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New advances on critical implications of tumor- and metastasis-initiating cells in cancer progression, treatment resistance and disease recurrence

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Summary

Accumulating lines of experimental evidence have revealed that the malignant transformation of multipotent tissue-resident adult stem/progenitor cells into cancer stem/progenitor cells endowed with a high self-renewal capacity and aberrant multilineage differentiation potential may be at origin of the most types of human aggressive and recurrent cancers. Based on new cancer stem/progenitor cell concepts of carcinogenesis, it is suggested that a small subpopulation of highly tumorigenic and migrating cancer stem/progenitor cells, also designated as cancer- and metastasis-initiating cells, can provide critical roles for primary tumor growth, metastases at distant tissues and organs, treatment resistance and disease relapse. Particularly, cancer initiation and progression to locally invasive and metastatic stages is often associated with a persistent activation of distinct developmental signaling pathways in these immature cells during epithelial-mesenchymal transition program. The signaling cascades that are often deregulated in cancer stem/progenitor cells include hedgehog, epidermal growth factor receptor (EGFR), Wnt/ β -catenin, NOTCH, polycomb gene product BMI-1 and/or stromal cell-derived factor-1 (SDF-1)/CXC chemokine receptor 4 (CXCR4). Importantly, the results from recent investigations have also indicated that different cancer subtypes may harbor distinct subsets and/or number of cancer-initiating cells during cancer progression as well as before or after therapy initiation and disease recurrence. Therefore, the identification of the molecular transforming events that frequently occur in cancer- and metastasis-initiating cells versus their differentiated progenies is of immense interest to develop new targeting approach for improving current therapies against aggressive, metastatic, recurrent and lethal cancers.

Keywords

Cancer stem/progenitor cells; Solid tumors; Epithelial-mesenchymal transition; Invasion; Metastases; Treatment resistance; Molecular therapeutic targets; Cancer therapies

Introduction

Major advances in the adult stem/progenitor cell biology have allowed researchers to identify certain specific physiological functions of these immature cells and their early progenies endowed with a self-renewal and multilineage differentiation potential (Kim et al., 2005; Bryder et al., 2006; Mimeault and Batra, 2006; Rizo et al., 2006; Wilson and Trumpp, 2006; Arai and Suda, 2007; Mimeault et al., 2007a; Zhao et al., 2008). The tissue-resident adult stem/progenitor cells with a long longevity generally provide critical roles in the replenishment of cells in homeostatic conditions and after intense injuries along the lifespan of individuals (Fig. 1) (Kim et al., 2005; Bryder et al., 2006; Rizo et al., 2006; Wilson and Trumpp, 2006; Mimeault and Batra, 2006; Arai and Suda, 2007; Mimeault et al., 2007a; Zhao et al., 2008). In counterpart, a growing body of experimental evidence has revealed that an accumulation of genetic abnormalities in tissue-resident adult stem/progenitor cells or their more committed progenies endowed with a self-renewal potential concomitant with the changes in their niches, may result in their malignant transformation into leukemic or tumorigenic cancer stem/progenitor cells (Figs. 1, 2) (Huntly et al., 2004; Bapat et al., 2005; Kim et al., 2005; Rizo et al., 2006; Ginestier et al., 2007; Mimeault et al., 2007b, 2008b; Nicolis, 2007; Nijhof et al., 2007; Vaish, 2007; Aubert and Lansdorp, 2008; Matsui et al., 2008; Mimeault and Batra, 2008a,b; Young et al., 2008; Barker et al., 2009; Zhu et al., 2009). Accordingly with the cancer stem/progenitor cell hypothesis, it is suggested that these immature cancer cells, endowed with a high self-renewal potential and aberrant differentiation ability, and which are also designated as cancer- or tumor-initiating cells, can provide critical roles for primary tumor growth, metastatic spread at distant tissues, resistance to current conventional therapies and disease relapse (Hope et al., 2004; Jamieson et al., 2004; Singh et al., 2004; Bao et al., 2006; Liu et al., 2006a; Ginestier et al., 2007; Hermann et al., 2007; Mimeault et al., 2007a^b; Nicolis, 2007; Eramo et al., 2008; Fillmore and Kuperwasser, 2008; Huang et al., 2008; Matsui et al., 2008; Mimeault and Batra, 2008c; Schatton et al., 2008; Wright et al., 2008; Yang et al., 2008a; Young et al., 2008; Zhang et al., 2008b). In support with these new concepts of carcinogenesis suggesting a major implication of cancer stem/progenitor cells in cancer initiation and progression, the small subpopulations of immature cancer cells with stem cell-like properties, comprising about 0.1–3% of the total cancer cell mass, have been isolated from primary malignant tissues of cancer patients and established cancer cell lines (Al-Hajj and Clarke, 2004; Hope et al., 2004; Jamieson et al., 2004; Matsui et al., 2004; Singh et al., 2004; Fang et al., 2005; Kim et al., 2005; Ponti et al., 2005; Maitland et al., 2006; Salmaggi et al., 2006; Ginestier et al., 2007; Hermann et al., 2007; Mimeault et al., 2007b; Ricci-Vitiani et al., 2007; Prince et al., 2007; Eramo et al., 2008; Schatton et al., 2008; She et al., 2008; Yang et al., 2008a; Zhang et al., 2008a; Huang et al., 2009). The cancer stem/progenitor cells typically expressed several stem cell-like markers including telomerase, aldehyde dehydrogenase (ALDH), CD133, CD44, CXC chemokine receptor 4 (CXCR4), stem cell factor (SCF) receptor KIT, ATP binding-cassette (ABC) multidrug transporters, and/or transcription factors such as OCT-3/4, Nanog and SOX2. The highly leukemic or tumorigenic cancer stem/progenitor cells were able to give rise in vitro and in vivo to the total mass of differentiated cancer cells that recapitulated the complex morphological characteristics and heterogeneous phenotype of original patient's tumors.

In addition, it has also been observed that the cancer progression to locally advanced and metastatic disease stages is usually associated with the inactivation of diverse tumor suppressor gene products and activation of a complex network of oncogenic signaling pathways in cancer cells including cancer-initiating cells concomitant with the changes in their microenvironment (Bapat et al., 2005; Brabletz et al., 2005a; Bao et al., 2006; Haraguchi et al., 2006; Li and Neaves, 2006; Liu et al., 2006a^b; Tso et al., 2006; Gray-Schopfer et al., 2007; Hermann et al., 2007; Katoh, 2007; Nicolis, 2007; Rich, 2007; Sato et

al., 2007; Sengupta et al., 2007; Chiba et al., 2008; Ma et al., 2008; Yang et al., 2008a; Mimeault and Batra, 2009). The stimulation of distinct tumorigenic cascades initiated, in an autocrine or a paracrine manner, through diverse hormones, growth factors and cytokines may contribute to the sustained growth, survival, migration, invasion and treatment resistance of cancer stem/progenitor cells and their differentiated progenies. Particularly, the acquisition of a migratory phenotype by tumorigenic cancer stem/progenitor cells and their progenies during the epithelial-mesenchymal transition (EMT) process concomitant with the changes in the activated stroma may lead to their invasion from primary neoplasms, dissemination through the peripheral circulation and formation of aggressive and metastatic cancers at distant sites (Figs. 1, 2) (Bapat et al., 2005; Brabletz et al., 2005a; Zhou and Hung, 2005; Haraguchi et al., 2006; Thiery and Sleeman, 2006; Tso et al., 2006; Hermann et al., 2007; Mimeault and Batra, 2007b; Moustakas and Heldin, 2007; Shah et al., 2007; Shipitsin et al., 2007; Spaderna et al., 2007; Wang et al., 2007b; Das et al., 2008; Sarkar et al., 2008; Storci et al., 2008; Turley et al., 2008). Consistent with this hypothesis, the tumorigenic and migrating cancer stem/progenitor cells, also designated as metastasis-initiating cells, have been detected at invasion sites in primary tumors as well as isolated from peripheral blood and secondary tumor samples of cancer patients and metastatic cancer cell lines (Galli et al., 2004; Brabletz et al., 2005a; Fang et al., 2005; Ponti et al., 2005; Balic et al., 2006; Patrawala et al., 2006, 2007; Hermann et al., 2007; Wei et al., 2007; Das et al., 2008; Fillmore and Kuperwasser, 2008; Moustakas and Heldin, 2008; Quintana et al., 2008; Schatton et al., 2008; She et al., 2008; Shmelkov et al., 2008; Yang et al., 2008a,b; Aktas et al., 2009). Importantly, the results from numerous recent studies have also indicated that the resistance of leukemic or tumorigenic and migrating cancer stem/progenitor cells to current clinical therapies could result in their persistence at primary and/or secondary neoplasms after treatment initiation (Bao et al., 2006; Liu et al., 2006a; Phillips et al., 2006; Hermann et al., 2007; Liu et al., 2007; Mimeault et al., 2007b, 2008a; Shah et al., 2007; Todaro et al., 2007; Wang et al., 2007a; Chen et al., 2008; Chiba et al., 2008; Fillmore and Kuperwasser, 2008; Friel et al., 2008; Johannessen et al., 2008; Levina et al., 2008; Loebinger et al., 2008; Ma et al., 2008; Matsui et al., 2008; Schatton et al., 2008; Sung et al., 2008; Zhang et al., 2008b). Thereby, cancer stem/progenitor cells can be responsible for the leukemia recurrence or tumor re-growth and disease relapse. In regard with this, we described here the recent investigations undertaken to establish the specific deregulated gene products induced in cancer stem/progenitor cells versus their differentiated progenies during cancer etiology and progression to locally invasive and metastatic disease stages. The provided information should help to design new targeting strategies for eradicating the total cancer cell mass including tumor- and metastasis-initiating cells and improving the current cancer treatments against aggressive, metastatic, recurrent and lethal cancers.

Critical functions of cancer stem/progenitor cells in cancer etiopathogenesis and progression to invasive and metastatic disease stages

Numerous factors may influence the risk of developing a cancer including inherited or somatic DNA mutations, intense oxidative stress, tobacco smoking, environmental carcinogens, radiation exposure, chronic inflammatory and fibrotic atrophies and age of individuals (Kim et al., 2005; Mimeault et al., 2007b; Nijhof et al., 2007; Sato et al., 2007; Vaish, 2007; Aubert and Lansdorp, 2008; Catassi et al., 2008; Friedman, 2008; Gumucio et al., 2008; Mimeault and Batra, 2008b; Mimeault et al., 2008b; Widera et al., 2008). Although the precise etiological causes responsible of cancer initiation remain not precisely established, the cancer development is usually associated with a cumulative genotoxic stress in cells that may cause chromosomal instability leading to the changes in the expression levels and/or activity of many deregulated gene products (Mimeault et al., 2007b; Shiras et

al., 2007; Tchernev and Orfanos, 2007; Vaish, 2007; Zhou et al., 2007; Gumucio et al., 2008; Marusyk and DeGregori, 2008; Widera et al., 2008). In this matter, it is worth mentioning that certain inherited genetic aberrations may notably result in developmental defects and a predisposition to develop certain cancer types in post-natal life (Fig. 1) (MacDonald et al., 2003; Tiffin et al., 2003; Nakagawara and Ohira, 2004; Walton et al., 2004; Tostar et al., 2006; Mannelli et al., 2007; Mimeault et al., 2007b; Park et al., 2007; Ross and Spengler, 2007). For instance, the germinal mutations in PTCH receptor gene leading to an aberrant activation of hedgehog signaling pathway may promote the incidence of embryonal rhabdomyosarcoma, fetal rhabdomyoma, basal cell carcinoma, medulloblastoma and meningioma (Tiffin et al., 2003; Tostar et al., 2006). In the same pathway, the primitive neuroectodermal tumors (PNETs) including neuroblastoma, pheochromocytoma, ependymoblastoma and pineoblastoma, which are particularly manifest in pediatric population, may also originate during the embryonic development from neuroectodermal stem cells such as neural crest stem cells (MacDonald et al., 2003; Nakagawara and Ohira, 2004; Walton et al., 2004; Mannelli et al., 2007; Ross and Spengler, 2007). Although the embryonal origin of certain cancer types, the most types of human cancers appear rather to originate from a sequential and progressive accumulation of genetic abnormalities occurring in tissue-resident adult stem/progenitor cells concomitant with the changes in their microenvironments that lead to their malignant transformation in tumorigenic cancer stem/progenitor cells (Figs. 1, 2) (Mimeault and Batra, 2007a,b, 2008c; Mimeault et al., 2008a,b). These molecular transforming events in adult stem/progenitor cells may promote their sustained proliferation and aberrant differentiation, and thereby disrupt the normal mechanisms of tissue regeneration.

Numerous recent studies suggest that the cancer development may generally derive from the clonal expansion of cancer stem cells (CSCs) and/or their early progenies endowed with a high self-renewal capacity but aberrant differentiation potential that trigger the tumor growth (Mimeault and Batra, 2007a,b, 2008c; Mimeault et al., 2008a,b). In analogy with the normal tissue regeneration process mediated through tissue-resident adult stem/progenitor cells, CSCs can generate, through an asymmetric division, the daughter cells designated as transit amplifying (TA) cells with a malignant phenotype (Figs. 1,2). The malignant TA cells, in turn, can give rise to a heterogeneous population of poorly-, moderately and terminally-differentiated cells with aberrant functions (Mimeault and Batra, 2007a,b, 2008c; Mimeault et al., 2008a,b). Furthermore, the migration of malignant TA cells at distant sites within tissues from which they originate concomitant with the changes in their local microenvironment may result in a populational asymmetry and generation of malignant daughter cells endowed with different phenotypic properties including a self-renewal potential. In spite this importance advance, it will be important to determine the implication of tumorigenic CSCs versus their early progenies endowed with stem cell-like properties in development of specific cancer subsets, and more particularly whether different immature cancer cells may persist along cancer progression and drive tumor development.

In support with the critical functions of leukemic and tumorigenic cancer stem/progenitor cells in cancer development, recent investigations have led to the identification and isolation of different cancer-initiating cells in the most common human cancers and established cancer cell lines (Mimeault and Batra, 2007a,b, 2008c 2008a; Mimeault et al., 2008b). Among the cancer types harboring a subpopulation of leukemic or tumorigenic cancer stem/progenitor cells, there are leukemias, lymphomas, sarcomas, melanoma, brain tumors and a variety of epithelial cancers including skin, head and neck, thyroid, lung, cervical, renal, hepatic, esophageal, gastrointestinal, colon, bladder, pancreatic, prostate, mammary and ovarian cancers (Al-Hajj and Clarke, 2004; Hope et al., 2004; Jamieson et al., 2004; Matsui et al., 2004; Singh et al., 2004; Fang et al., 2005; Kim et al., 2005; Ponti et al., 2005; Maitland et al., 2006; Salmaggi et al., 2006; Ginestier et al., 2007; Hermann et al., 2007;

Mimeault et al., 2007b; Prince et al., 2007; Ricci-Vitiani et al., 2007; Zhong et al., 2007; Eramo et al., 2008; Fillmore and Kuperwasser, 2008; Huang et al., 2008; Schatton et al., 2008; Shi et al., 2008; Wright et al., 2008; Yang et al., 2008a; Yu et al., 2008; Zhang et al., 2008a). It has been shown that the small subpopulations of isolated cancer stem/progenitor cells with stem cell-like properties displayed a greater clonogenic potential *in vitro* and generated leukemias or tumors with a higher incidence as compared to their differentiated progenies in animal model *in vivo*. In addition, the cancer progression is usually associated with the acquisition of a more malignant behavior by leukemic or tumorigenic stem/progenitor cells and their progenies.

Molecular transforming events in cancer stem/progenitor cells and their progenies associated with cancer initiation and progression

The transition from non-malignant hyperproliferative lesions to well established cancers has been associated with the occurrence of some oncogenic events in tissue-resident adult stem/progenitor cells and their microenvironment resulting into their acquisition of a malignant behavior (Li and Neaves, 2006; Mimeault and Batra, 2007a,b, 2008c; Das et al., 2008; Gumucio et al., 2008; Huang et al., 2008; Mimeault et al., 2008a, 2008b; Griffero et al., 2009). The transforming events include the stimulation of telomerase and inactivating mutations in numerous tumor suppressor gene products [p16^{INK4A}, pRb, p53 and/or phosphatase and tensin homolog deleted on chromosome 10 (PTEN)]. Moreover, a constitutive activation of diverse growth factors and oncogenic signaling products [Ras, Myc, NF- κ B, PI₃K/Akt/mTOR, Bcl-2, survivin and/or fusion proteins resulting from chromosomal rearrangements] frequently occurs during cancer progression (Fig. 2) (Mimeault et al., 2007b; Sato et al., 2007; Sharpless and DePinho, 2007; Shiras et al., 2007; Tchernev and Orfanos, 2007; Vaish, 2007; Zhou et al., 2007; Dumont et al., 2008; Huang et al., 2008; Ma et al., 2008; Marusyk and DeGregori, 2008; Mimeault and Batra, 2009). In this regard, a growing body of evidence suggests that a progressive accumulation of different transforming events in tissue-resident adult stem/progenitor cells or pre-cancerous stem/progenitor cells with advancing age could be associated with an enhanced incidence of certain age-related cancer types including some epithelial cancers such as breast, prostate, pancreatic and colorectal cancers (Fig. 1) (Sharpless and DePinho, 2007; Griffero et al., 2009; Mimeault and Batra, 2009). Particularly, an activation of telomerase activity in the pre-cancerous cells could lead to their immortalization while the up-regulated expression and/or activity of diverse hormones, growth factors, cytokines and/or their cognate receptors in these immortalized cells could culminate to cancer development during chronological aging (Sharpless and DePinho, 2007; Griffero et al., 2009; Mimeault and Batra, 2009). In this regard, it has been proposed that the age-related increase in the number of cancer stem celllike markers such as CD44, CD166 and ESA as well as EGFR in macroscopically normal colonic mucosa could be associated with a predisposition to developing colorectal cancer with advancing age (Griffero et al., 2009). Further investigations are necessary to establish the phenotypic changes occurring in tissue-resident adult stem/progenitor cells versus their differentiated progenies during chronological aging that predispose to cancer formation in older individuals.

In addition, the up-regulation of distinct oncogenic signaling pathways in tumorigenic cancer stem/progenitor cells during cancer progression may also contribute to their sustained growth, survival and/or invasion (Bao et al., 2006; Haraguchi et al., 2006; Liu et al., 2006a; Onoue et al., 2006; Tso et al., 2006; Katoh, 2007; Mimeault and Batra, 2007a, b, 2008b, c, 2009; Mimeault et al., 2007b, 2008b; Nicolis, 2007; Sato et al., 2007; Glinsky, 2008; Huang et al., 2008; Moustakas and Heldin, 2008). These signaling pathways include hedgehog, epidermal growth factor (EGF)-EGFR system, Wnt/ β -catenin, Notch, SCF/KIT, hyaluronan (HA)/CD44 receptor, interleukin-4 (IL-4)/IL-4R α , stromal cell-derived factor-1

(SDF-1)/CXCR4, transforming growth factor- β (TGF- β) superfamily cytokines, and/or polycomb group (PcG) proteins. For instance, the stimulation of HA/CD44 signaling cascade may lead to the activation of multiple receptor tyrosine kinases (RTKs) such as EGFR, ErbB2, insulin growth factor 1 receptor- β (IGF1R- β), platelet-derived growth factor- β (PDGFR- β) and/or the c-MET receptor of hepatocyte growth factor (HGF) and up-regulation of distinct anti-apoptotic factors and ABC multidrug transporters (Misra et al., 2006; Gilg et al., 2008; Toole and Slomiany, 2008). Hence, these effects mediated through the stimulation of HA-CD44 axis may promote the survival, invasion, multidrug resistance and/or protection against DNA oxidative damages of cancer stem/progenitor cells and their progenies. In general, the cancer development is also accompanied by an enhanced glycolysis in cancer cells including cancer stem/progenitor cells (Das et al., 2008; Gerlee and Anderson, 2008; Kondoh, 2008; Olivotto and Dello, 2008). This phenomenon known as Warburg effect may contribute to the resistance of cancer cells to oxidative stress as well as their survival in intratumoral hypoxic conditions (Das et al., 2008; Kondoh, 2008; Gerlee and Anderson, 2008; Olivotto and Dello, 2008). Moreover, the EMT phenomenon, which occurs in embryonic stem cells (ESCs) through the tissue and organ morphogenesis and patterning during embryonic development as well as tissue regeneration and wound healing in adult, is also re-activated during the progression of numerous aggressive cancers (Figs. 1, 2). Among them, there are brain, skin, prostate, ovarian, mammary, hepatic, gastrointestinal, pancreatic and colorectal carcinomas (Bapat et al., 2005; Brabletz et al., 2005a; Zhou and Hung, 2005; Ratajczak et al., 2006; Thiery and Sleeman, 2006; Tso et al., 2006; Bailey et al., 2007; Hermann et al., 2007; Mimeault and Batra, 2007a,^b, 2008b,^c; Mimeault et al., 2007b, 2008a,^b; Moustakas and Heldin, 2007; Shipitsin et al., 2007; Wang et al., 2007b; Das et al., 2008; Turley et al., 2008).

Molecular events in tumorigenic cancer stem/progenitor cells and their differentiated progenies during the EMT process associated with the formation of locally invasive and metastatic cancers

The acquisition of a more malignant behavior by tumor-initiating cells and their progenies in primary malignant neoplasms, including a migratory phenotype during EMT process, represents a determinant factor that may contribute to disease progression to locally invasive and metastatic cancer subtypes (Abraham et al., 2005; Bapat et al., 2005; Brabletz et al., 2005a; Zhou and Hung, 2005; Balic et al., 2006; Ratajczak et al., 2006; Thiery and Sleeman, 2006; Tso et al., 2006; Bailey et al., 2007; Hermann et al., 2007; Mimeault and Batra, 2007b; Moustakas and Heldin, 2007; Shah et al., 2007; Shipitsin et al., 2007; Spaderna et al., 2007; Wang et al., 2007b; Das et al., 2008; Mani et al., 2008; Morel et al., 2008; Storci et al., 2008; Turley et al., 2008). The occurrence of EMT program in cancer stem/progenitor cells and their progenies may lead to the changes in their morphology and differentiation including a loss of polarity and epithelial cell markers concomitant with a gain of mesenchymal phenotypes that promote their migratory ability. This process is generally associated with a disruption of cell-cell junctions, loss of contact inhibition and extensive reorganization of the actin cytoskeleton and remodeling of extracellular matrix (ECM) components that lead to an increase of the motile and invasive abilities of cancer cells (Fig. 2). A complex network of different oncogenic cascades initiated through activating mutations in signaling components such as Ras and persistent activation of different growth factor pathways may cooperate for inducing a more complete EMT program in cancer stem/progenitor cells and their progenies (Fig. 2) (Zhou and Hung, 2005; Ratajczak et al., 2006; Thiery and Sleeman, 2006; Tso et al., 2006; Mimeault and Batra, 2007b; Shah et al., 2007; Spaderna et al., 2007; Wang et al., 2007b; Das et al., 2008; Kong et al., 2008; Levina et al., 2008; Morel et al., 2008; Storci et al., 2008; Turley et al., 2008). Among them, there are diverse developmental pathways such as sonic hedgehog SHH/PTCH/SMO/GLIs, EGF/

EGFR/Ras/MAPKs, Wnt/ β -catenin, Notch, bone morphogenic proteins (BMPs), fibroblast growth factor (FGF)/FGFR, TGF- β /TGFR β and PDGF/PDGFR pathways (Fig. 2) (Zhou and Hung, 2005; Ratajczak et al., 2006; Thiery and Sleeman, 2006; Tso et al., 2006; Feldmann et al., 2007; Mimeault and Batra, 2007b; Moustakas and Heldin, 2007; Shah et al., 2007; Spaderna et al., 2007; Wang et al., 2007b; Das et al., 2008; Kong et al., 2008; Levina et al., 2008; Storci et al., 2008; Turley et al., 2008; DiMeo et al., 2009). The sustained stimulation of these growth factor pathways may result in an up-regulation of diverse gene products in cancer stem/progenitor cells and their differentiated progenies during the EMT program. The signaling elements that are frequently altered in cancer cells during the EMT process include a decreased expression of E-cadherin concomitant with an up-regulation of different signaling elements such as N-cadherin, vimentin, tenascin C, NF- κ B, snail, slug, twist, β -catenin, CXCR4 and anti-apoptotic factors (Brabletz et al., 2005a; Zhou and Hung, 2005; Onoue et al., 2006; Ratajczak et al., 2006; Thiery and Sleeman, 2006; Tso et al., 2006; Feldmann et al., 2007; Hermann et al., 2007; Mimeault and Batra, 2007b; Moustakas and Heldin, 2007; Shah et al., 2007; Spaderna et al., 2007; Wang et al., 2007b; Das et al., 2008; Dumont et al., 2008; Sarkar et al., 2008; Storci et al., 2008; Chen et al., 2009; DiMeo et al., 2009; Kurrey et al., 2009). These signaling elements may contribute to the invasive and metastatic phenotypes of cancer cells and enhanced resistance to radiation and chemotherapies.

Consistent with the critical functions of tumorigenic and migrating cancer stem/progenitor cells in invasion and metastases at distant sites, the cancer cells with the stem cell-like properties have been detected at invasive front in primary neoplasms, in blood and metastatic tissues from cancer patients (Brabletz et al., 2005a; Fang et al., 2005; Ponti et al., 2005; Balic et al., 2006; Hermann et al., 2007; Das et al., 2008; Fillmore and Kuperwasser, 2008; Schatton et al., 2008; Shi et al., 2008; Yang et al., 2008a). Particularly, the migrating cancer cells involved in several cancers such as melanoma, brain, breast, ovarian, prostate and pancreatic cancer express the CXCR4 receptor that may promote their invasion and migration to distant sites such as lymph nodes, bones, lungs and/or liver that are characterized by high expression levels of its ligand, SDF-1 (Geminder et al., 2001; Muller et al., 2001; Sun et al., 2003, 2005, 2007; Ratajczak et al., 2006; Hermann et al., 2007). The stimulation of SDF-1-CXCR4 signaling cascade may contribute to activate diverse signaling elements involved in survival, ECM degradation (metalloproteinases) and motility of cancer cells at the primary neoplasm. Moreover, the chemoattractant SDF-1 ligand gradient may promote the migration and adhesion of CXCR4⁺ circulating cancer cells including cancer stem/progenitor cells to endothelium and stromal ECM components at distant metastatic sites by modulating intracellular signaling pathways mediated by adhesion molecules and ECM/integrins (Ratajczak et al., 2004; Hartmann et al., 2005). For instance, the data from immunohistochemical analyses of primary pancreatic adenocarcinoma specimens from patients have revealed the presence of two different subpopulations of pancreatic cancer stem/progenitor cells, including tumorigenic CD133⁺/CXCR4⁻ and migrating CD133⁺/CXCR4⁺ subsets localized in the bulk mass and invasive front of pancreatic tumor, respectively (Hermann et al., 2007). It has also been shown that CD133⁺ cells isolated from patient's tumor tissues were able to form tumors after orthotopic injection in the pancreas of immunodeficient nude mice. Importantly, the depletion of CD133⁺/CXCR4⁺ migrating cancer stem/progenitor cells in the total cancer cell mass constituting highly metastatic human pancreatic cancer cell line L3.6 pl or inhibition of CXCR4 cascade effectively abrogated their metastatic capacity without altering their tumorigenic potential in animal models *in vivo* (Hermann et al., 2007). Hence, these observations suggest that the migrating CD133⁺/CXCR4⁺ cells may correspond to a more malignant cell subpopulation than tumorigenic CD133⁺/CXCR4⁻ cells, which may have acquired a migratory phenotype during the EMT program, and thereby they may be involved in invasion and metastases to distant sites.

In addition, it has been shown that the occurrence of EMT may promote the metastatic dissemination of cancer stem/progenitor cells and their progenies (Abraham et al., 2005; Balic et al., 2006; Morel et al., 2008). For instance, the circulating tumor cells expressing the stem cell-like marker ALDH1 and EMT-associated molecules (twist, PI3K α and Akt2) have been detected in blood samples from metastatic breast cancer patients (Aktas et al., 2009). Importantly, it has also been reported that the putative CD44⁺CD24^{-/low} breast cancer stem cells were detected in all bone marrow patient's specimens from early breast cancer patients expressing cytokeratin (CK⁺) with a high mean prevalence of 72% relative to a low prevalence inferior to 10% of cells in primary tumors (Balic et al., 2006). These data support the concept that the enhanced expression levels of specific gene products, and more particularly EMT-associated molecules, in cancer-initiating cells at primary neoplasms may promote their invasion and metastatic spread at definite distant sites.

Molecular events in the tumor stroma associated with the formation of locally invasive and metastatic cancers

The cancer progression is also accompanied by an extensive tumor stromal remodeling of ECM components and changes in gene expression patterns in activated tumor-associated stromal cells including myofibroblasts and/or stellate cells as well as infiltrating circulating endothelial progenitor cells (EPCs) and immune cells such as macrophages (Fig. 2) (Mimeault and Batra, 2007a,b,2008d;Mimeault et al., 2007b,2008b;Friedman, 2008;Kirkland, 2009). These changes in tumor stroma may contribute to the acquisition of stem cell-like phenotypes, a more malignant behavior and invasive ability by cancer cells. Especially, a variety of motomorphogens and chemoattractant factors may be secreted by stromal cells and promote the migration of cancer stem/progenitor cells and their progenies during the transition from localized cancers to invasive and metastatic disease stages. The soluble growth factors, cytokines and chemokines released by tumor stromal cells in reactive stroma include EGF, insulin-like growth factor (IGF), hepatocyte growth factor (HGF), TGF- β and SDF-1 as well as matrix metalloproteinases (MMPs) and urokinase plasminogen activator (uPA) (Brabletz et al., 2005a;Mimeault and Batra, 2006;Ratajczak et al., 2006;Kleeff et al., 2007;Mimeault and Batra, 2007a,^b,2008b;Mimeault et al., 2007b, 2008b;Spaderna et al., 2007). The release of these soluble factors in reactive stroma may promote, in a paracrine manner, the malignant transformation of cancer stem/progenitor cells and their differentiated progenies during the EMT process (Fig. 2). Moreover, the secretion of diverse angiogenic factors by tumor cells and myofibroblasts may stimulate the tissue-resident endothelial cells or EPCs as well as the recruitment of BM-derived hematopoietic cells including circulating EPCs at tumoral sites that can cooperate to induce the tumor neovascularization process. Hence, these molecular events may promote the invasion of tumorigenic and migrating cancer stem/progenitor cells into reactive stroma, dissemination through the near lymph nodes and peripheral circulation and metastasis at distant tissues and organs.

Importantly, the phenotypic changes of cancer stem/progenitor cells, including a re-differentiation towards the occurrence of mesenchymal-epithelial transition (MET) process, may also occur at certain metastatic tissues and organs including BM (Fig. 2) (Thiery and Sleeman, 2006;Spaderna et al., 2007). Moreover, the immature metastasis-initiating cells like adult stem/progenitor cells may exhibit a quiescent and less metabolically active state in their novel local microenvironment, niches prevalent at secondary anatomical sites. The adoption of a quiescent state by metastasis-initiating cells may explain, at least in part, the long-term dormancy phenomenon. The quiescence of these immature cancer cells may be associated with their persistence under form of micrometastases for a prolonged period of time without clinical or histopathologic signs of apparent macrometastases as well as their resistance to radiation or cytotoxic drugs targeting the proliferative cancer cells (Fig. 2)

(Cleardin and Teti, 2007; Mimeault and Batra, 2007a; Mimeault et al., 2007b; Horak et al., 2008; ix-Panabieres et al., 2008; Mimeault et al., 2008a; Rak et al., 2008; Riethdorf et al., 2008; Trumpp and Wiestler, 2008; Vincent-Salomon et al., 2008). The changes in the local microenvironment of metastasis-initiating cells leading to the repression of metastasis suppressor genes and/or re-activation of mitogenic signaling pathways could however trigger their proliferation, secondary tumor growth and culminate to disease recurrence (Fig. 2) (Cleardin and Teti, 2007; Mimeault and Batra, 2007a; Mimeault et al., 2007b; Horak et al., 2008; ix-Panabieres et al., 2008; Rak et al., 2008; Riethdorf et al., 2008; Trumpp and Wiestler, 2008; Vincent-Salomon et al., 2008).

Collectively, these observations suggest that the tumorigenic and metastatic potentials of cancer stem/progenitor cells may be enhanced along disease progression through a sequential and progressive accumulation of oncogenic transforming events occurring during EMT process. Especially, the acquisition of a migratory phenotype and survival advantages by tumorigenic cancer stem/progenitor cells during the EMT process at primary neoplasm, concomitant with the changes in their local tumor microenvironment may result to their invasion and progression from organ-confined cancers to metastatic disease states (Fig. 2). Hence, the intrinsic and acquired resistance of tumor- and metastasis-initiating cells to current cancer therapies may lead to their persistence at primary and secondary neoplasms and disease relapse. In respect with this, we discussed accumulating lines of evidence suggesting that the acquisition of distinct phenotypes by tumor-initiating cells during cancer progression may lead to the formation of different cancer subtypes that differently respond to current therapeutic treatments.

Heterogeneity of cancers derived from distinct tumorigenic and migrating cancer stem/progenitor cells and their differential response to current clinical treatments

Numerous investigations revealed that the occurrence of different malignant transforming events in adult stem/progenitor cells during cancer initiation as well as an accumulation of distinct genetic and/or epigenetic alterations in cancer stem/progenitor cells along cancer progression may lead to the development of different cancer subtypes (Dontu et al., 2004; Brabletz et al., 2005a; Clarke et al., 2005; Teuliere et al., 2005; Anderson and Matsuno, 2006; Asselin-Labat et al., 2006; Buyse et al., 2006; Laakso et al., 2006; Sorlie et al., 2006; Tang et al., 2006; Tso et al., 2006; Dalerba et al., 2007; Fodde and Brabletz, 2007; Kelly et al., 2007; Kennedy et al., 2007; Langley and Fidler, 2007; Mimeault and Batra, 2007b; Zhou et al., 2007; Agelopoulos et al., 2008; Ben-Porath et al., 2008; Das et al., 2008; Huang et al., 2008; Quintana et al., 2008). This heterogeneity of cancers has important repercussion since these distinct cancer subtypes with variable degrees of differentiation and/or aggressivity may harbor distinct cancer-initiating cells exhibiting different phenotypic and functional properties, and differentially respond to current cancer therapies. More specifically, the expression of embryonic stem cell (ESC)-associated and mesenchymal genes and the acquisition of a migratory phenotype by poorly- or moderately-differentiated tumorigenic cancer stem/progenitor cells during the EMT program may result in the formation of highly invasive and metastatic cancer subtypes characterized by a poorly- to moderately-differentiated state (Abraham et al., 2005; Brabletz et al., 2005b; Anderson and Matsuno, 2006; Asselin-Labat et al., 2006; Balic et al., 2006; Buyse et al., 2006; Mimeault and Batra, 2007b; Ben-Porath et al., 2008; Aktas et al., 2009). In contrast, the tumorigenic cancer stem/progenitor cells that do not undertake the EMT transition could rather give rise to weakly invasive cancer subtypes (Mimeault and Batra, 2007b).

In addition, the changes in the local microenvironment of tumorigenic cancer stem/progenitor cells and their differentiated progenies including their localization within hypoxic zones of solid tumors may result in the expression of a different subset of oncogenic gene products during cancer development and be responsible, at least in part, for the intratumoral heterogeneity (Mimeault and Batra, 2007b; Das et al., 2008). This line of thought is well-supported by the observations indicating that certain invasive cancer types, such as mammary, ovarian, prostate, pancreatic, gastric, colorectal and squamous cell carcinomas, harbored an intratumoral heterogeneity (Galli et al., 2004; Brabletz et al., 2005a; Kalluri and Zeisberg, 2006; Hermann et al., 2007; Mimeault and Batra, 2007a,b, 2008c). These invasive cancer subtypes typically display distinct proliferating and differentiating regions, including a preferential localization of migrating cancer stem/progenitor cells at intratumoral hypoxic zones and invasive front (Fig. 2). In support with this model of heterogeneity of cancers, we are reporting in a more detailed manner, accumulating lines of experimental evidence obtained for breast and brain cancer subtypes underlining the complexity of clinical classification based on specific phenotypic features of cancer cells. The therapeutic significance of the heterogeneity of these cancer subtypes is also discussed.

Heterogeneity of breast cancers

At least five subtypes of breast cancer have been identified by comparing gene expression signatures of cancer cells and their invasive phenotype. The classification of breast cancer includes basal-like (estrogen receptor “ER α ”, progesterone receptor “PR”, erbB2^{-low}, CK5/6⁺ and EGFR⁺); erbB2/HER2⁺ overexpressing (ER α ⁻ and PR⁻); luminal A (ER α ⁺ and/or PR⁺ and erbB2⁻); luminal B (ER α ⁺ and/or PR⁺ and erbB2⁺) and normal breast cancer subtype (high expression of normal epithelium genes and low expression of luminal epithelial gene products) (Clarke et al., 2005; Anderson and Matsuno, 2006; Asselin-Labat et al., 2006; Buyse et al., 2006; Calza et al., 2006; Carey et al., 2006b; Laakso et al., 2006; Lacroix, 2006; Sorlie et al., 2006; Tang et al., 2006; Agelopoulos et al., 2008). More recently, two other subtypes designated as metaplastic and claudin^{low} breast cancers, have also been characterized as triple negative for ER- α , PR and erbB2 (Fig. 2) (Hennessy et al., 2009). The triple-negative breast cancers (TNBCs), encompassing basal-like, metaplastic, claudin^{low} and normal breast cancer subtypes, represent one of the most aggressive and lethal breast cancer subgroup (Nielsen et al., 2004; Carey et al., 2006a; Haffty et al., 2006; Bauer et al., 2007; Rakha et al., 2007; Reis-Filho and Tutt, 2008; Schneider et al., 2008; Dawson et al., 2009; Jaspers et al., 2009). The patients with TNBC lacking ER- α , PR and erbB2 receptor expression are unresponsive to the treatments with anti-estrogen, hormonal therapies and/or specific inhibitors of erbB2 cascade such as trastuzumab (Schneider et al., 2008; Jaspers et al., 2009). Moreover, although TNBC patients initially respond to taxaneand/or anthracycline-based chemotherapies, the development of chemoresistance usually leads to disease recurrence and the death of patients (Dontu et al., 2004; Carey et al., 2006b; Sorlie et al., 2006; Rakha et al., 2007; Cheang et al., 2008; Li et al., 2008; Reis-Filho and Tutt, 2008; Dawson et al., 2009; Jaspers et al., 2009). In addition, the erbB2-overexpressing subtype has also been associated with a poorer prognosis and patient survival relative to differentiated ER α ⁺ breast cancer subtypes despite patients are generally responsive to adjuvant treatment with trastuzumab (Carey et al., 2006b).

Collectively, these observations suggest that the malignant transformation of immature ER α ⁻stem/progenitor cells in the basal compartment of breast epithelium may result in highly aggressive breast cancer subtypes that are less responsive to current clinical treatments. In support with this, it has been reported that the targeted expression of stabilized β -catenin in basal myoepithelial cells of mouse mammary epithelium resulted in an enhanced proliferation of basal-type celllike progenitors possessing an abnormal differentiation potential. This oncogenic event led to the development of invasive basal-type

carcinomas (Teuliere et al., 2005). The ER α ⁻ breast cancer cells, which did not express the metastasis-associated gene 3 (MTA3) that inhibits snail transcriptional activity, also possessed a lower level of E-cadherin and displayed a higher migratory capacity than the ER α ⁺ breast cancer cells (Fujita et al., 2003). Moreover, an increased expression of NF- κ B in ER α ⁻ breast cancer cells may lead to an induction of EMT program throughout the stimulation of transcription factor RelB and enhanced expression of the anti-apoptotic protein, Bcl-2 that may contribute to treatment resistance (Wang et al., 2007b). In this matter, a differently expressed gene pattern, designated as invasiveness 186-gene signature (IGS), has also been detected in CD44⁺/CD24^{-/low} tumorigenic breast cancer stem/progenitor cells relative to that of normal breast epithelial cells and associated with a poor overall survival of patients with breast cancer (Liu et al., 2007). Among the genes expressed in this very little population of CD44⁺/CD24^{-/low} tumor-initiating cells, there are the gene products associated with the NF- κ B and MAPK pathways, and epigenetic control of gene expression (Liu et al., 2007). In this regard, it noteworthy that the metaplastic and claudin^{low} breast cancer subtypes, which are usually enriched for the markers linked to EMT and tumorigenic CD44⁺/CD24⁻ stem cell-like features and chemoresistance, frequent display an aberrant activation of PI₃K/Akt pathway (Hennessy et al., 2009). Moreover, the results from another study have also revealed that the TGF- β pathway may be specifically activated in CD44⁺ breast cancer cells and its inhibition induced a more epithelial phenotype (Shipitsin et al., 2007). These data support the therapeutic interest of targeting NF- κ B, MAPK, PI₃K/Akt and/or TGF- β signaling elements to prevent the EMT process, eradicate breast cancer-initiating cells and improve the current clinical chemotherapies.

Heterogeneity of brain cancers

Among the brain cancer subtypes, the glioblastoma multiformes (GBMs), which represent a heterogeneous population of cancer cells, may arise from the malignant transformation of neural stem cells (NSCs) into brain tumor stem cells (BTSCs) (Galli et al., 2004; Yuan et al., 2004; Tso et al., 2006). BTSCs can acquire the mesenchymal properties like mesenchymal stem cells and give rise to further differentiated progenies. Primary GBMs, which are aggressive brain cancers that are frequently accompanied with EGFR overexpression, typically progress rapidly without evidence of a transitory step of lower-grade tumor. In this regard, it has been reported that CD133⁺ BTSCs found in three primary cell lines established from glioblastoma patients, expressed high mRNA levels of diverse stem cell-like markers (Liu et al., 2006a). These stemness gene products include CD44 and OCT-3/4 biomarkers, MGMT, BCRP/ABCG2 transporter, anti-apoptotic factors such as Bcl-2, survivin and inhibitor of apoptosis proteins (IAPs), hedgehog signaling elements SHH/PTCH/GLI and the transcriptional repressor of the E-cadherin, snail (Liu et al., 2006a). The isolated CD133⁺ glioblastoma cells were also more resistant to chemotherapeutic agents, such as temozolomide, carboplatin, etoposide and paclitaxel, as compared to the CD133⁻ cell fraction (Liu et al., 2006a). Importantly, higher levels of CD133 stem cell-like surface marker, which may be associated with the presence of CD133⁺ BTSCs, were also detected in recurrent GBM tissues obtained from five patients relative to their respective newly diagnosed tumors (Liu et al., 2006a). It has also been reported that CD133⁺ glioma stem/progenitor cells from a recurrent tumor after malignancy progression displayed more aggressive and invasive phenotypes *in vivo* than CD133⁺ glioma stem/progenitor cells from the primary tumor of same patient (Huang et al., 2008). Together these observations suggest that the acquisition of a more malignant phenotype by CD133⁺ BTSCs during disease progression and their high chemoresistance may contribute, at least in part, to the recurrence of highly aggressive primary GBMs. On the opposite end, the secondary or progressive GBMs, which are often characterized by mutations in the p53 suppressor gene, appear to derive from low-grade tumors that did not show the changes in the gene expression pattern that are usually associated with the EMT program (Tso et al., 2006). Hence, the clinical

management of these two different brain cancer subtypes with different aggressivity and phenotypic markers generally require distinct types of therapeutic treatments.

In light of together these observations, it appears that different cancer subtypes may originate from distinct tumorigenic cancer stem/progenitor cells that acquire a specific oncogenic gene profile during cancer development and EMT process. Therefore, the management of these cancer subtypes may require different therapeutic strategies. In respect with this, we describe new targeting approaches that have been developed to eradicate the total cancer cell mass including tumor- and metastasis-initiating cells and their differentiated progenies.

Novel cancer therapies

The progression of organ-confined cancers to locally invasive and metastatic disease stages that are resistant to current anti-hormonal, radiation and/or chemotherapeutic treatments represents one of the major causes of disease recurrence and cancer-related deaths (Lacroix, 2006; Gray-Schopfer et al., 2007; Mimeault et al., 2007b, 2008a; Sathornsumetee et al., 2007; Sorscher, 2007; Mimeault and Batra, 2008c). Therefore, the molecular targeting of cancer- and metastasis-initiating cells that can contribute in a substantial manner to drive tumor growth and metastases at distant tissues and organs, resistance to current conventional therapies and disease relapse, constitute a promising approach to develop novel effective combination therapies against aggressive and recurrent cancers (Horak et al., 2008; Le Tourneau et al., 2008; Mimeault and Batra, 2008c,d; Mimeault et al., 2008a; Steeg and Theodorescu, 2008). Especially, the molecular targeting of the EMT- and multidrug resistance-associated molecules in cancer- and metastasis-initiating cells and their local microenvironment represent potential therapeutic strategies for overcoming treatment resistance and improving current cancer therapies (Wang et al., 2002; Horak et al., 2008; Steeg and Theodorescu, 2008). Consistently, numerous recent studies have revealed that molecular targeting of hedgehog, EGFR, Wnt/ β -catenin, Notch, HA/CD44 and/or SDF-1/CXCR4 pathway as well as PI₃K/Akt/mTOR, NF- κ B, snail or twist signaling components and ABC multidrug transporters could be effective to eradicate the total cancer cell mass, including cancer- and metastasis-initiating cells, and thereby prevent disease relapse (Mimeault et al., 2007b, 2008b; Mimeault and Batra, 2008b,c,d, 2009; Chen et al., 2009; DiMeo et al., 2009).

Conclusions and perspectives

Recent advances in basic and clinical oncology have revealed that the tumorigenic and migrating cancer stem/progenitor cells can provide critical functions in tumor formation, metastases at distant sites, treatment resistance and disease relapse. Consequently, the molecular targeting of tumor- and metastasis-initiating cells and their microenvironment may represent a potential strategy for improving the efficacy of current cancer treatments. Future investigations are still essential to more precisely determine the specific biomarkers and altered gene products regulating the self-renewal, differentiation and/or treatment resistance of tumor- and metastasis-initiating cells during cancer initiation and progression to locally invasive and metastatic disease stages. Especially, a comparative analysis of gene expression profiles observed for tumor-initiating cells during primary cancer development and EMT process versus their respective normal tissue-resident stem/progenitor cells should shed light on the molecular transforming events occurring in these immature malignant cells and their pathological consequences. These studies should allow to more precisely define the altered gene products that can contribute to the acquisition of a migratory phenotype by tumor-initiating cells during the EMT program and formation of invasive and metastatic

cancer subtypes and help to develop new molecular targeting strategies that selectively eradicate these immature tumor-initiating cells.

In addition, a comparative analysis of signaling elements deregulated in tumorigenic and metastatic cancer stem/progenitor cell subpopulation relative to tumorigenic but not metastatic cancer stem/progenitor cell subset isolated from primary and metastatic patient tissues and well established cancer cell lines should permit to characterizing their specific phenotypic features. The determination of molecular events involved in the transcriptional up-regulation of SDF-1 and CXCR4 and downstream signaling effectors activated through SDF-1/CXCR4 axis in tumor- and metastasis-initiating cells during cancer progression is also of particular therapeutic interest. Furthermore, the establishment of molecular mechanisms associated with the specific migration of metastasis-initiating cells to pre-determinate metastatic sites, dormancy phenomenon and re-activation of metastasis-initiating cells at distant sites after a long latency, is also essential to identify new potential therapeutic targets to counteract the metastasis formation and disease relapse. Additional investigations to establish the signaling elements deregulated in local microenvironment of tumor- and metastasis-initiating cells is also important to design novel adjuvant cancer treatments for reversing MDR phenotype and preventing disease recurrence. These studies should lead to identification of new biomarkers and molecular therapeutic targets that could be exploited to develop new diagnostic and prognostic methods and preventive and therapeutic approaches for treating and even curing the patients diagnosed with locally advanced, metastatic, recurrent and lethal cancers.

Abbreviations

ABC	ATP-binding cassette
ALDH	aldehyde dehydrogenase
BPH	benign prostatic hyperplasia
BTSCs	brain tumor stem cells
CDK	cyclin-dependent kinase
COX-2	cyclooxygenase 2
CXCR4	chemokine receptor 4
ECM	extracellular matrix
EGF	epidermal growth factor
EGFR	epidermal growth factor receptor
EMT	epithelial-mesenchymal transition
Fzd	Frizzled receptor
KIT	stem cell factor receptor
HA	hyaluronan
MAPKs	mitogen-activated protein kinase
MEK	extracellular signal-related kinase kinase
MET	mesenchymal-epithelial transition
MDR	multidrug resistance
MMPs	matrix metalloproteinases

MGMT	O6-methylguanine DNA methyltransferase
NF-κB	nuclear factor-kappa B
NSCs	neural stem cells
PI₃K	phosphoinositide 3'-kinase
PTCH	hedgehog patched receptor
PTEN	phosphatase and tensin homolog deleted on chromosome 10
Rb	retinoblastoma
RTK	receptor tyrosine kinase
SDF-1	stromal cell-derived factor-1
SHH	sonic hedgehog ligand
SMO	smoothened co-receptor
TERT	telomerase reverse transcriptase
TK	tyrosine kinase
TR	telomere RNA component
uPA	urokinase-type plasminogen activator
VEGF	vascular endothelial growth factor
VEGFR	vascular endothelial growth factor receptor and Wnt, Wingless ligand

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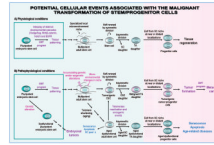


Fig. 1.

Hierarchical model of the clonal expansion and differentiation of adult stem/progenitor cells during epithelial tissue regeneration in physiological conditions and cancer initiation and progression through their malignant transformation. This scheme shows **(A)** the tissue patterning derived through the differentiation of pluripotent embryonic stem cells during normal embryonic development. The symmetric or asymmetric division of normal tissue-resident adult stem cells (SC) into transit-amplifying (TA)/intermediate cells that in turn can regenerate the bulk mass of poorly, moderately and terminally differentiated cells in the tissue from which they originate in homeostatic conditions or after tissue injury is also illustrated. Moreover, this scheme shows **(B)** the possibility of embryonal tumor formation derived of the malignant transformation of pluripotent embryonic stem cells. The malignant transformation of adult stem/progenitor cells into cancer stem/progenitor cells (CSCs), which may be induced through genetic and/or epigenetic alterations in these immature cells and changes in their local microenvironments including activated stromal cells. The tumorigenic cancer stem/progenitor cells endowed with an aberrant differentiation potential may give rise to malignant TA cells that in turn can generate the bulk mass of poorly, moderately and/or terminally differentiated cancer cells forming tumor. In addition, the chronological aging of tissue-resident adult stem/progenitor cells concomitant with telomere shortening, which may lead to their dysfunction and loss *via* senescence and apoptosis and age-related diseases is also indicated. In contrast, the re-activation of telomerase and accumulation of genetic and/or epigenetic alterations in adult stem/progenitor cells with advancing age may result in their malignant transformation and cancer development.

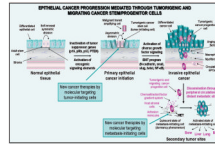


Fig. 2.

Model of epithelial cancer initiation and progression mediated through tumorigenic and migrating cancer stem/progenitor cells. The scheme shows the cancer initiation through the accumulation of genetic abnormalities in tissue-resident adult stem cells. The asymmetric division of cancer stem cells localized in the basal compartment into transit-amplifying (TA) cancer progenitor cells can generate the bulk mass of differentiated cancer cells constituting the solid tumor. Furthermore, the transformation of tumorigenic stem/progenitor cells into migrating cancer stem/progenitor cells, which may be induced by the sustained activation of distinct growth factor signaling pathways during the epithelial-mesenchymal transition (EMT) program, is also shown. The possible invasion of certain tumorigenic and migrating cancer stem/progenitor cells in the activated stroma which may lead to their dissemination through the peripheral circulation at distant sites along chemoattractant ligand gradient systems such as SDF-1/CXCR4, and adhesion to ECM components is also illustrated. Moreover, the possible loss of the migratory phenotype of cancer stem/progenitor cells *via* the occurrence of mesenchymal-epithelial transition (MET) at secondary tumor sites is indicated. The dormancy phenomenon of metastasis-initiating cells and their possible re-activation associated with the formation of secondary tumor formation under specific microenvironmental conditions at distant sites is also indicated. The new cancer therapies by molecular targeting of tumor- and metastasis-initiating cells to counteract cancer progression and metastases at distant sites are also indicated.