REVIEW

Properties and feasibility of using cancer stem cells in clinical cancer treatment

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ABSTRACT

Cancer treatment failure, drug resistance, or metastatic recurrence are thought to be caused mainly by the existence of a very small number of cancer stem cells (CSCs). The characteristics of this subgroup of cells include self-renewal, tumorigenesis, multiple differentiation and high invasiveness, metastasis, and drug resistance potential. Many studies have demonstrated that CSCs play important roles in tumor growth, spread and metastatic relapse after treatment, and are closely related to the prognosis of patients. From a therapeutic viewpoint, deep insights into the CSCs biology, development of specific therapeutic strategies for targeting CSCs, and characterization of their microenvironment could be an ideal way to combat cancer.

KEYWORDS

Cancer stem cells; cancers; treatment failure; metastasis; cancer therapy

Introduction

As one of the most serious diseases that afflict humans, cancer accounts for 15% of global mortality each year. Furthermore, 90% of patients with solid tumor die of metastasis, which consists of multiple processes associated with many factors, including epithelial-mesenchymal transition (EMT) and cancer stem cells (CSCs)1. Even in a hematopoietic system, where the cancers can be eradicated, cancer relapse always occurs2. Although patients are diagnosed as disease-free after cancer treatment, cancer frequently recurs several months or years later because of the redifferentiation and colonization of a small group of CSCs. Thus, at present, CSCs are considered as the cause of tumor initiation, relapse, metastasis, and drug resistance.

The exploration of CSCs can be traced back to the experiment on teratomas in the 18th century. At present, CSCs are widely considered as the root of tumors. Hematopoietic malignancies are the focus of basic research and clinical trials on CSCs. However, over the past several decades, with respect to stem cell biology, an effective and satisfactory cure for cancer has not been developed yet. From this perspective, the complexity of CSCs is considered as the main barrier for curing cancer. Thus, the origins, functions during tumor metastasis, and differentiation of CSCs should be revealed with urgency. Cutting-edge studies on CSCs have already progressed from seeking and screening related biomarkers to placing greater emphasis on the mechanism of molecular regulation and clinical transformation.

Until today, some CSCs biological features, which are regarded as distinct from the group’s parental counterparts, continue to be revealed. CSCs always remain quiescent to suppress their own outgrowth and to maintain their long-term survival until the stress from extra- or intra-tumor cells interferes and induces them to enter the cell cycle from G0/G1 phase to aberrant rapid proliferation3,4. Moreover, some signal pathways, such as phosphatidylinositol 3-kinase (PI3K)/AKT5, interleukin 6 (IL6)/stat36, and Wnt/β-catenin7, which are closely associated with drug resistance, are abnormally activated. After CSCs or non-CSCs transform via an EMT and enter into circulation to become circulating tumor cells (CTCs), many proteins are ectopically expressed for the purposes of anti-apoptosis, immune escape, and even dormancy8,9. Currently, many clinical trials and pre-clinical experiments that target CSCs are ongoing. Molecular and pharmacological targeting of proline-rich tyrosine kinase 2 specifically induces cell death in aldehyde dehydrogenase (ALDH)-positive myeloma CSCs10. However, CSCs could not be completely destroyed, and they allow the regrowth of tumor because of uncovered complicated characteristics.
Therefore, CSCs are the cause of cancer treatment failure; on the other hand, CSCs could be promising clinical targets for improving future cancer therapy. In this review, we discuss the role of CSCs in seeding tumor and the possible strategies for using them in clinical cancer treatment (Figure 1).

Overview of CSCs

The theory on CSCs was proposed in 1967 by Bergsagel, who discovered that leukemic stem cells might exist in Philadelphia chromosome positive chronic myelogenous leukemia (CML). This observation challenged the traditional Darwinian “clonal selection theory,” which advocates that tumor cells are derived from a single cell in the subclone. During the past decades, more researchers have mentioned that when a group of sub-clonal cells are on the verge of extermination, about 5% of these cells narrowly escape apoptosis and acquire more malignant phenotypic properties such as tumorigenesis, metastasis, self-renewal, and drug resistance. Eventually, in 1997, Bonnet et al. ascertained for the first time the existence of CSCs with the marker for CD34+/CD38- in acute myeloid leukemia (AML). In 2003, Al-Hajj et al. successfully obtained CSCs from breast carcinoma by using the marker for CD44+/CD24-. Thereafter, CSCs were found in other kinds of solid malignancies, including those of the brain and the liver.

Although increasing evidence has indicated the existence of CSCs, the origin of CSCs remains unknown. At present, three hypotheses are used to explain the origin of CSCs. First, adult stem cells or progenitor cells from normal tissues become CSCs because of somatic mutation accumulation at the genetic or epigenetic level, or differentiated tumor cells become CSCs through EMT. This hypothesis attracts more researchers to exert effort on discovering the differences between CSCs and transformed cells. Learning about such differences may help modify current cancer treatments more precisely or even develop a completely effective cure for cancer. Second, the differentiated cell and tissue stem cell fuse into a euploid cell. The euploid cell has an unstable genome, which is prone to mutating, and it can finally transform into CSCs. This phenomenon is a rare event, but it might contribute to tumor progression. Third, the normal cells absorb exogenous DNA from nearby tumor cells and integrate the genetic material into their own genomes, resulting in the suppression or ectopic expression of some genes and ultimately leading to the transformation of CSCs. This process is not easily recognized. Given the diversity of the hypotheses, the origin of CSCs still remains a fundamental issue to be addressed.

CSCs: controversial biomarkers in solid tumor

The CSCs hypothesis contradicts the classical theory of clonal selection in which only CSCs is considered as the origin and seed of tumors. However, the heterogeneity of CSCs is universal. Thus, selecting a single specific CSCs biomarker for a certain type of cancer is arduous. For example, CD138, ALDH1+, and side population cells have been used to identify multiple myeloma stem cells. CD133+, CD24+, CD90, and CD47 are all approved to be the markers in hepatic CSCs; leucine-rich repeat-containing G-protein coupled receptor 5 (Lgr5), CD133, and CD44 are used in colonic carcinoma. These markers indicate that distingui-
The communication between CSCs and the microenvironment in primary tumor affects the stemness factors or pathways facilitating the invasiveness and metastatic abilities of CSCs. After the formation of pre-metastases, some attractors could further guide CSCs to go “home” to specific distal tissues. Finally, the changing metabolism of CSCs and the cross-talk of the cells with the new environment during the formation of metastatic cells also regulate the self-renewal and settlement of CSCs. The researchers aim to find intrinsic driving factors within these processes and to perform more effective targeted therapies. In hepatocellular carcinoma, when CD44+ CSCs are co-cultured with CD14+ tumor-associated macrophage (TAM), their migration and tumorigenic abilities are enhanced by IL6 from TAM. In breast cancer, when CD90+ cells interact with TAM, EphA4 is upregulated to modulate the activity of CD90+ breast carcinoma cells. In addition, the exosome, which is secreted by mesenchymal cells in the primary tumor of pancreatic cancer, reaches the targeted organ in advance to remodel the microenvironment, thereby facilitating the adaptability of metastatic pancreatic cancer cells and elevating their survival rate.

Treatment strategies that target CSCs

After cytotoxic treatments, such as chemotherapy, radiotherapy, or targeted therapy, the tumor initially shrinks or disappears; however, relapse may occur and cause the death of cancer patients. The theory on CSCs tries to identify the reason for this phenomenon. CSCs are considered as the “beating heart” of tumors. Thus, possibly, only by exterminating CSCs could cancer be cured. Actually CSCs can be targeted through various strategies, such as by inhibiting the molecular pathway(s) of CSCs, targeting CSCs or the microenvironment, combining CSC-targeted therapy with chemotherapuy, or differentiating inducers to promote the differentiation of CSCs into bulk tumor cells, which can be eradicated by chemotherapy.

Targeted treatments always focus on the cell surface or abnormally activated signaling pathway of CSCs or the cells in the niche. CD133, CD135, and CD24 could be targets for eliminating CSCs. The embryonic signaling pathways always abnormally activate CSCs, which are inhibited by different inhibitors, such as cyclopamine, the prototype of Hedgehog (Hh) pathway-specific inhibitors; iCG001, which selectively inhibits Wnt/β-catenin signaling; and γ-secretase inhibitors, which block notch signaling. Adoptive therapy with ALDH1 family member A1 (ALDH1A1)-specific CD8+ T cells eliminates ALDH1A1-specific CD8+ T cells, which have the
characteristics of cancer-initiating cells. The curative effect of classical treatments may be increased when combined with the targeted therapy of CSCs. CSCs resist drugs because of their quiescent state, whereas most drug treatments target proliferating cells. This characteristic suggests that stimulating CSCs to differentiate may increase their sensitivity to drugs. Oncostatin M induces EpCAM+ liver CSCs to differentiate into a hepatocytic lineage, causing the CSCs to be more sensitive to 5-fluorouracil.

In a hematological system, given that CSCs were first identified in AML in 1994, eradicating the leukemia stem cells (LSCs) has been regarded as the optimal way of decreasing the possibility of relapse. Similar to the solid tumor, LSCs can be targeted with two kinds of treatment strategies, such as by interfering in the abnormal signal pathways or by targeting the specific surface biomarkers of LSCs. Along with their unique characteristics, distinct alterations in many signal pathways, such as Wnt, Notch, NF-κB, and PI3K-AKT, which play important roles in cell self-renewal, proliferation, motility, stemness maintenance, and survival, have been identified in LSCs compared with their normal corresponding hematopoietic stem cells (HSCs). Some inhibitors, such as asbortezomib, parthenolide, and rapamycin, which target abnormally activated pathways, have been developed. Bruton tyrosine kinase (BTK), which could govern or at least participate in Wnt/β-catenin and AKT signaling in myeloma stemness, is not expressed in normal plasma cells. Thus, the BTK inhibitors, including ibrutinib or CGI1746, are promising therapeutic candidates. However, the major concern regarding these inhibitors is their potential toxicity to the normal hematopoietic system because these pathways are also functional in HSCs. Thus, discovering a target that could selectively impair the survival, self-renewal, or proliferation pathways of LSCs is important.

The discovery of the specific surface molecules of LSCs provides some important insights. An ideal therapeutic target should be expressed on LSCs but not on the normal HSCs. Several candidate antigens, such as CD33, CD44, CD47, CD123, and CLL-1, have been identified. The monoclonal antibodies against these markers show more specificity in vitro experiments than the inhibitors of signal pathways. However, the expression levels of certain LSCs surface markers may vary greatly in patients. Even within the same patient, more than one phenotypically distinct subpopulation of LSCs might be found. This information suggests that the cell surface antigen profiles of LSCs may vary. Another challenge in the treatment of hematological malignancies is that targeted drugs cannot reach the appropriate concentration in circulation to induce an effective action. To deal with this problem, the surface markers are combined with nano-delivery systems. The multifunctional delivery using targeting ligands (e.g., hyaluronic acid to target CD44) and other combination therapeutics (e.g., siRNA with small molecular drug) can enhance the treatment efficacy.

Table 1  Treatment strategies for targeting CSCs

<table>
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<tr>
<th>Cancer types</th>
<th>Therapeutic strategies</th>
<th>Examples</th>
<th>Limitation</th>
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<tr>
<td>Solid tumor</td>
<td>Targeting specific surface biomarkers</td>
<td>CD133, CD13, CD24</td>
<td>Provided that CSCs and normal stem cells share many common characteristics, the treatments for targeting CSCs could also have side effects on normal stem cells</td>
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<td>Targeting abnormal signaling pathways</td>
<td>Cycloheximide (11-deoxojervine), the prototype of Hh pathway-specific inhibitors; IC801, which selectively inhibits Wnt/β-catenin signaling; γ-secretase inhibitors, which block notch signaling</td>
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<td>Targeting the stem cell niche</td>
<td>Adoptive therapy with ALDH1A1-specific CD8+ T cells eliminates ALDH bright cells</td>
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<td>Inducing differentiation</td>
<td>Oncostatin M induces EpCAM+ liver CSCs to differentiate into a hepatocytic lineage, making CSCs more sensitive to 5-fluorouracil</td>
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<td>Hematologic neoplasm</td>
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<td></td>
<td>Targeting abnormal signaling pathways</td>
<td>Inhibitor of BTK, ibrutinib, inhibits Wnt/β-catenin and AKT signaling</td>
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CSCs and normal stem cells share many common characteristics, including stemness-related gene expression profiles and capabilities for self-renewal, development into lineages, and proliferation. Thus, the inhibitors that target CSCs could also have side effects on normal stem cells. The side effects may be due to the inaccurate experimental methods that simply regard CSCs as adult stem cells. However, non-CSCs progeny could revert to CSCs under certain conditions, implying the dynamic transition between CSCs and non-CSCs. In addition, multi-lineage differentiation is not a fixed feature of CSCs, because most papers published in recent years have shown that CSCs may only possess mono-potential, which could result in non-CSCs. However, if all of these key points are addressed, specific targeting of CSCs is possible. Robert A. Weinberg, an expert in CSCs-targeted therapy, is taking the risk with treatment through his Verastem Incorporated, which has over 2 trillion dollars in investment and is conducting more than 60 CSCs-related clinical trials.

Problems and prospects

CSCs models have been established for most types of cancer, and related discoveries have revealed important information on them. However, their applications in clinical practices still face many obstacles. First, given the uncertainty of CSCs-specific biomarkers, the criteria for CSCs are not simple and accurate enough. Second, the gold standard for approving CSCs is introducing them to immuno-deficient mice at various doses by injection, which is time-consuming and expensive. Thus, administration is another difficulty in the study of CSCs. Third, the origin of CSCs may vary, and a dynamic transition exists between CSCs and non-CSCs. Therefore, eradicating the existing CSCs does not guarantee that non-CSCs will never transform into CSCs. Furthermore, provided that human cancer is a systemic disease under unbalanced homeostasis, the passive attitudes demonstrate that tumors may not disappear even after CSCs are eliminated. In consequence, although CSCs are promising therapeutic targets, more comprehensive studies are needed to uncover all myths involved in CSCs and to establish a bridge between the progress in basic studies and clinical application in routine cancer diagnosis and treatment of CSCs.

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Conflict of interest statement

No potential conflicts of interest are disclosed

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