

CD133 is a temporary marker of cancer stem cells in small cell lung cancer, but not in non-small cell lung cancer

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Abstract. Lung cancer is the most common cause of cancer-related death worldwide. Current investigations in the field of cancer research have intensively focused on the 'cancer stem cell' or 'tumor-initiating cell'. While CD133 was initially considered as a stem cell marker only in the hematopoietic system and the nervous system, the membrane antigen also identifies tumorigenic cells in certain solid tumors. In this study, we investigated the human lung cancer cell lines A549, H157, H226, Calu-1, H292 and H446. The results of real-time PCR analysis after chemotherapy drug selection and the fluorescence-activated cell sorting analysis showed that CD133 only functioned as a marker in the small cell lung cancer line H446. The sorted CD133⁺ subset presented stem cell-like features, including self-renewal, differentiation, proliferation and tumorigenic capacity in subsequent assays. Furthermore, a proportion of the CD133⁺ cells had a tendency to remain stable, which may explain the controversies arising from previous studies. Therefore, the CD133⁺ subset should provide an enriched source of tumor-initiating cells among H446 cells. Moreover, the antigen could be used as an investigative marker of the tumorigenic process and an effective treatment for small cell lung cancer.

Introduction

Lung cancer incidence has increased sharply and the disease has become the most common cause of cancer-induced death worldwide (1). Lung cancer consists of two main pathological types. Small cell lung cancer (SCLC) accounts for 13-15% of all lung cancers, but its clinical features tend to be more aggressive than the other type, non-small cell lung cancer (NSCLC) (2,3). Recently, increasing evidence has rapidly accumulated all over the world showing a small subset of

cells in the solid tumors. These so-called cancer stem cells can initiate the tumors with performing unique functions such as, self-renewal, asymmetric division, drug resistance and quiescence.

CD133, a highly conserved protein, has been proven and generally accepted as a marker for tumor stem cells in leukemia and glioblastoma (4,5). This membrane antigen also contributes to the identification of the tumorigenic cells in some solid tumors, including prostate cancer, hepatocellular carcinoma and colon carcinoma (6-9).

With regard to lung cancer, both clinical analyses and laboratory studies have shown that CD133 plays a critical role in tumorigenesis in SCLC and NSCLC (10,11). However, this theory is still controversial because CD133 is not a reliable tumor stem cell marker in lung cancer according to some studies, especially in NSCLC (12,13). Therefore, we attempted to demonstrate the role of CD133 in SCLC and NSCLC and explore the possibility of sorting the tumor-initiating cells from lung cancer cells based on CD133 expression.

To exclude any synergistic effects (i.e., those effects resulting from angiogenesis), we worked with pure cell lines in this study. The tumor-initiating cells were initially enriched using chemotherapy agents, followed by analysis of CD133 mRNA expression in drug-selected cells (DSCs). We then performed a fluorescence-activated analysis to study the CD133⁺ subsets in all cell lines and selected the positive fractions. The specific cell lines with a CD133⁺ subset were sorted into a positive subset and a negative subset. With subsequent assays, we analyzed their respective biological and immune capacities. The study was designed to investigate the possibility of isolating tumor-initiating cells with the CD133 antibody and to provide evidence of the role of CD133 in lung cancer.

Materials and methods

Cell lines. Human lung cancer lines included in this study were: A549 (adenocarcinoma); H157 and H226 (squamous cell carcinoma); Calu-1 (epidermoid carcinoma); H292 (mucoepidermoid pulmonary carcinoma); and H446 (small cell lung cancer). Cells were obtained from the Shanghai Life Science Cell Bank of the Chinese Academy of Sciences and the American Type Culture Collection (ATCC). Cells were

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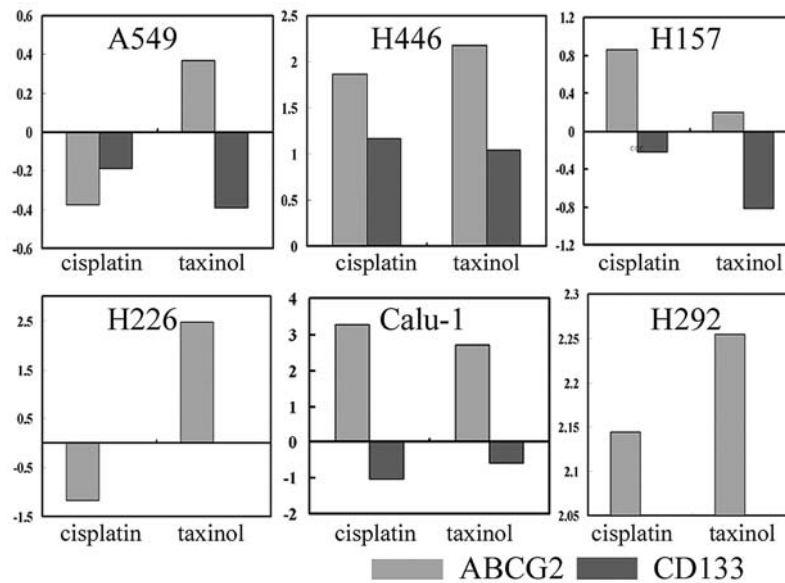


Figure 1. The mRNA expression of ABCG2 and CD133 in drug selected cells. The values were normalized with 18S and converted to the common logarithm. ABCG2 mRNA expression up-regulated in all the respectively lines after treatment with cisplatin or taxinol, but only the CD133 mRNA expressed significantly higher in H446 cell line ($p < 0.05$).

seeded in RPMI-1640 medium (Gibco), as recommended by the ATCC, that was supplemented with 1% penicillin/streptomycin and 10% fetal bovine serum (FBS, Invitrogen-Life Technologies). All cells were incubated in a humidified incubator at 37°C in 5% carbon dioxide.

Chemotherapy drug selection. During the proliferative stage, cisplatin and taxinol were added at a final concentration of 1.25 and 2.5 μM , respectively (based on a preliminary experiment). After 72 h, the living adherent cells (DSC, drug-selected cells), ~50% of the original number, were collected for the subsequent tests.

RNA extraction and real-time PCR analysis. We followed the standard real-time protocol to design the primers into product of <200 bp. The sequences were referred to code AF027208 (AC133 antigen mRNA, CD133, GAAGAGCTTGCA CCAACAAA, AGATGACCGCAGGCTAGTTT) and NM_004827 (ATP-binding cassette sub-family G member 2 mRNA, ABCG2, GTTGTGATGGGCACTCTGAC, CCCTGTAAATCCGTTTCGTTT) in gene bank.

Cells were harvested, and RNA extracted by the TRIzol (Invitrogen). Total RNA was treated with SuperScript III First-Strand Synthesis System on the iCycler PCR system (Bio-Rad) to process the synthesis of first strand followed by cDNA amplification according to the manual. Real-time PCR was operated on the iCycler iQ real-time PCR detection system (Bio-Rad) with SYBR supermix and Taq (Invitrogen) according to the manufacturer's instructions. The programmed thermal cycling conditions were 95°C for 30 sec followed by 45 cycles of 5 sec at 95°C, 34 sec at 60°C. Levels of expression were normalized to the 18S RNA (CGGCTACCA CATCCAAGGAA, GCTGGAATTACCGCGGCT).

Fluorescence-activated analysis and fluorescence-activated cell sorting (FACS). Parental cells were dissociated and

suspended in phosphate-buffered saline (PBS) containing 2% bovine serum albumin and adjusted to a final concentration of 10^7 cells/ml. The specimens were labeled with anti-human CD133-PE antibody (eBioscience, USA) and incubated in the dark at 4°C for 30 min. After washing twice with ice-cold PBS, the cells were sorted with the flow cytometer (BD FACS Calibur, USA). We only found a CD133⁺ subset in the H446 cell line (see Results), therefore the CD133⁺ and CD133⁻ subsets from the H446 cells were collected and cultured for the subsequent tumorigenesis assays. Cells were also stained with isotype control antibody (mIgG2b-PE, Caltag Laboratories) and analyzed by flow cytometry. We followed the same protocol described above to analyze the expression of ABCG2 in cell groups with purified mouse anti-human ABCG2 (BD Pharmingen™, USA) and stained with PE-labeled goat anti-mouse IgG (BD Pharmingen).

Tumorigenicity assay

Growth curve. The sorted positive subset, negative subset and unsorted parental cells were cultured in 96-well plates, with 10^3 cells per well. We assessed three wells from each group and performed the assay daily, and recorded the average data. The growth curve graphs were generated according to cell counts versus time.

Invasion assay. Cellular potential for invasion was determined for each subset of cells using 24-well plates and 8- μm Matrigel invasion chambers (Corning). The chambers were coated with Matrigel, and the cells were dissociated and resuspended in RPMI-1640 medium containing 0.5% bovine serum albumin. Cells were seeded into upper chambers at 1×10^5 cells per chamber. Lower wells were filled with DMEM containing 10% FBS as a chemoattractive agent. Cells were incubated at 37°C with 5% carbon dioxide for 48 h, and then non-invading cells were removed by swabbing the top layer of Matrigel. The membranes containing invading cells were fixed with methanol followed by staining with methyl violet.

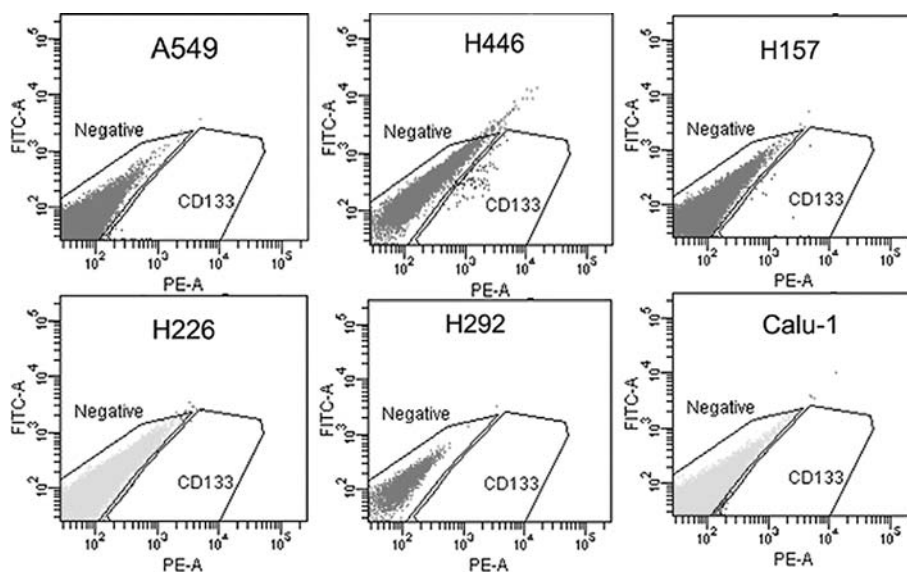


Figure 2. The CD133 analysis in the cell lines. The proportion was 0.4% in H446, and <0.1% in the other 5 cell lines.

The membranes were then washed and mounted on slides. The invading cells were counted under a light microscope.

In vivo assay. We performed the *in vivo* experiment by following our institutional guidelines for the use of laboratory animals. The sorted positive subset, the negative subset and the parental cells were subcutaneously injected into male nude mice (CByJ.Cg-Foxn1 nu/J). The mice were supplied by the National Resource Center for Mutant Mice in Nanjing University of China at three weeks of age. The mice were inoculated with 1×10^3 , 5×10^3 , 1×10^4 , 5×10^4 , 1×10^5 and 5×10^5 cells, respectively, and three mice were inoculated per group. When the xenograft tumors grew to 8 mm, or after 6 weeks of inoculation, the tumor was resected under intraperitoneal anesthesia. The tumor tissues were embedded in paraffin and sectioned, followed by H&E staining. The final slides were viewed with a microscope. The positive subset cells were also analyzed with the CD133 antibody by fluorescence cytometry.

Statistical analysis. Data are presented as the mean \pm SD. The data were analyzed with SPSS 17.0 software. T-test and ANOVA were used to compare differences between groups. $p < 0.05$ was considered statistically significant.

Results

Increased mRNA expression in the drug-selected cells. The CD133 mRNA (Prominin-I) was expressed at very low levels in the cell lines. Nevertheless, the level of CD133 mRNA varied after treatment with cisplatin and taxinol. Only H446 cells showed a significant increase in CD133 mRNA expression after treatment. However, the ABCG2 mRNA was expressed at much higher levels in all six treated cell lines compared to untreated cells (Fig. 1).

Isolation and differential characterization of the CD133⁺ H446 cells. We analyzed CD133 protein expression in various cell lines by flow cytometry and found that the positive rate was $0.883 \pm 0.491\%$ in H446 cells (Figs. 2 and 5), but the positive

subset had nearly disappeared by the 15th generation (more than one month). No positive fraction was found in the other five cell lines (Fig. 2). At the same time, ABCG2 expression, in the H446 CD133⁺ subset, was much higher than the parental cells. The positive and negative subsets were sorted by labeling with the anti-human CD133 antibody and cultured in 96-well plates under identical conditions to compare their repopulation capacity. According to the above results in H446 cells, the DSCs of this line were analyzed as well. This assay demonstrated a similar level of CD133 expression and a higher level of ABCG2 expression in these DSCs as the parental H446 cells.

We also re-analyzed both subsets within two weeks. The positive proportion of the sorted positive subset of the cells had decreased from 100 to 18% after 7 days with a similar reduction in ABCG2 expression. The proportion declined to 2.1% by the end of day 14 (Fig. 5). In contrast, the positive proportion of the negative-sorted subset had risen from zero to 1.0% after 14 days. Since the CD133⁺ cells regenerated both CD133⁻ cells and CD133⁺ cells, the result indicated that the CD133⁺ cells had renewal and differentiation capacities.

The CD133⁺ subset has a greater capacity for proliferation than the CD133⁻ subset. In the same medium of a 96-well plate, the respective proliferative ability of the two subsets and the parental cells were investigated via growth curve analysis (Fig. 6B). Based on the cell counts versus time graph, the parental cells initially multiplied faster than both subsets. After several population doublings, the parental cells and both subsets entered an exponential growth period. As expected, the CD133⁺ subset showed greater proliferative ability than the negative subset ($p=0.011$). However, the number of positive subset cells only slightly exceeded the parental cells after the 10th day of seeding, without a statistically significant difference ($p=0.129$). Our observation period lasted two weeks, and the multiplication rate (shown as the percent of the slope in the graph) began to decline after the 12th day due to contact inhibition.

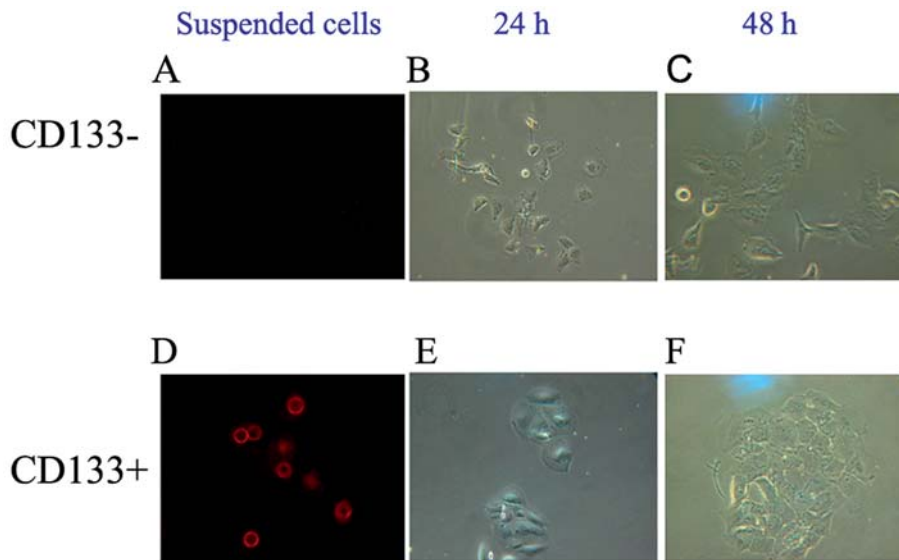


Figure 3. Photographs of sorted CD133⁺ and CD133⁻ cells in H446. The images of suspended CD133⁻ (A) and CD133⁺ (D) cells in H446 were acquired by a fluorescence microscope. There were more streak impurities (arrows) and fragments attaching the cells in CD133⁺ cell cultures (E and F) than in CD133⁻ cultures (B and C). Later, the impurities were washed off by PBS when the medium was changed. The negative cell clones proliferated much faster than the positive after sorting for 24-48 h.

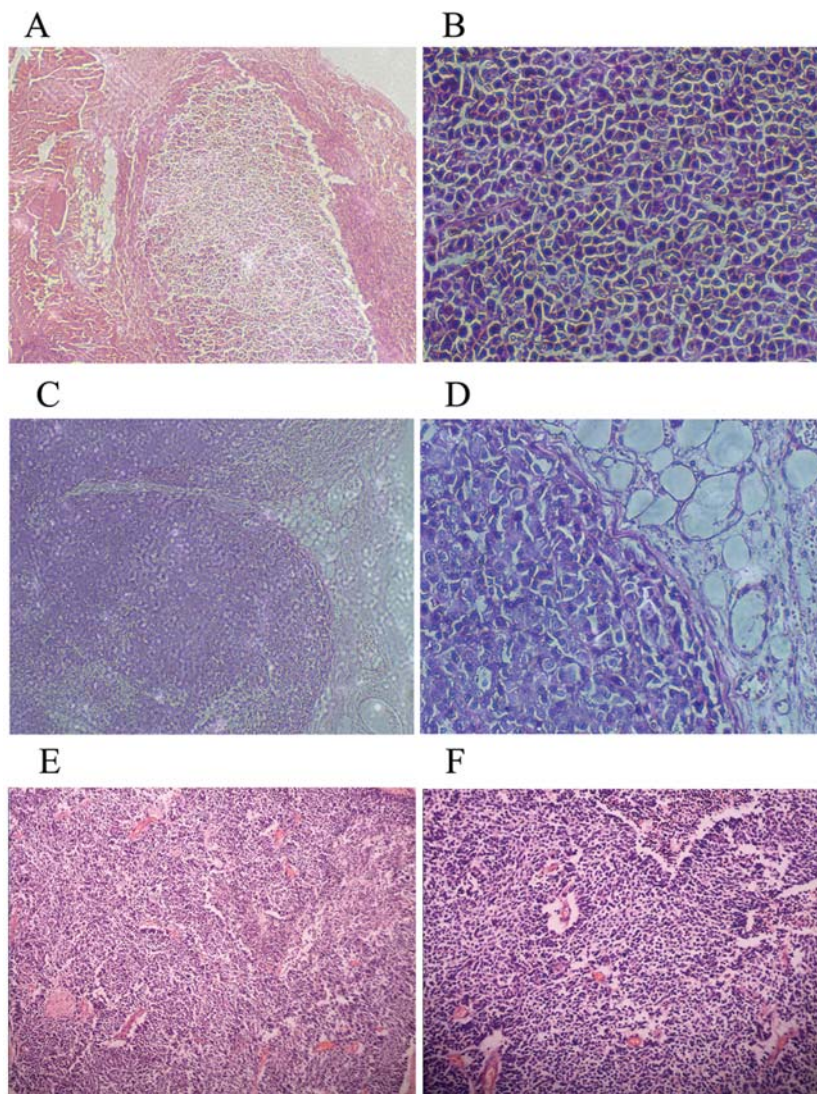


Figure 4. Histopathology of nude mouse xenografts and human small cell lung cancer. Both CD133⁻ (A and B) and CD133⁺ (C and D) nude mouse tumor lost the appearance of typical human small cell lung cancer (E and F) and displayed the characteristics of undifferentiated tumor cells.

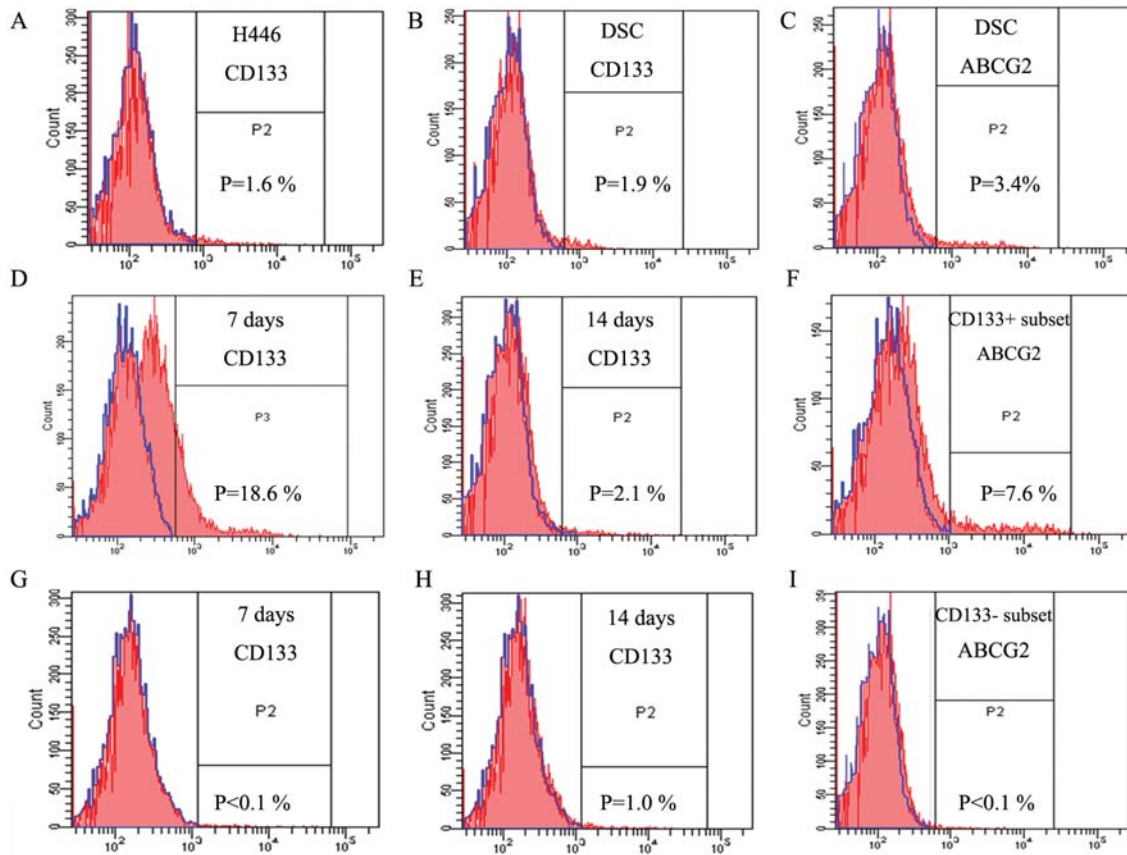


Figure 5. The variation of CD133 and ABCG2 expression in H446 (the outline of control was drawn in blue line to exclude the control result). The CD133⁺ proportion of H446 was $0.883 \pm 0.491\%$ (A), and the CD133⁺ proportion of H446 drug selected cells (DSC) was $0.933 \pm 0.565\%$ (B), which equaled to the parental cells ($p=0.504$; >0.05). The expression of ABCG2 in H446 DSCs: the proportion of ABCG2 expression was 3.4% (C). The CD133 expression in sorted positive subset did not remain stable: the proportion had dropped to 18.6%. After 14 days, it was only 2.1% (D and E). On the contrary, the proportion rose from 0.1 to 1.0% in the CD133⁻ subset after 14 days (G and H). The ABCG2 expression in both sorted subsets after 14 days: the proportion of sorted CD133⁺ subset was 7.6%, and the proportion of the CD133⁻ subset was $<0.1\%$ (F and I).

Increased invasive ability and greater *in vivo* tumorigenic potential of CD133⁺ cells. In the invasion assay, we counted the number of cells within the membrane. We found that over 3 times more CD133⁺ cells ($n=1258 \pm 582$) had traversed the Matrigel than CD133⁻ cells ($n=380 \pm 200$), and the positive cells were ~ 1.2 times more than the parental cells ($n=1045 \pm 377$). There was a statistically significant difference between the positive subset and the negative subset ($p < 0.01$), but there was no significance between the positive and parental cells ($p=0.331$). Therefore, the positive fraction showed more active invasive ability than the negative fraction (Fig. 6A).

The tumorigenic potential of CD133⁺ cells was so distinct compared with the other groups that tumors could form using only 5×10^3 of these cells (2/3 mice) in immune-deficient nude mice within 3 weeks of hypodermic injection (Fig. 6C). Even so, it is surprising that the hypodermal tumor never expanded >2 mm after having emerged. We did not plant $>10^5$ positive cells for two reasons: 1) The culture time was too long to obtain that number without a conclusive higher CD133 expression (proven by the CD133 analysis in the 7th day and 14th day), and 2) Since all these cells are malignant, 10^5 cells are generally enough to form a tumor, that is why injecting $>10^5$ cells had no significance in comparison to tumorigenesis between tumor cells. The assay showed that the CD133⁻ subset had less tumor-initiating capacity (Table I)

because this subset could only form a tumor when 10^5 cells were injected (1/3 mice). This number is at least 20-fold more than the positive cells. We resected the xenograft tumor derived from the positive subset and reanalyzed CD133 expression. We found that the positive proportion had returned to normal levels (Fig. 6D), which was equal to the level seen in the parental H446 cells. The histopathology of the xenografts is shown in Fig. 4. Although there was no difference between the positive tumors and negative tumors, the tumor cells were obviously discernable from the normal human SCLC cells under a magnification of $\times 200$.

Discussion

In the present study, we only found a CD133⁺ subset in the H446 SCLC cell line. The CD133⁺ subset possessed a greater tumorigenic potential than the CD133⁻ subset in the *in vivo* assay. The positive fraction could differentiate into positive and negative fractions as asymmetric division. Furthermore, compared with the negative subset, the positive subset displayed a more aggressively invasive and proliferative ability, as shown with the proliferation assays. The above results demonstrated that the tumor stem cells could be enriched in the CD133⁺ cells, in accordance with the current standards to cancer stem cells (14).

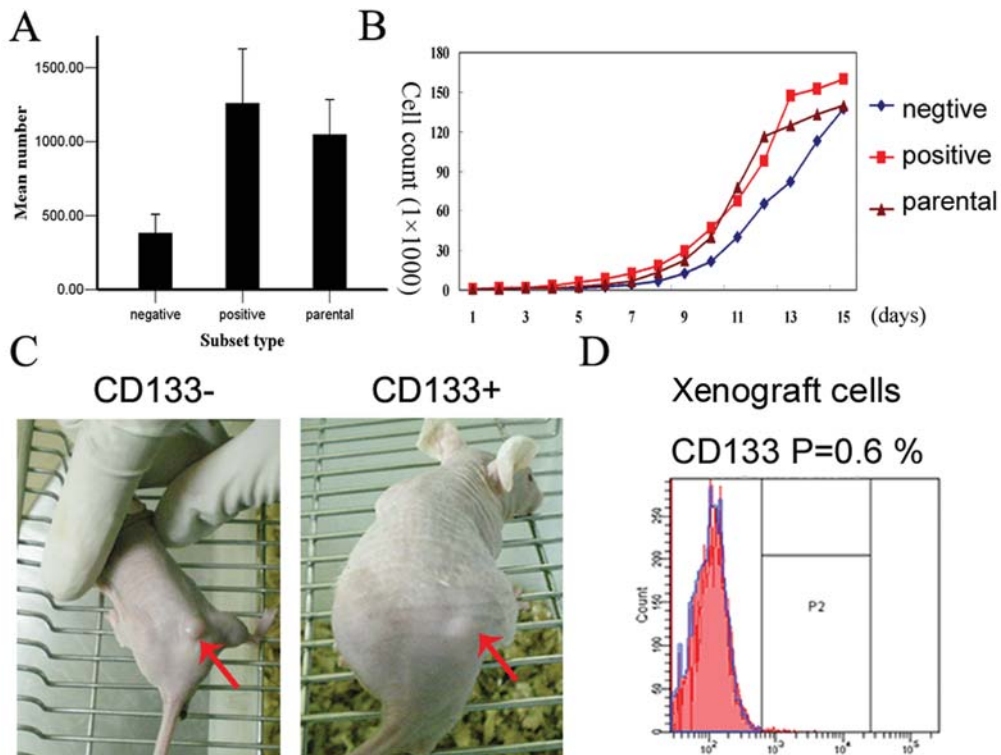


Figure 6. Tumorigenic features of the *in vitro* and *in vivo* assays. In the invasion assay, the data (A) showed CD133⁺ subset traversed more than both CD133⁻ subset ($p < 0.01$), therefore this subset had more active invasive ability. However, there was no difference between positive subset and parental cells ($p = 0.33$; > 0.05). (B) The growth curve graph, the parental cells multiplied faster than both subsets at first because of the cell injury of sorting process. Then the CD133⁺ subset proliferated significantly faster than the CD133⁻ subset during the whole observation period ($p < 0.05$). Even the CD133⁺ subset had exceeded the parental cells, but the slope began to decline after 12 days. Groups came to plateau after the population after 1.4×10^5 . (C) Just 5×10^3 CD133⁺ cells could grow to a tumor (arrow) in the nude mouse, however, it needed 10^5 parental cells to form a tumor compared with the former. The CD133 expression of nude xenograft tumor cells was 0.6% (D).

Table I. The result of *in vivo* assay.

	Cell number for injection (tumor diameter, mm)					
	10^3	5×10^3	10^4	5×10^4	10^5	5×10^5
CD133 ⁺		2 mm (2/3)	2 mm (3/3)	2 mm (3/3)	N/A ^a	N/A
Parent	-	-	-	-	6 mm (3/3)	8 mm (3/3)
CD133 ⁻	-	-	-	-	4 mm (1/3)	6 mm (3/3)

^aExplained in Results.

In contrast, we did not find a CD133⁺ subset in NSCLC cell lines such as A549, H157, H226, H292 and Calu-1. Therefore, CD133 should not be considered a reliable tumor stem cell marker in NSCLC.

The lung cancer stem cell hypothesis is controversial. Some believe that these so-called stem cells come from bone marrow stem cells or embryonic stem cells (15-17), while others insist that they may derive from bronchoalveolar progenitor cells (BASCs) (18). SCLC and NSCLC, the two subtypes of lung cancer, are completely different from one another in terms of tumor biological behavior, histological appearance and treatment. SCLC may be more like a systemic disease that affects the respiratory system rather than a

respiratory disease, especially in the advanced stages. All these results suggest that the sources of progenitor cells in SCLC and NSCLC are not the same, necessitating two separate methods for isolating the tumor-initiating cells, but this hypothesis has not been confirmed. There is also some evidence that NSCLC may be related to BASCs but not to CD133⁺ cells (13,18). The abnormal expression of CD133 in clinical NSCLC patients may not be due to an independent effect but rather to some synergetic effects.

It has not been confirmed whether the cancer stem cells in SCLC come from bone marrow stem cells or embryonic stem cells. CD133 was initially identified as a stem cell marker in the hematopoietic system. It is also found among immature

cells in some human organs, such as the bone marrow and spleen (19). The role of CD133 in leukemia is undoubtedly important. Since some chemotherapy protocols are similar between leukemia and SCLC, understanding the role of CD133 in SCLC may be valuable. Hayashi *et al* reported on the existence of a CD133⁺ subset in two SCLC lines, NCI-H82 and NCI-H69, with higher tumorigenicity than the CD133⁻ subset (20). Also, according to our present results, the CD133⁺ subset only was found in the SCLC cell line. We found only the CD133 mRNA up-regulation after the treatment with cisplatin and taxinol in the H446 SCLC line, whereas the ABCG2 mRNA was overexpressed in all drug-selected lines.

ABCG2 is one of the confirmed markers in a wide variety of stem cells (21), and several studies have shown that the co-action of ABCG2 and CD133 may induce tumor chemoresistance (22,23). The drug selection process enriches stem-like cells in tumors (24). Theoretically, although the cancer stem cell hypothesis comes from the definition of human stem cells, clinicians are interested in it because some particular fraction of cells can survive treatment with cytotoxic agents. The existence of cancer stem cells explains why chemotherapy cannot completely eliminate solid tumors. Thus, chemoresistance should be one of the features of the so-called cancer stem cells. Some studies have suggested that this specific fraction possesses this function for two reasons: first, that there are some membrane transport proteins, such as ABCG2, that can transport the chemotherapeutic agents out of the cell, and secondly due to quiescence (discussed below). Thus, according to our results, the co-increase in CD133 mRNA and ABCG2 mRNA in H446 drug-selected cells implies that CD133 may play a role in the enriched tumor-initiating cells, but not in other cell lines (A549, H157, H226, H229 and Calu-1). Although we cannot yet explain why normal CD133 protein expression is maintained in the drug-selected H446 cells (Fig. 5B), this result may be logically considered as a sign of the induction or enrichment of immature cells.

We also found some additional interesting phenomena related to so-called quiescence. Although the correlation between CD133 and quiescence has not been confirmed, Dooley and colleagues have demonstrated that CD34 expression increases as CD34⁻ cells shift from quiescence to proliferation (27). Since CD133 cells are regarded as a subset of the CD34⁺ cells, the role of CD133 in quiescence is likely to attract attention. Based on our results, some interesting phenomena were found, which might be related to quiescence.

A small number of positive cells were present in the sorted negative subsets after several generations, and their proportion rose to the level of the positive ratio of the parental cell population. This phenomenon suggested that the potential CD133⁺ cells may be latent in the negative fraction or that quiescent CD133 cells may be revived after micro-environmental changes. On the contrary, the proportion of CD133 cells in the sorted positive subset was reduced to normal level at the same time, which indicated that the positive ratio tended to remain stable. The positive cells decrease in proportion due to the asymmetrical division, which means that one cancer stem cell can divide to produce one cancer

stem cell and one mature tumor cell (14). Therefore, their reduction makes the cells generally appear quiescent.

The xenograft tumors generated from the CD133⁺ subset could not expand infinitely in the *in vivo* assay. As few as thousands of cells were enough to form a tumor up to 2 mm in diameter within 3 weeks. However, the xenograft tumor could not expand beyond this size, whereas the unsorted parental cells could form a tumor up to 6 mm in diameter upon the injection of 10⁵ cells. We can also interpret this phenomenon as a general down-regulation of the CD133⁺ subset because the positive rate of CD133 expression in the CD133⁺ nude mouse xenograft tumors was 0.6% (Fig. 6D). This result indicated that the proportion of CD133-positive cells had decreased significantly within six weeks, even in an *in vivo* microenvironment.

The above results suggest that CD133 may be a temporary cancer stem cell marker that is not always expressed throughout the tumor proliferation process. Even in normal organs, CD133 is rapidly down-regulated and disappears completely during the differentiation and maturation period (25). Hence, it is typically quiescent, which may contribute to drug resistance (26). When the quiescence is broken by certain factors, the CD133 mRNA is up-regulated, and the protein will be expressed, after which it will play a role in some unknown signaling pathway. If the microenvironment later reverses to a more stable condition, the active tumor-initiating cells will again become dormant and down-regulate CD133 mRNA. Our results can be illustrated with this hypothesis: The status of quiescence makes CD133⁺ cells more like stem cells than the negative one. Based on the above points, CD133 may play a role in the tumor initiation process rather than the tumor maintenance process. Thus, we refer to its cancer stem cell marker status as temporary.

Some other controversies involving CD133 can also be resolved if the role of CD133 in quiescence can be confirmed. For example, Shmelkov *et al* found that both CD133⁺ and CD133⁻ metastatic tumor subpopulations can form colonies in *in vitro* cultures and were capable of long-term tumorigenesis in a NOD/SCID serial xenotransplantation model. Moreover, metastatic CD133⁻ cells form more aggressive tumors than the positive cells and express typical phenotypic markers of cancer-initiating cells (28). We consider all of these results to be reliable and reasonable as long as the point mentioned above is accepted. *In vivo* and *in vitro* assays take at least two weeks, which is long enough to stimulate the up-regulation of CD133 after quiescence or to down-regulate it from the active situation. We believe these dynamics explain why there is almost no difference between the positive and negative xenografts in terms of pathologic tumor morphology. From our study, we can see that the variations in CD133 expression tend towards stability not only in the positive subset but also in the negative subset. The malignant cells are likely to lose some stem cell-like features when the proportion of CD133⁺ cells decreases. Thus, in our assay, the tumors generated from CD133⁺ cells could not grow any larger after three weeks. This area of study will require further investigation to determine the causes of this phenomenon.

In accordance with the opinion that CD133 is a temporary marker, any future targeted therapies directed against CD133

should be considered for the maintenance of the quiescent status of the malignant cells (particularly including the latent CD133⁺ cells) instead of merely eliminating certain cells. This treatment approach may transform cancer into a chronic disease, much like hypertension and diabetes.

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