>	M
Stroke	<u>36</u>	<u>F</u>
mesial temporal sclerosis Wyler 4	<u>36</u>	M
CNS + cavernoma	<u>38</u>	M



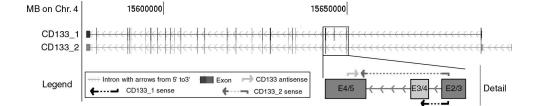


Figure 1: Position of sense and antisense primers on the different positions for the CD133_1 and CD133_2 splice variants generating similar amplicon lengths. The pairs were constructed spanning at least one intron to avoid priming possible genomic DNA contaminations. 157x33mm (300 x 300 DP1)

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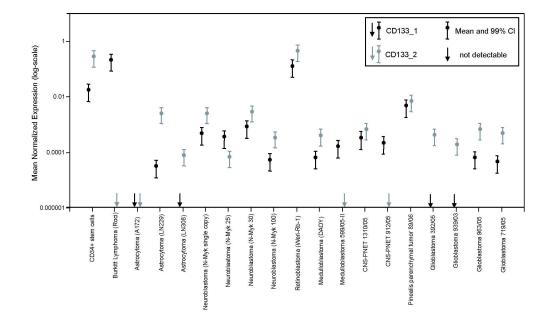


Figure 2: Realtime RT-PCR results. Course of expression for the different CD_133 splice variants in cultured cells and excised tumour samples. Means and 99% CIs for three measurement repetitions are given. 685x407mm (72 x 72 DPI)

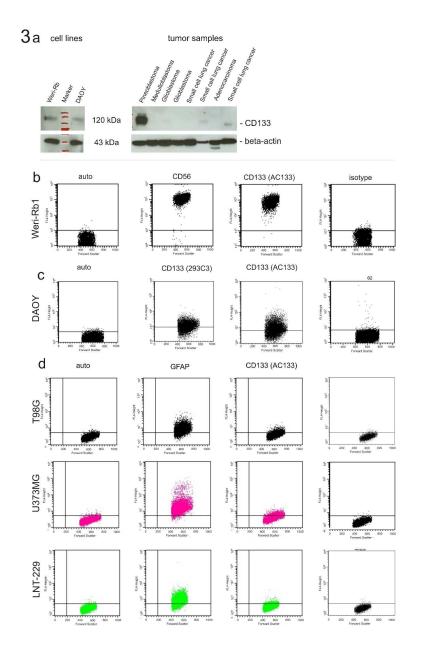


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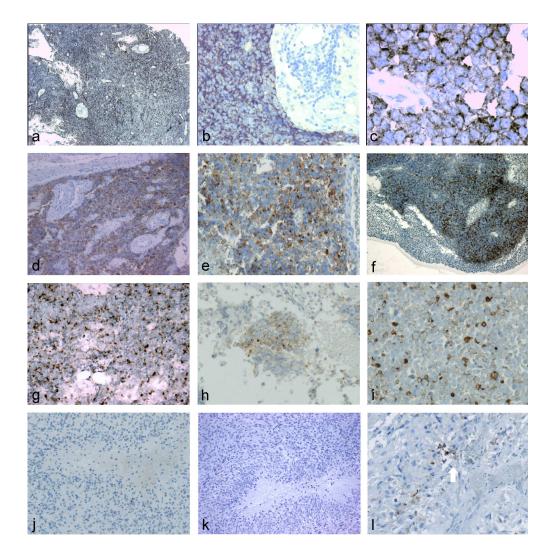
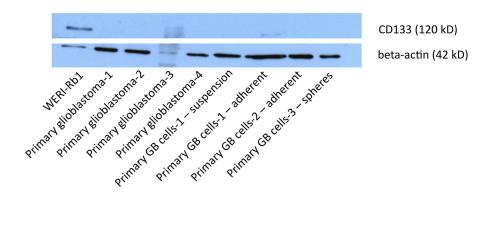
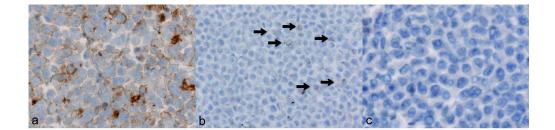


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 In contrast only small round and blue cells of a glioblastoma partially exhibiting small cell differentiation show a rather diffuse pattern.
 874x893mm (72 x 72 DPI)

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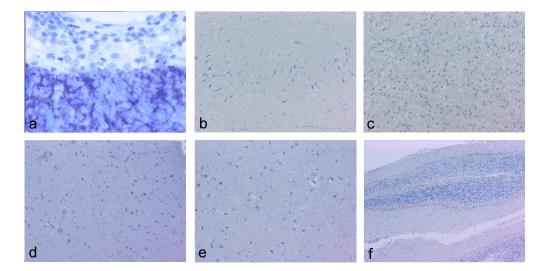


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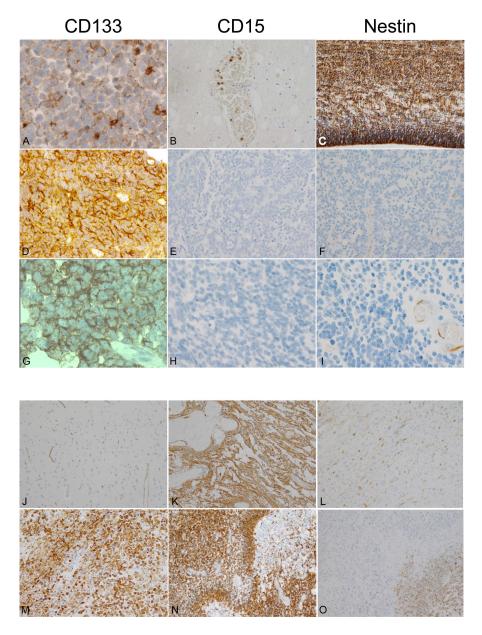


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1064x265mm (72 x 72 DPI)



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