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# The Role of Mesenchymal Stem Cells in the Tumor Microenvironment

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## 1. Introduction

Currently, there are many promising clinical trials using mesenchymal stem cells (MSCs) in cell-based therapies of diseases ranging widely from graft-versus-host to joint and cartilage disorders (Salem and Thiemermann 2010; Tolar, Le Blanc et al.). Increasingly, however, there is a concern over the clinical use of MSCs because they are also known to home to tumors and once resident in the tumor microenvironment (TME) to support tumor growth and spread (Karnoub, Dash et al. 2007; Kidd, Spaeth et al. 2008; Coffelt, Marini et al. 2009; Klopp, Gupta et al. 2010; Klopp, Gupta et al. 2011). Conversely, other studies have reported that MSCs found in the TME diminish tumor growth, which has further generated some controversy in this field (reviewed in (Klopp, Gupta et al. 2010; Klopp, Gupta et al. 2011). Either way as a result of the MSC propensity for the TME, genetically modified MSCs that can act as “Trojan horses” and deliver anti-cancer therapeutics into the tumor stroma are being evaluated as a promising new specific cell-based therapy for cancer.

Our group established that MSCs in the ovarian tumor microenvironment promoted tumor growth and favored angiogenesis (Zwezdaryk, Coffelt et al. 2007; Coffelt and Scandurro 2008; Coffelt, Marini et al. 2009). We also developed new methodology to induce the conventional mixed pool of MSCs into two uniform but distinct phenotypes, *MSC1* and *MSC2* (Waterman, Tomchuck et al. 2010). We based their classification on several parallel observations reported within the monocyte literature. Like MSCs, heterogeneous bone marrow-derived monocytes respond to stress or “danger” inflammatory signals and home to tissue injury. Monocyte polarization into pro-inflammatory macrophages (M1) occurs early on in tissue repair whereas, monocyte polarization into anti-inflammatory macrophages (M2) follows later to help in tissue injury resolution (Mantovani, Sozzani et al. 2002; Martinez, Gordon et al. 2006). Although, this is a much simplified view of what occurs in the complex process of wound healing and repair, it provides a convenient paradigm to begin to dissect critical components within this biological process (Mantovani, Sica et al. 2007; Mosser and Edwards 2008; Mosser and Zhang 2008). Likewise, we believe that pro-inflammatory *MSC1* and anti-inflammatory *MSC2* provide convenient tools with which to begin to interrogate the role of MSCs in the tumor microenvironment.

In recent studies we found that *MSC2* supported ovarian cancer growth and spread while surprisingly *MSC1* had an opposite anti-tumor effect (Waterman 2011). We suggest that by

more closely studying the distinct tumor effects observed for these MSC phenotypes we may figure out why in the studies mentioned above MSCs favor tumor growth while in others MSCs attenuate tumors. In other words, induction into each discrete but uniform phenotype may help resolve some of the controversies surrounding the use of MSCs in cell based-therapies.

It is known that MSCs resident in the TME contribute mitogens, extracellular matrix proteins, angiogenic, and inflammatory factors. These contributions are not trivial to tumor growth and spread and serve to recruit specific subsets of leukocytes and endothelia to the TME that profoundly influence tumors. *MSC1* in the TME are expected to attenuate tumor growth by secretion of anti-tumor factors and recruitment of anti-tumor immunity. *MSC2* found in TME should promote tumor growth and spread by secretion of mitogens and suppressing anti-tumor immune responses. We expect that by identifying the differences between these two phenotypes we will shed some light on the growing controversy on the role of MSCs in tumors, and provide a means to safely deliver MSCs in cell-based therapies. We have attempted to provide all relevant information that is available concerning these issues in the sections included in this chapter.

## 2. Current understanding of MSCs function in the TME

Mesenchymal stem cells (MSCs) are a group of heterogeneous multipotent cells that can be easily isolated from many tissues throughout the body. Though initially isolated from the bone marrow, they are now recognized to be mostly in perivascular regions throughout the body (Feng, Mantesso et al. ; Zvezdaryk, Coffelt et al. 2007; da Silva Meirelles, Caplan et al. 2008). The discovery of these cells dates back to the 1960s (Friedenstein, Piatetzky et al. 1966). In recent years, MSCs have been widely studied due to their ability to be expanded in culture and stored without losing their capacity to differentiate into many different cells of mesodermal origin such as osteoblasts, chondrocytes, and adipocytes (Bruder, Jaiswal et al. 1997; Jaiswal, Haynesworth et al. 1997; Digirolamo, Stokes et al. 1999; Phinney, Kopen et al. 1999; Pittenger, Mackay et al. 1999). MSCs can also transdifferentiate into cells of ectodermal (Kopen, Prockop et al. 1999) and endodermal (Sun, Chen et al. 2007; Ju, Teng et al. 2010) origins. As a result, many preclinical studies have focused on evaluating the capacity of MSCs to repair and replace injured or diseased tissues of all origins.

Despite these research efforts however, there is growing evidence that the clinical benefit of MSCs in cell-based therapies is not the replacement of the injured tissue, but rather their efficiency in modulating aberrant host immune responses (Pittenger, Mackay et al. 1999; Prockop 2003; Prockop 2009). Following the remarkable clinical observations by the Le Blanc group who used the successful delivery of MSCs as a last resort to stave off graft-versus-host disease in a young boy, the immune modulating capability of MSCs is now more widely recognized (Le Blanc, Rasmusson et al. 2004). Further evidence indicating that immunomodulation is the primary activity of MSCs can be gleaned from the observation in many studies that although infused MSCs home to sites of injury and provide treatment benefit in widely ranging diseases, they can rarely be detected within the repaired tissue. Subsequent research efforts are beginning to identify the myriad ways that MSCs affect host immune responses. These appear to be mediated both by direct cell-to-cell contact and indirectly by the secretion of inflammatory factors (further discussed below) (Aggarwal and Pittenger 2005; Abdi, Fiorina et al. 2008; Uccelli, Moretta et al. 2008; Nemeth, Mayer et al. 2009; Bunnell, Betancourt et al. 2010; Singer and Caplan 2011).

Thus far, the immune modulating effects of MSCs include inhibition of the proliferation of activated CD8<sup>+</sup> and CD4<sup>+</sup> T lymphocytes and natural killer (NK) cells, recruitment and support of regulatory T cells, suppression of Th17 lymphocytes and immunoglobulin production by plasma cells, inhibition of maturation of dendritic cells (DCs), as well as attenuation of mast cells (Aggarwal and Pittenger 2005; Abdi, Fiorina et al. 2008; Uccelli, Moretta et al. 2008; Nemeth, Mayer et al. 2009; Nemeth, Keane-Myers et al. 2010). MSCs secrete various inflammatory factors including TNF- $\alpha$ -induced protein 6 (TNAIP6 or TSG-6), prostaglandin E2 (PGE2), human leukocyte antigen G5 (HLA-G5), hepatocyte growth factor (HGF), inducible nitric oxide synthase (iNOS), indoleamine-2,3-dioxygenase (IDO), transforming growth factor  $\beta$  (TGF- $\beta$ ), leukemia-inhibitory factor (LIF), and interleukin (IL)-10 (Krampera, Pasini et al. 2006; Gur-Wahnon, Borovsky et al. 2009; Bunnell, Betancourt et al. 2010; Singer and Caplan 2011).

MSCs express low levels of human leukocyte antigen (HLA) major histocompatibility complex (MHC) class I, do not express co-stimulatory molecules (B7-1/CD80 and -2/CD86, CD40, or CD40L), and must be induced to express MHC class II and Fas ligand that likely allows the safe delivery of these cells in non-self (allogeneic) hosts (Aggarwal and Pittenger 2005; Bunnell, Betancourt et al. 2010). Indeed, MSCs stand alone among the other types of stem cells such as embryonic or induced pluripotent (iPS) cells being considered in regenerative medicine for their safe, non-immune provoking, allogeneic host delivery capability. This has prompted many new and established businesses to amass expanded stockpiles of MSCs ready for use in the treatment of many human diseases including cancer (Salem and Thiemermann 2010).

Given the ability to deliver expanded, stockpiled clinical grade MSCs, knowing that they specifically home to the TME, and that they secrete mitogens, extracellular matrix proteins, angiogenic and inflammatory factors, it is not hard to conceive that MSCs might on the one hand influence tumors, and on the other hand, be used as vehicles to deliver anti-cancer agents. At issue is that despite intense study over the past few years, the effect of MSCs on tumors or their function in the TME is far from clear. Some studies report that MSCs promote tumor growth and spread while others report that MSCs attenuate tumor growth (Table 1). The distinct effects by MSCs on tumors has recently been attributed to differences in the experimental cancer model, the heterogeneity of MSC preparations, the dose or timing of the delivered MSCs, the animal host, or some as yet unknown factor (Klopp, Gupta et al. 2010; Klopp, Gupta et al. 2011). Also at play may be that the primary immunomodulatory function of MSCs is not realized in the context of most of these studies, which rely on immune compromised animal models. It is clear however, that with all of their unique properties MSCs make attractive candidates in cell therapies of cancer. In fact, a few promising pre-clinical reports have shown the delivery by MSCs of several anti-cancer therapeutics such as interferon (IFN)- $\beta$ , cytosine deaminase, tumor necrosis factor-related apoptosis-inducing ligand (TRAIL), and oncolytic viruses to tumors (Pittenger, Mackay et al. 1999; Studeny, Marini et al. 2002; Prockop 2003; Studeny, Marini et al. 2004; Nakamizo, Marini et al. 2005; Ren, Li et al. 2007; Kim, Lim et al. 2008; Ren, Kumar et al. 2008; Ren, Kumar et al. 2008; Mader, Maeyama et al. 2009; Prockop 2009). Though it would seem from these reports that any pro-tumor MSC effect is outweighed by the anti-cancer strategy, it is important to fully understand all of the contributions that MSCs have in the TME of immune competent tumors to safely use them in cell-based therapies of human disease.

It is appreciated that MSCs contribute in a number of ways within the TME. As mentioned above, it has long been documented that MSCs elaborate a number of factors directly, after

stimulation, or after contact with adjacent cells. These include mitogens, extracellular matrix (ECM) proteins, angiogenic factors, and inflammatory factors, all of which could potentially influence tumor growth and spread. These are summarized below along with some of the pro-tumorigenic and anti-tumorigenic evidence for MSCs.

## 2.1 Pro-tumorigenic evidence

There are a growing number of studies implicating a role for MSCs derived from various tissues in tumor growth and spread. Upon review of these studies and the anti-tumorigenic

Study	MSC Source	MSC:Tumor Ratio	Immune Status of animal model	Tumor Model	MSC Effect
(Muehlberg, Song et al. 2009)	Hu, Mu ASCs	10:1	-	Br	Larger tumor, increased SDF-1
(Karnoub, Dash et al. 2007)	Hu BMSCs	3:1	-	Br	Larger tumor, increased spread, CCL5-mediated
(Galie, Konstantinidou et al. 2008)	Mu ASCs	1:1	+	Br	Larger tumors, pro-angiogenesis
(Yu, Ren et al. 2008)	Hu ASCs	1:1, 1:2, 1:10	-	Lu, Glioma	Larger tumor, anti-apoptosis
(Djouad, Plerce et al. 2003; Djouad, Fritz et al. 2005)	Mu BMSCs	1:1	+	Melanoma	Larger tumors, inflammation
(Kucerova, Matuskova et al.)	Hu ASCs	1:5-1:10	-	Melanoma Glioblastoma	Larger tumors, VEGF and SDF1-CXCR4
(Coffelt, Marini et al. 2009)	Hu BMSCs	1:10	-	Ova	Larger tumors, pro-angiogenesis
(Lin, Yang et al. 2010)	Hu ASCs	1:2	-	Pr	Larger tumors, pro-angiogenesis and CXCR4
(Prantl, Muehlberg et al.)	Hu ASCs	1:10	-	Pr	Larger tumors, pro-angiogenesis
(Zhu, Xu et al. 2006)	Hu BMSCs	10:1, 1:1	-	Co	Larger tumors, pro-angiogenesis
(Shinagawa, Kitadai et al.)	Hu BMSCs	1:2	-	Co	Larger tumors, anti-apoptosis

Abbreviations: Hu- human, Mu- murine, ASC- adipose-derived MSCs, BMSCs- bone marrow-derived MSCs, Immune Status of animal model- - immune compromised +- immune competent, Br- breast, Lu- lung, Ov- ovarian, Pr- prostate, and Co- colon cancer cell lines.

Table 1. Pro-tumorigenic evidence for MSCs in the TME.

ones below it is tempting to speculate that cancers of endo- and ectodermal tissue origin are likely supported by MSCs whereas cancers of mesodermal tissue origin are likely inhibited by MSCs. However, as stated above, the fact that most of the studies are for technical reasons conducted in immune compromised animals greatly limits these conclusions and our understanding of the final outcome of MSCs in cancer. Evidence that MSCs promote tumor growth and their stated mechanism(s) is given by the studies summarized in Table 1. MSCs supported growth of breast, brain, lung, ovary, prostate, and colon, as well as lymphoma and melanoma (Kucerova, Matuskova et al. ; Shinagawa, Kitadai et al. ; Djouad, Plence et al. 2003; Djouad, Fritz et al. 2005; Zhu, Xu et al. 2006; Karnoub, Dash et al. 2007; Galie, Konstantinidou et al. 2008; Yu, Ren et al. 2008; Coffelt, Marini et al. 2009; Muehlberg, Song et al. 2009; Lin, Yang et al. 2010). The MSCs delivered at high ratios to the experimental tumor cell lines most commonly promoted tumor growth and metastasis. Most studies reported an increase in angiogenesis as a result of increased VEGF production by the MSCs in the TME. Some studies reported attenuation of tumor apoptosis. Chemokines such as Chemokine Ligand-5 (CCL5 or RANTES) and stromal-derived factor-1 (SDF-1)-C-X-C chemokine receptor-4 (CXCR4) axis effects by the MSCs were associated with elevated tumor migration and spread.

The secretion of pro-angiogenic molecules by the MSCs likely assist the tumors in capturing essential nutrients – perhaps also explaining the anti-apoptosis effects-- and in gaining the ability to spread to remote tissues – explaining the role of the chemokines. MSCs are known to secrete pro-angiogenic factors such as VEGF and possibly erythropoietin (Epo) thus this chief effect is not unexpected (Zwezdaryk, Coffelt et al. 2007; Singer and Caplan 2011). More studies are needed that focus on whether MSC conditioned medium is sufficient to elicit these responses and to test whether cell-to-cell contact by the MSCs, leukocytes, and/or cancer cells is required for the promotion of tumor growth and spread by MSCs.

## 2.2 Anti-tumorigenic evidence

While the pro-tumorigenic activity of MSCs is largely characterized by the secretion of pro-angiogenic molecules, the anti-tumorigenic activity of these cells is exemplified by modulation of members of the Wnt-signaling family (Table 2). MSCs inhibited the growth of tumors in several different models (Maestroni, Hertens et al. 1999; Ohlsson, Varas et al. 2003; Khakoo, Pati et al. 2006; Lu, Yuan et al. 2008; Qiao, Xu et al. 2008; Qiao, Xu et al. 2008; Cousin, Ravet et al. 2009; Otsu, Das et al. 2009; Zhu, Sun et al. 2009; Dasari, Kaur et al. ; Dasari, Velpula et al. ; Secchiero, Zorzet et al.). For instance, in studies that used fetal tissue derived MSCs, their secretion of the Wnt-signalling inhibitor Dickkopf-related protein-1 (DKK-1) inhibited breast and liver cancer cell lines (Qiao, Xu et al. 2008; Qiao, Xu et al. 2008). When the researchers used a neutralizing antibody or small interfering RNA to block DKK-1 within MSCs, the inhibitory tumor effects were attenuated. In the DKK-1 associated inhibition of primary leukemia by adipose-derived MSCs (ASCs), the stem cell transcription factor NANOG was also implicated (Zhu, Sun et al. 2009).

Interestingly, in an immune competent model, MSCs typically believed to be immune suppressive, recruited leukocytes and appeared to favor pro-inflammatory monocyte/granulocyte infiltration, which promoted rat colon carcinoma growth (Ohlsson, Varas et al. 2003). In the other immune competent model studies, one reported lack of immune suppression or attenuation of T-cell activation by the admixed MSCs but did not report the changes in any other pro-inflammatory leukocytes, and the other study was

focused more on the effect on angiogenesis by the MSCs rather than on inflammatory cells (Lu, Yuan et al. 2008; Otsu, Das et al. 2009).

Study	MSC Source	MSC:Tumor Ratio	Immune Status of animal model	Tumor Model	MSC Effect
(Khakoo, Pati et al. 2006)	Hu BMSCs	1:1, 2:1	-	Kaposi's Sarcoma	Smaller tumors, E-cadherin dependent AKT-inhibition
(Secchiero, Zorzet et al.)	Hu BMSCs	1:2, 1:10	-	NH-Lymphoma	Smaller tumors, increased animal survival
(Lu, Yuan et al. 2008)	Mu BMSCs	2-4:1	+	Insulinoma Li	Decreased ascites, pro-apoptosis
(Zhu, Sun et al. 2009)	Hu ASCs	1:10	-	Leukemia	DKK-1 mediated anti-proliferation
(Cousin, Ravet et al. 2009)	Hu ASCs	10 <sup>3</sup> ASCs/mm <sup>3</sup> tumor	-	Pan	Smaller tumors
(Otsu, Das et al. 2009)	Mu BMSCs	10 <sup>6</sup> MSCs/700mm <sup>3</sup> tumor	+	Melanoma	Smaller tumors, anti-angiogenesis
(Maestroni, Hertens et al. 1999)	Hu BMSCs	1:1	-	Melanoma, Lu	Smaller tumors and mets with GM-CSF tx MSCs
(Dasari, Kaur et al. ; Dasari, Velpula et al.)	Hu UCSCs	1:4	-	Glioma	Smaller tumors, ↑PTEN, ↓PI3K,AKT
(Qiao, Xu et al. 2008)	Hu MSCs-TERT tx	1:100	-	Br	Smaller tumors, less mets, DKK-1 mediated Wnt1 inhibition
(Qiao, Xu et al. 2008)	Hu MSCs-TERT tx	1:1	-	Li	Smaller tumors, less mets, DKK-1 mediated Wnt1 inhibition
(Ohlsson, Varas et al. 2003)	Mu BpMSCs- <i>c-myc</i>	1:1-10	+	Co	Smaller tumors, ↑inflammation

Abbreviations: Hu- human, Mu- murine, ASC- adipose-derived MSCs, BMSCs- bone marrow-derived MSCs, UCSCs- umbilical cord-derived MSCs, MSCs-TERT tx -MSC cell line immortalized with telomerase vectors, BpMSCs-*c-myc*-bone marrow-derived MSC progenitor cells immortalized with *c-myc*, Immune Status of animal model- - immune compromised +- immune competent, Br- breast, Co- colon, Li- liver, Lu- lung, NH- Non-Hodgkin's lymphoma, and Pan- pancreas cancer cell lines. DKK-1- dickkopf-related protein 1, GM-CSF-granulocyte/ monocyte-colony stimulating factor, PTEN- phosphatase and tensin homolog 10, PI3K-phosphoinositol-3-kinase.

Table 2. Anti-tumorigenic evidence for MSCs in the TME.

### 2.3 Controversies

Greater than a 100 clinical trials are underway or completed that investigate MSC-based therapy of human disease, and thus far the reports of adverse effects related to the therapy have been unremarkable (Salem and Thiemermann 2010; Tolar, Le Blanc et al. 2010; Singer and Caplan 2011). Therapy-related tumorigenicity has not been found, yet the preclinical studies presented above argue that we should carefully study this MSC potential. The question is why did MSCs promote cancer growth and spread in some studies, while in others MSCs diminished growth and spread? To begin to address this question there are a few important issues that have to be considered. First is the fact that surprisingly the chief effect of MSC-based therapies on disease is the modulation of the inflammatory host responses and not the replacement of injured tissue. Secondly, this observed therapeutic benefit is carried out by a few lingering MSCs that survive the relatively quick clearance of the cell bolus from the circulation—given that very small numbers of MSCs are ever detected at the sites of injury (Prockop 2009). Thirdly, it is known that both the adaptive and innate immune response arms profoundly influence tumor growth and spread by a complex interplay between inflammation and immunosurveillance (Frese and Tuveson 2007; Cheng, Ramesh et al. 2010). To resolve some of this controversy and to better understand the complex nature of the MSC-tumor interaction these issues need to be taken into account in future studies.

It is difficult to accurately model tumorigenesis with human tumor xenograft models in immunodeficient mice to finally resolve the effect that MSC-based therapy will have on cancer (Frese and Tuveson 2007; Cheng, Ramesh et al. 2010). Moreover, the number of MSCs interacting with the tumor must reflect more closely what is observed by the clinical experience. To more precisely model tumorigenicity attempts have been made at humanizing the murine immune system by eliminating the endogenous immune system followed by engraftment of human bone marrow or immune cells (Frese and Tuveson 2007). The problem with this approach has been that species-specific differences in both arms of the immune system confound interpretations. Immunocompetent autochthonous mouse models of human cancer provide a valuable tool that better addresses some of these issues. Though far from perfect, these models more closely parallel human carcinogenesis by allowing intrinsic tumor formation with immune surveillance and offer a better alternative system to study MSC-tumor interactions.

Apart from the limitations of current cancer models there are many other reasons that have been suggested to explain the divergent effects of MSCs in tumors (Klopp, Gupta et al. 2010; Klopp, Gupta et al. 2011). These include the heterogeneity of cells present in current MSC preparation protocols. Convention dictates that more homogeneous preparations of MSCs will also yield more consistent therapeutic outcomes with these cells. However, provided that we can overcome this hurdle and deliver more uniform cells, we may never get away from the variability that comes from the human donors. The age, gender, weight, and disease status of the donor may always affect efficacy outcomes and needs to be investigated more closely. Differences in the tissue source of the MSCs, whether bone marrow, adipose, umbilical cord, or other, also appear to affect a number of MSC functions (Sakaguchi, Sekiya et al. 2005; Hass, Kasper et al. 2011). Further complicating matters in all MSC-based therapy is the cell number and dosing frequency used to achieve a particular therapeutic efficacy. Cancer is a complex disease and to fully understand the contribution of MSCs, which are also intricate, more careful consideration of all these issues needs to be given. Despite these hurdles, MSCs remain an intriguing vehicle that can specifically target tumors.



### 3. Contributions by MSCs to tumors

In spite of all the limitations described, there is agreement about certain factors that MSCs elaborate that are important to tumorigenesis. It has long been known that MSCs synthesize a broad spectrum of growth factors, extracellular matrix proteins (ECM), cytokines, chemokines, and angiogenic molecules that have effects on cells in their vicinity. The effects of the bioactive molecules that MSCs secrete can be either direct, indirect, or even both: direct by causing intracellular signaling or indirect by causing another cell in the vicinity to secrete a bioactive factor. The indirect activity is typically termed "trophic", based on the original use of this word in neurobiology to distinguish neurotransmitters from other bioactive molecules released from nerve terminals (Caplan and Dennis 2006; Meirelles Lda, Fontes et al. 2009; Singer and Caplan 2011).

Typically, the bioactive molecules that are released from MSCs are reported to be relatively constant between different donors, regardless of age or health status of the donor. However, there can be some donor-specific differences in the levels of the secreted molecules-- that can be as high as a ten-fold difference. Moreover, the specific bioactive agents secreted by individual MSCs are also controlled by their functional status, level of differentiation, and the influence of their local microenvironments (Phinney, Kopen et al. 1999; Djouad, Fritz et al. 2005; Caplan and Dennis 2006; Krampera, Pasini et al. 2006; Tomchuck, Zvezdaryk et al. 2008; Nemeth, Mayer et al. 2009; Prasanna, Gopalakrishnan et al. ; Singer and Caplan 2011). It is expected that MSCs, as multipotent stem cells, will elaborate different levels and arrays of bioactive molecules as they differentiate into defined lineages. Additionally, the pattern and quantity of these secreted factors is well known to feed back on the MSC itself and change both its functional status and physiology.

These MSC paracrine and autocrine factors can have profound effects on local cellular dynamics. For instance, the marrow stroma derived from MSCs not only provides the matrix that supports cell anchorage, but also helps to maintain nearby endothelia and hematopoietic cells. In stroma poor niches within the marrow the hematopoietic stem cells (HSCs) will begin distinct programs of differentiation. The interdependence of MSCs and HSCs in the marrow is governed by the secretion of bioactive molecules such as the stromal-derived factor-1 (SDF1) to C-X-C chemokine receptor-4 (CXCR4) axis that helps support full hematopoietic lineage progression (Lopez Ponte, Marais et al. 2007).

#### 3.1 Soluble, Extracellular Matrix (ECM), and angiogenic factors

The secretion of these broad range bioactive molecules is now believed to be the main mechanism by which MSCs achieve their therapeutic effect and that likely most affect the tumor microenvironment. These are typically divided by the processes they affect, such as mitogenic, angiogenic, apoptotic, or inflammatory/immune modulating (Table 3). We have added exosomes as a new category to these bioactive factors. Exosomes appear to be a previously unrecognized secretory vesicle that can affect neighboring cells. We include mitogens, Extracellular Matrix (ECM) proteins, and angiogens, exosomes and inflammatory/immune modulating bioactive factors as molecules potentially contributed by MSCs but caution that this is not an exhaustive list of all MSC products. Some of the molecules overlap in function, some of the molecules play greater roles in one species versus another (e.g.-mouse vs. human), and some of the molecules are released only following

specific stimulation or activation (Tomchuck, Zwezdaryk et al. 2008; Klopp, Gupta et al. 2010; Waterman, Tomchuck et al. 2010; Klopp, Gupta et al. 2011). These have been recently reviewed (da Silva Meirelles, Caplan et al. 2008; Klopp, Gupta et al. 2010; Klopp, Gupta et al. 2011; Singer and Caplan 2011).

Molecule Types	Molecules	Study
Mitogens	bFGF, G-CSF, GM-CSF, HGF, IGF-I, IL6, Leptin, LIF, SCF, SDF-1, stanniocalcin-1, TGF $\beta$ , VEGF	(Zwezdaryk, Coffelt et al. 2007; Block, Ohkouchi et al. 2009; Meirelles Lda, Fontes et al. 2009[Tomchuck, 2008 #621; Klopp, Gupta et al. 2010; Waterman, Tomchuck et al. 2010; Klopp, Gupta et al. 2011)
Extracellular Matrix Proteins	Collagens, Fibronectin, Laminin	(Zuckerman and Wicha 1983; Hashimoto, Kariya et al. 2006; Zwezdaryk, Coffelt et al. 2007; Tomchuck, Zwezdaryk et al. 2008; Meirelles Lda, Fontes et al. 2009; Waterman, Tomchuck et al. 2010)
Angiogens	Angiopoetin-1, bFGF, IL6, IL8, Leptin, stanniocalcin-1, VEGF	(Zwezdaryk, Coffelt et al. 2007; Tomchuck, Zwezdaryk et al. 2008; Meirelles Lda, Fontes et al. 2009; Waterman, Tomchuck et al. 2010)
Exosomes	Pro-inflammatory molecules, miRNAs	(Anand 2010; Chen, Lai et al. 2010; Lai, Arslan et al. 2010)
Inflammatory/ Immune Modulating	galectin-3, galectin-1, HGF, HLA-G, IDO, IL1 $\beta$ , IL1RA, IL6, IL8, IL12, iNOS, IP-10, LIF, MCP-1, MIP-1, PGE2, semaphorin-3A, RANTES, SDF-1, stanniocalcin-1, TGF $\beta$ , TSG-6	(Zwezdaryk, Coffelt et al. 2007; Tomchuck, Zwezdaryk et al. 2008; Block, Ohkouchi et al. 2009; Meirelles Lda, Fontes et al. 2009; Bartosh, Ylostalo et al. 2010; Bunnell, Betancourt et al. 2010; Klopp, Gupta et al. 2010; Waterman, Tomchuck et al. 2010; Danchuk, Ylostalo et al. 2011; Klopp, Gupta et al. 2011)

Abbreviations: bFGF- basic fibroblast growth factor, CCL- C-C motif chemokine ligand, CXC- C-X-C-motif chemokine, CXCL-CXC-ligand, G-CSF-granulocyte-colony stimulating factor, GM-CSF-granulocyte-macrophage-colony stimulating factor, HGF-hepatocyte growth factor (scatter factor), HLA-G- human leukocyte antigen-G, IDO- indoleamine 2,3-dioxygenase, IGF-I-insulin-like growth factor-1, IL-interleukin, IL-1RA- interleukin-receptor 1 antagonist, iNOS-inducible nitric oxide synthase, IP-10-interferon-gamma-inducible protein 10 (CXCL10), LIF-leukemia inhibitory factor, MCP-1-monocyte chemoattractant protein-1 (CCL2), MIP1-macrophage inflammatory protein-1 (CCL3), PGE2-prostaglandin-E2, PlGF-placental-derived growth factor, RANTES- regulated upon activation normal T cell expressed and secreted (CCL5), SCF-stem cell factor, SDF-1-stromal-derived factor-1, TGF $\beta$ -transforming growth factor- $\beta$ , TSG-6- TNF-alpha stimulated gene/protein 6, VEGF-vascular-derived endothelial growth factor (vascular permeability factor, VPF).

Table 3. Molecules Contributed by MSCs.

### 3.2 Exosomes

A recently described form of intercellular communication that may also be important in MSC-tumor exchanges is exosomes. These are endosome-derived vesicles of about 40-100 nm that are formed by the involution of endosome membranes resulting in the formation of

multi-vesicular bodies (MVBs). Following certain physiological conditions, the MVBs fuse with the plasma membrane and release the exosomes into the circulation or tissue microenvironment. Exosomes have a “saucer-shaped” morphology as determined from electron microscopy analyses. Various methods have been developed to enrich for exosomes derived from a number of cell types including antigen-presenting cells (APCs), monocytes, T-lymphocytes, reticulocytes, mast cells, platelets, fibroblasts, tumor cells, and MSCs (Anand 2010; Lai, Arslan et al. 2010; Tan, De La Pena et al. 2010).

Investigators studying the cardioprotective effect of human embryonic stem cell-derived MSC-conditioned medium (CM) on myocardial ischemia/reperfusion injury reasoned based on proteomic analyses that exosomes were responsible for the beneficial effect (Sze, de Kleijn et al. 2007; Lai, Arslan et al. 2010). Their unbiased proteomic profiling of proteins secreted by MSCs revealed an abundance of membrane and cytosolic proteins. This suggested to them that the trophic effects of MSCs were not mediated by soluble growth factors and cytokines alone. Sze *et al.* proceeded to enrich for particles by size-exclusion fractionation on HPLC. Based on the size and the composition of the particles they figured exosomes were present in the condition medium of MSCs. Moreover they demonstrated that the enriched fraction of exosomes reduced infarct size in a mouse model of myocardial ischemia/reperfusion injury.

The particles could be visualized by electron microscopy and were shown to be phospholipid vesicles consisting of cholesterol, sphingomyelin, and phosphatidylcholine. Moreover, they were composed of known exosome-associated proteins-- CD81, CD9, and Alix. Exosomes are known to have a specific protein composition, including CD9, CD81, Alix, TSP-1, SOD-1, and pyruvate kinase. CD9 and CD81 are tetrapannin membrane proteins that are also localized in the membrane of exosomes. Consistent with the presence of exosomes in the CM of the MSCs this study further demonstrated that CD9 in the CM was a membrane-bound protein while SOD-1 was localized within a lipid vesicle. They eliminated the possibility of immune cells or platelets as sources of exosomes with an *ex vivo* mouse model of myocardial ischemia/reperfusion injury.

Similarly in human ESC-derived MSC conditioned medium other investigators found exosomes that contained small RNAs (less than 300 nt) encapsulated in cholesterol-rich phospholipid vesicles. The small RNAs were identified by a number of biochemical and genetic criteria to be microRNAs (miRNAs). Of interest the *Let-7* family of miRNAs figured prominently in these studies (Chen, Lai et al. 2010; Koh, Sheng et al. 2010). It is becoming increasingly clear that miRNAs are potent global gene regulators of many diverse cell functions including adaptation to mitogens, low oxygen (hypoxia), and inflammation. Perhaps this might explain why exosomes are potent immune modulators (Anand 2010). Apart from the molecules present inside the lumen of exosomes, it has been suggested that certain exosomal membrane molecules can interact with their surface receptors on the target cells thereby inducing an immunomodulatory response or activating the immune system. Consistent with this notion, exosome release is enhanced following pathologies where immune activation is required. It has been suggested that immunogenic molecules on the exosomal membrane can activate leukocytes. In support of this idea is the fact that exosomes are analogous to inverted endosomes and thus display inflammatory intracellular factors present normally within plasma membrane. Taking advantage of this inflammatory nature

of exosomes, clinicians are developing cancer vaccines based on loading dendritic cells (DCs) with tumor antigens, expanding the DCs *ex vivo*, and subsequently isolating their enriched exosomes (Tan, De La Pena et al. 2010). The tumor antigen loaded exosomes are then reintroduced into patients to elicit tumor specific anti-tumor immunity.

Lastly, highlighting the interactions of tumors and MSCs, exosomes derived from tumors appear to drive adipose-derived MSC differentiation toward tumor associated myofibroblasts that can then contribute to tumor growth and spread (Webber, Steadman et al. 2010; Cho, Park et al. 2011; Cho, Park et al. 2011). Interestingly and perhaps providing a mechanism for the Wnt-signaling mediated anti-tumor effect of MSCs mentioned above,  $\beta$ -catenin was found to be contained within exosomes (Chairoungdua, Smith et al. 2010). Furthermore, exosomal release of  $\beta$ -catenin antagonized Wnt-signaling in the recipient cell. These studies emphasize the need for more intense investigations that clarify the role of both tumor- and MSC-derived exosomes in tumorigenesis. Besides identifying new components of tumor biology such studies may identify new therapeutic interventional agents.

### 3.3 Immune modulation

Apart from the ability of MSCs to contribute mitogens, ECM proteins, pro-angiogenic molecules, inflammatory agents, and exosomes to the TME, their most significant contribution may be modulating specific subsets of immune cells (Table 4)(Fibbe, Nauta et al. 2007; Nauta and Fibbe 2007; Bunnell, Betancourt et al. 2010; Roddy, Oh et al. 2011; Singer and Caplan 2011; Weiss, Bertoncello et al. 2011). The specific mechanism for this MSC role is not completely understood and may involve direct immune cell-MSC cell contact or indirect effects such as by the contribution of the factors just described or both. However, knowing the importance of immune and inflammatory cells in cancer growth and metastasis, the manner that MSCs in the TME might influence this process deserves closer attention and study.

Though initially described as an *ex vivo* phenomena requiring the stimulation of the MSCs to lead to suppression of T-lymphocyte activation or proliferation, many clinical trials have asserted immune modulation to be a primary effect of MSC-based therapies (Di Nicola, Carlo-Stella et al. 2002; Krampera, Glennie et al. 2003; Le Blanc, Rasmusson et al. 2004; Aggarwal and Pittenger 2005). In addition, these early observations prompted a number of studies to explore the distinct immune modulatory effects of MSCs derived from a variety of sources and species. Of note, although MSCs influence many immune cells, part of what makes them attractive candidates in cell-based therapies is their muted host immune responses even when delivered into a non-self (allogeneic) host. This is partly due to the fact that MSCs express low levels of human leukocyte antigen (HLA) major histocompatibility complex (MHC) class I, do not express co-stimulatory molecules (B7-1/CD80 and B7-2/CD86, CD40, or CD40L), and express MHC class II and Fas ligand only after specific stimulation.

MSCs are now known to inhibit dendritic cell maturation, B and T cell proliferation and differentiation, attenuate natural killer cell and mast cell activity, as well as support the production of suppressive T regulatory cells (Tregs) while attenuating pro-inflammatory Th17 cells (Table 4) (Najar, Raicevic et al. ; Di Nicola, Carlo-Stella et al. 2002; Krampera, Glennie et al. 2003; Aggarwal and Pittenger 2005; Beyth, Borovsky et al. 2005; Ramasamy, Fazekasova et al. 2007; Ren, Zhang et al. 2008; Uccelli, Moretta et al. 2008; Gur-Wahnon, Borovsky et al. 2009; Meirelles Lda, Fontes et al. 2009; Nemeth, Mayer et al. 2009; Bunnell, Betancourt et al. 2010; Salem and Thiernemann 2010; Tolar, Le Blanc et al. 2010; Brown, Nemeth et al. 2011; Singer and Caplan 2011).

Immune Response Arm	Cells	MSC effects
Innate	Dendritic Cells (APC)	Inhibition of maturation (CD80/86 expression) by STAT3 and IL10 (Beyth, Borovsky et al. 2005; Gur-Wahnon, Borovsky et al. 2009; Mezey, Mayer et al. 2009; Nemeth, Leelahavanichkul et al. 2009)
	Monocyte/Macrophages (APC)	PGE2 mediated increased IL10 secretion and attenuation of maturation (Beyth, Borovsky et al. 2005; Gur-Wahnon, Borovsky et al. 2009; Mezey, Mayer et al. 2009; Nemeth, Leelahavanichkul et al. 2009)
	Natural Killer Cells	Inhibition of proliferation and cytolytic activity (Giuliani, Oudrhiri et al. 2011)
	Mast Cells	COX-2 mediated suppression (Brown, Nemeth et al. 2011)
Adaptive	Th1	Inhibition of proliferation/activation (class switching) by HLA-G5, HGF, iNOS, COX2, IDO, PGE2, TGF $\beta$ and indirectly through support of immature APCs reviewed in (Singer and Caplan 2011)
	Th2	Inhibition of proliferation/activation (class switching) by HLA-G5, HGF, iNOS, COX2, IDO, PGE2, TGF $\beta$ and indirectly through support of immature APCs reviewed in (Singer and Caplan 2011)
	Tregs	Recruitent and support (class switching) IL10, TGF $\beta$ , LIF
	Th17	Inhibition of proliferation/activation (class switching) by COX-2 and PGE2 (Duffy, Pindjakova et al. 2011; Duffy, Ritter et al. 2011)
	B lymphocyte	Suppression of terminal differentiation to plasma cell (Asari, Itakura et al. 2009)

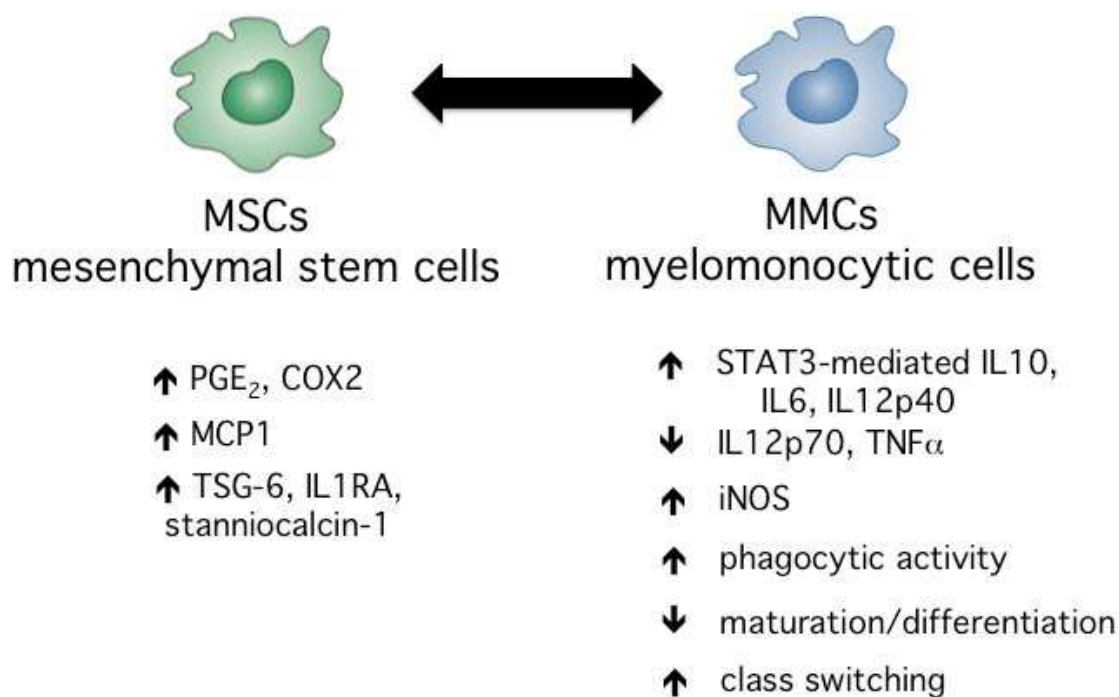
Abbreviations: COX-2- cyclooxygenase-2, HGF-hepatocyte growth factor (scatter factor), HLA-G- human leukocyte antigen-G, IDO- indoleamine 2,3-dioxygenase, iNOS-inducible nitric oxide synthase, IL10-interleukin-10, LIF-leukemia inhibitory factor, PGE2- prostaglandin-E2, STAT3- signal transducer and activator of transcription-3, TGF $\beta$ -transforming growth factor- $\beta$ .

Table 4. Immune cells modulated by MSCs.

### 3.3.1 MSCs and myelomonocytic cells

Although the details of the interactions of MSCs with T lymphocytes, B lymphocytes, natural killer cells, and dendritic cells have been investigated in some detail, the effects of MSCs on cells of myelomonocytic lineages (MMCs) observed early on by the Rachmilewitz group remained under investigated until recently (Figure 1. Beyth, Borovsky et al. 2005). The growing clinical evidence for MSCs as major regulators of immune and inflammatory processes and the central role played by MMCs (including monocytes and granulocytes) within them has sparked new interest in studies on the interplay between MSCs and MMCs. Kim and Hematti (2009) reported that human macrophages generated *in vitro* after co-culture with MSCs assume an immunophenotype defined as IL-10-high, IL-12-low, IL-6-high, and

## Consequences of MSC-Myelomonocytic Cell Interaction



Abbreviations: COX-2- cyclooxygenase-2, IL-interleukin, IL-1RA- interleukin-receptor 1 antagonist, iNOS-inducible nitric oxide synthase, MCP-1-monocyte chemoattractant protein-1 (CCL2), PGE2-prostaglandin-E2, STAT3- signal transducer and activator of transcription-3, TSG-6- TNF-alpha stimulated gene/protein.

Fig. 1. The Consequences of the Interaction Between MSCs and Myelomonocytic Cells. Though still in their infancy the studies that have begun to identify the effect of the interactions between MSCs and MMCs whether cell-cell contact dependent or not have so far described those included in the figure. Please refer to the text for details.

TNF- $\alpha$ -low secreting cells (Kim and Hematti 2009). They proposed that these MSC-educated monocytes represent a unique and novel type of alternatively activated macrophage with a potentially significant role in tissue repair. Initially, Beyth *et al.* reported that human MSCs affect monocytes or dendritic antigen-presenting cell (APC) maturation in a contact-dependent manner (Beyth, Borovsky et al. 2005). Later, it was reported that the MSCs co-cultured with the APCs induced the expression of the anti-inflammatory IL10 and that activation of the signal transducer and activator of transcription 3 (STAT3) within APCs is linked to abnormal APC differentiation and function by a new contact-dependent mechanism, that plays a critical role in mediating the immunomodulatory effects of MSCs (Gur-Wahnon, Borovsky et al. 2007; Gur-Wahnon, Borovsky et al. 2009). In order to understand this process better, they further extended their studies to tumor cells since tumors secrete a variety of bioactive factors that activate STAT3 within infiltrating APCs. Their studies demonstrated that in at least certain cellular microenvironments, cell-to-cell dependent interactions represent a novel way to activate STAT3 signaling different from the activation of STAT3 seen with soluble bioactive factors. As such this observation suggests an uncoupling of APC activation events and that may consequently independently regulate

immunity and tolerance. In agreement with these studies, the Mezey group identified other pathways involved in MSC-murine macrophage interactions (Nemeth, Leelahavanichkul et al. 2009). They also showed that LPS-stimulated macrophages produced more IL-10 when cultured with MSCs, but this effect was eliminated if the MSCs lacked the genes encoding TLR4, myeloid differentiation primary response gene-88 (MyD88), TNF-receptor-1 $\alpha$  or cyclooxygenase-2 (COX-2). Their observations demonstrated that MSCs reprogram macrophages by releasing PGE2 that then acts on the macrophages through the prostaglandin EP2 and EP4 receptors. A unique population of MSCs isolated from human gingiva (GMSCs) with similar stem cell-like properties, immunosuppressive, and anti-inflammatory functions as bone marrow-derived MSCs were also studied in this context with similar effects (Zhang, Su et al. 2010).

When co-cultured with GMSCs, macrophages acquired an anti-inflammatory M2 phenotype similarly characterized by an increased expression of IL10 and IL6, mannose receptor (MR; CD206), a suppressed production of TNF $\alpha$ , and also decreased the ability to induce Th-17 cell expansion. Interesting to the discussion on tumors and their microenvironments, they demonstrated that systemically infused GMSCs could home to wounds-- specifically to sites where host macrophages were found-- promoted M2 polarization of the co-localized monocytes, significantly enhanced wound repair, and thus presumably could promote tumor growth by similar mechanisms. In addition, they noted that GMSC treatment suppressed local inflammation by reducing the infiltration of inflammatory cells and the production of IL6 and TNF $\alpha$ , and by increased expression of IL10. Another complementary study used murine macrophages stimulated with LPS and co-cultured with MSCs and found the suppression of TNF $\alpha$ , IL6, IL12p70 and interferon $\gamma$  but increased levels of secreted IL10 and IL12p40. They noted that the murine MSC effect could be reproduced with MSC conditioned medium suggesting that bioactive factors constitutively released by the murine MSCs may be sufficient for the monocyte effect in this animal species (Maggini, Mirkin et al. 2010). They also found in cell-based therapy of mouse models that MSCs supported macrophages that showed a low expression of CD86 and MHC class II, and with a high ability to secrete IL10 and IL12p40, but not IL12 p70. They suggested in agreement with the other studies, that MSCs switch monocytes into a regulatory profile characterized by enhanced IL10 secretion, reduced inflammatory cytokine elaboration and enhanced phagocytic activity. Apart from elevated IL10 and related signaling mechanisms, other new players in the effects observed for MSCs on monocytes were recently advanced (Block, Ohkouchi et al. 2008; Block, Ohkouchi et al. 2009; Danchuk, Ylostalo et al. 2011; Prockop and Youn Oh 2011)]. Anti-inflammatory effects supported by MSC-monocyte interactions were suggested to also be partly mediated by elevated IL1 receptor antagonist (IL1RA) and by a negative feedback loop in which TNF $\alpha$  and other pro-inflammatory cytokines from resident macrophages activate MSCs to secrete the anti-inflammatory protein TNF $\alpha$  stimulated gene/protein 6 (TSG-6). These reports demonstrate that MSC derived TSG-6 acts to repress NF- $\kappa$ B signaling in the resident macrophages causing attenuation of pro-inflammatory cytokine synthesis. The investigators of these studies also proposed that MSC secreted PGE2 promotes monocytes toward an IL10 secreting phenotype as well as, that anti-inflammatory effects may also be mediated by stanniocalcin-1.

Finally, in another recent report using pre-clinical murine models it was shown that MCP1 secreted by activated MSCs contributes to the bone marrow egress, trafficking, and

recruitment of monocytes towards remote sites (Shi, Jia et al. 2011). This elegant study demonstrated the intimate and complex cooperation that exists between MSCs and myelomonocytic cells that occurs not only in peripheral tissues or tumors but also in their originating bone marrow niche. It is widely recognized that tumor infiltrating cells can include macrophages, myeloid-derived suppressor cells (MDSCs), MSCs, and TIE2-expressing monocytes that are all mostly derived from the bone marrow. MDSCs represent a heterogeneous population of cells of myeloid origin that are expanded and activated in response to growth factors and cytokines released by tumors much like MSCs. The details of the effects of MDSCs on tumors are better understood. It is known that once MDSCs are activated, they accumulate in lymphoid organs and tumors where they exert specific T cell mediated immune suppression. However, not much is known about whether MDSCs and MSCs cooperate at tumor sites or the nature of that interaction. It is tempting to suggest that MSC-myelomonocytic cell interactions including MSC-MDSC ones represent an intriguing new target for cancer therapies that would break the anti-inflammatory tumor tolerance mechanisms established by these two cell types however, there is still much left to learn before this can come to fruition. Furthermore, while the vast majority of these reports demonstrate the ability of MSCs to suppress immune responses or act in an anti-inflammatory manner, there is emerging evidence that supports their contrasting ability to elicit pro-inflammatory responses-- which may also be mediated by their interaction with myelomonocytic cells. Both anti-inflammatory and pro-inflammatory effects will be important to know in dissecting their specific roles in tumors. This information will ultimately help in the design of more effective and targeted cancer therapeutics.

### 3.3.2 Immune suppressive or anti-inflammatory responses

The expression of IDO and iNOS by MSCs has been associated with its immune suppression of T-cell proliferation. Recently, secretion of IDO by MSCs therapeutically delivered in an experimental autoimmune myasthenia gravis model inhibited the proliferation of acetylcholine receptor-specific T cells and B cells and normalized the distribution of Th1, Th2, Th17 and Treg cells (Kong, Sun et al. 2009). IDO catalyzes the conversion of tryptophan, an essential amino acid for T-cell proliferation, into kynurenine. Immune suppression by IDO results from the local accumulation of tryptophan metabolites, rather than through tryptophan depletion (Ryan, Barry et al. 2007). Expression of IDO by MSCs was thought to be IFN- $\gamma$  dependent (Krampera, Cosmi et al. 2006; Ryan, Barry et al. 2007; Bunnell, Betancourt et al. 2010). However, Opitz and colleagues recently demonstrated that IDO expression in MSCs can also be induced by activation of Toll-like receptor 3 (TLR3) and TLR4 via induction of an autocrine IFN- $\beta$  signaling loop involving protein kinase R and independent of IFN- $\gamma$  (Opitz, Litzenburger et al. 2009). Interestingly, when MSCs were treated with IFN- $\gamma$  *in vitro*, they expressed extremely high levels of IDO and very low levels of iNOS, whereas mouse MSCs expressed abundant iNOS and very little IDO. These data suggest there is species variation in the mechanisms of MSC immunosuppression (Opitz, Litzenburger et al. 2009).

Prostaglandin E2 (PGE2) is emerging as a central mediator of many of the anti-inflammatory properties of MSCs (Nauta and Fibbe 2007; Uccelli, Moretta et al. 2008). PGE-2 is synthesized from arachidonic acid by cyclooxygenase (COX) enzymes COX-1 and COX-2.



COX-1 is constitutively expressed in MSCs and COX-2 expression can be induced by inflammatory cytokines such as IL-1 $\beta$ , IL-6, IFN- $\gamma$ , and TNF- $\alpha$  (Chen, Wang et al. 2010). Inhibitors of PGE2 synthesis attenuated MSC suppression of T cells and natural killer cells (Sotiropoulou, Perez et al. 2006; Chen, Wang et al. 2010). PGE2 is associated also with the MSC-mediated inhibition of dendritic cell maturation. Nemeth *et al.* reported that activated MSCs released PGE2 causing increased production of IL10 by macrophages, and decreased production of the pro-inflammatory cytokines TNF- $\alpha$  and IL-6 in a murine sepsis model (Sotiropoulou, Perez et al. 2006). Maggini *et al.* similarly reported macrophage alterations by PGE2 (Maggini, Mirkin et al. 2010).

Mezey's group demonstrated that COX-2 is also involved in MSCs ability to suppress mast cell activation (Brown, Nemeth et al. 2011). Mast cells (MCs) have a key role in the induction of allergic inflammation and contribute to the severity of certain autoimmune diseases. An increasing body of literature also implicates MCs in the TME to affect tumor inflammation, angiogenesis, and growth (Ribatti, Nico et al. 2011). To date, few studies have investigated the potential of mast cell-MSC interactions. Since MCs are critical effector cells in allergic inflammation and they represent an important cell type to therapeutically target using the immune modulatory properties of MSCs, Mezey's group set out to study murine MC-MSCs effects. They reported that MSCs effectively suppressed specific MC functions *in vitro* and in animal models. MCs co-cultured with MSCs in direct contact, had dampened MC degranulation, pro-inflammatory cytokine production, chemokinesis, and chemotaxis. They also found that MC degranulation within mouse skin or the peritoneal cavity was suppressed following delivery of MSCs. Lastly, they discovered that these inhibitory effects were dependent on COX2 in MSCs (Brown, Nemeth et al. 2011).

Transforming growth factor- $\beta$  (TGF $\beta$ ) is an anti-inflammatory cytokine that is constitutively expressed by MSCs. The immune modulatory function of MSCs on T cells and natural killer cells can be impaired by treatment with neutralizing antibodies to TGF $\beta$  (Di Nicola, Carlo-Stella et al. 2002; Sotiropoulou, Perez et al. 2006). In contrast, several studies have also established that TGF $\beta$  had no effect on the immunosuppressive properties of MSCs (Tse, Pendleton et al. 2003; Xu, Zhang et al. 2007). These discrepancies are likely explained by differences in species or experimental conditions. The importance of TGF $\beta$  in MSC therapy was recently established in a mouse model of ragweed-induced asthma. Mezey's group again demonstrated this assertion with neutralizing antibodies and the use of MSCs derived from TGF $\beta$  knockout mice (Nemeth, Keane-Myers et al. 2010). Notably, the number of Tregs in this model was elevated by the MSC-therapy. However, the role of TGF $\beta$  in this process was not directly studied, as was done by Patel *et al.* who showed that in co-cultures of peripheral blood mononuclear cells (PBMCs) with MSCs, TGF $\beta$  produced by MSCs resulted in increased numbers of Tregs (Patel, Meyer et al. 2010).

Several other factors are associated with the potential anti-inflammatory properties of MSCs including HLA-G, hepatocyte growth factor (HGF), leukemia inhibitory factor (LIF), IL1 receptor antagonist (IL1RA), CCL2, galectin-3, galectin-1 and semaphorin-3A, most of which attenuate T lymphocyte activation and are highly expressed by MSCs (Di Nicola, Carlo-Stella et al. 2002; Ortiz, Dutreil et al. 2007; Di Ianni, Del Papa et al. 2008; Kang, Kang et al. 2008; Nasef, Ashammakhi et al. 2008; Rafei, Hsieh et al. 2008; Lepelletier, Lecourt et al. 2009; Selmani, Naji et al. 2009; Sioud, Mobergslien et al. 2010; Volarevic, Al-Qahtani et al. 2010). A recently advanced culprit is TNF- $\alpha$ -induced protein 6 TNAIP6 or TSG-6 (Lee, Pulin et al.

2009; Prockop and Youn Oh 2011). TSG-6 secretion is known to suppress inflammation through the inhibition of the inflammatory network of proteases primarily by increasing the inhibitory activity of inter- $\alpha$ -inhibitor, sequestration of hyaluronan fragments, and decreasing neutrophil infiltration into sites of inflammation. In a model of acute inflammation induced by myocardial infarction, TSG-6 knockdown in MSCs significantly reduced their anti-inflammatory therapeutic effect. The administration of recombinant TSG-6 protein largely duplicated the therapeutic effects of the delivered MSCs on inflammatory responses and infarct size (Getting, Mahoney et al. 2002; Wisniewski and Vilcek 2004; Milner, Higman et al. 2006; Forteza, Casalino-Matsuda et al. 2007; Lee, Pulin et al. 2009). Together these results make TSG-6 an interesting new factor in the anti-inflammatory effects of MSCs.

### 3.3.3 Pro-inflammatory MSC responses

Though we are beginning to better understand the many complex mechanisms associated with the secretion by MSCs of immune suppressive mediators like TSG-6, so far only a few reports have described a contrasting pro-inflammatory activity of MSCs that could be important in understanding the distinct role of MSCs in tumors. Indeed, the observation of this distinct MSCs immune effect came from studies primarily focused on the downstream consequences of TLR stimulation within these cells. TLRs are a conserved family of receptors that recognize pathogen-associated molecular patterns (PAMPs) and promote the activation of immune cells (Wright 1999-76; Triantafilou, Triantafilou et al. 2001; Sabroe, Read et al. 2003; Anders, Banas et al. 2004; Miggin and O'Neill 2006; West, Koblansky et al. 2006; Bunnell, Betancourt et al. 2010). Many TLRs (TLR1 to TLR13) have been identified and characterized in a variety of immune cell types and species. Agonists for TLRs include exogenous microbial components, such as LPS (TLR2 and 4), lipoproteins and peptidoglycans (TLR1, 2, 6), viral RNA (TLR3), bacterial and viral unmethylated CpG-DNA (TLR9), and endogenous molecules shed following cell injury, including heat shock proteins and extracellular matrix molecules (Wright 1999-77; Triantafilou, Triantafilou et al. 2001; Sabroe, Read et al. 2003; Anders, Banas et al. 2004; Miggin and O'Neill 2006; West, Koblansky et al. 2006; Bunnell, Betancourt et al. 2010). Specific agonist engagement of TLRs leads to the expression of inflammatory cytokines or co-stimulatory molecules by a MyD88 (a TLR adapter protein)-dependent or MyD88-independent signaling pathways and can promote chemotaxis of the stimulated cell. TLRs are differentially expressed on leukocyte subsets and non-immune cells and may regulate important aspects of innate and adaptive immune responses (Mempel, Voelcker et al. 2003; Hwa Cho, Bae et al. 2006; Nagai, Garrett et al. 2006; Pevsner-Fischer, Morad et al. 2006; West, Koblansky et al. 2006; Tomchuck, Zvezdaryk et al. 2008).

MSCs are among the cells that express an array of TLRs, including TLR2, 3, 4, 5, 6 and 9 (Hwa Cho, Bae et al. 2006; Pevsner-Fischer, Morad et al. 2006; Tomchuck, Zvezdaryk et al. 2008). Furthermore, studies by our group established that the stimulation of MSCs with TLR agonists led to the activation of downstream signaling pathways, including NF- $\kappa$ B, AKT, and mitogen-activated protein kinase (MAPK). Consequently, activation of these pathways triggers the previously unreported induction and secretion of pro-inflammatory cytokines, chemokines, and related TLR gene products. Interestingly, the unique patterns of affected genes, cytokines, and chemokines measured identified the TLRs as potential players in the

established MSC immune modulatory properties, as well as their ability to migrate towards injured tissues. Surprisingly, we noted that TLR4 stimulation with LPS led to the secretion of primarily pro-inflammatory mediators, such as IL-1 $\beta$  and IL6 (Tomchuck, Zvezdaryk et al. 2008). Though unexpected, previous observations reported by Beyth *et al.* recognized that LPS priming affected co-cultures of leukocytes with human MSCs and attenuated the expected human MSC-mediated inhibition of T-lymphocyte activation as well as affected their capacity to secrete interferon (Beyth, Borovsky et al. 2005). More recently, Romieu-Mourez *et al.* showed that TLR stimulation in murine MSCs similarly resulted in the production of inflammatory mediators, such as IL-1, IL-6, IL-8, and CCL5 (Romieu-Mourez, Francois et al. 2009). Furthermore, they demonstrated that TLR and IFN activated murine MSCs injected within Matrigel matrices into mice resulted in the formation of an inflammatory site attracting innate immune cells and resulting in a dramatic recruitment of neutrophils. Raicevic *et al.*, studying the effect of TLR activation within MSCs in an inflammatory milieu, observed that this environment shifted the cytokine profile to a pro-inflammatory one rather than the expected immunosuppressive one (Raicevic, Rouas et al. 2010). They similarly observed an increase in IL-1 $\beta$ , IL-6, and IL-12 after TLR activation in this inflammatory context.

Though somewhat confounding, this recent body of work on the downstream consequences of TLRs provides emerging evidence for a new pro-inflammatory immune modulating role for MSCs. The identification of the molecular details for this new pro-inflammatory MSC role, and whether it is innate or just an *in vitro* artifact, awaits further investigation. However, this novel observation is important to consider given the accelerated use of MSCs in anti-inflammatory cell-based therapies. Additionally, as Raicevic *et al.* suggest targeting of TLRs in MSCs, may avoid deleterious consequences in their use as anti-inflammatory therapies (Raicevic, Rouas et al. 2010). By contrast, TLR-activated pro-inflammatory MSCs could prove useful in breaking tolerance in the therapy of immune evasive diseases, such as cancer.

#### **4. New MSC paradigm: Pro-inflammatory MSC1 and anti-inflammatory MSC2**

Our recent studies are partly an attempt to resolve some of the controversy surrounding the potential of MSCs to be anti-inflammatory in some cases and pro-inflammatory in others or to be pro-tumor in some cancers and anti-tumor in others, as described above. These studies led us to propose a new paradigm for MSCs based on the premise that these heterogeneous cells can be induced to polarize into two distinct but homogeneously acting phenotypes--that we modeled after monocytes, the other heterogeneous bone marrow-derived cells (Figure 2. Verreck, de Boer et al. 2006).

It is established that stimulation of monocytes with known cytokines or agonists to their TLRs, including IFN- $\gamma$  and endotoxin (LPS, TLR4-agonist), polarizes them into a classical M1 phenotype that participates in early pro-inflammatory responses. IL-4 treatment of monocytes yields the alternative M2 phenotype that is associated with anti-inflammatory resolution responses (Verreck, de Boer et al. 2006). We proposed that MSCs, like monocytes, are polarized by downstream TLR signaling into two homogeneously acting phenotypes, classified as MSC1 and MSC2, following the monocyte nomenclature. We reported that TLR4 agonists polarized MSCs toward a pro-inflammatory MSC1 phenotype while the downstream consequences of TLR3 stimulation of MSCs was a skewing toward an anti-inflammatory MSC2 phenotype. This novel MSC polarization paradigm is based on the

consistent but novel outcomes observed for *MSC1* when compared with *MSC2* for several parameters, including dissimilar patterns of secretion of cytokines and chemokines and differences in differentiation capabilities, extracellular matrix deposition, TGF- $\beta$  signaling pathways, and Jagged, IDO and PGE-2 expression (Waterman, Tomchuck et al. 2010). The most compelling outcome was opposite effects of each cell type on T-lymphocyte activation (Waterman, Tomchuck et al. 2010).

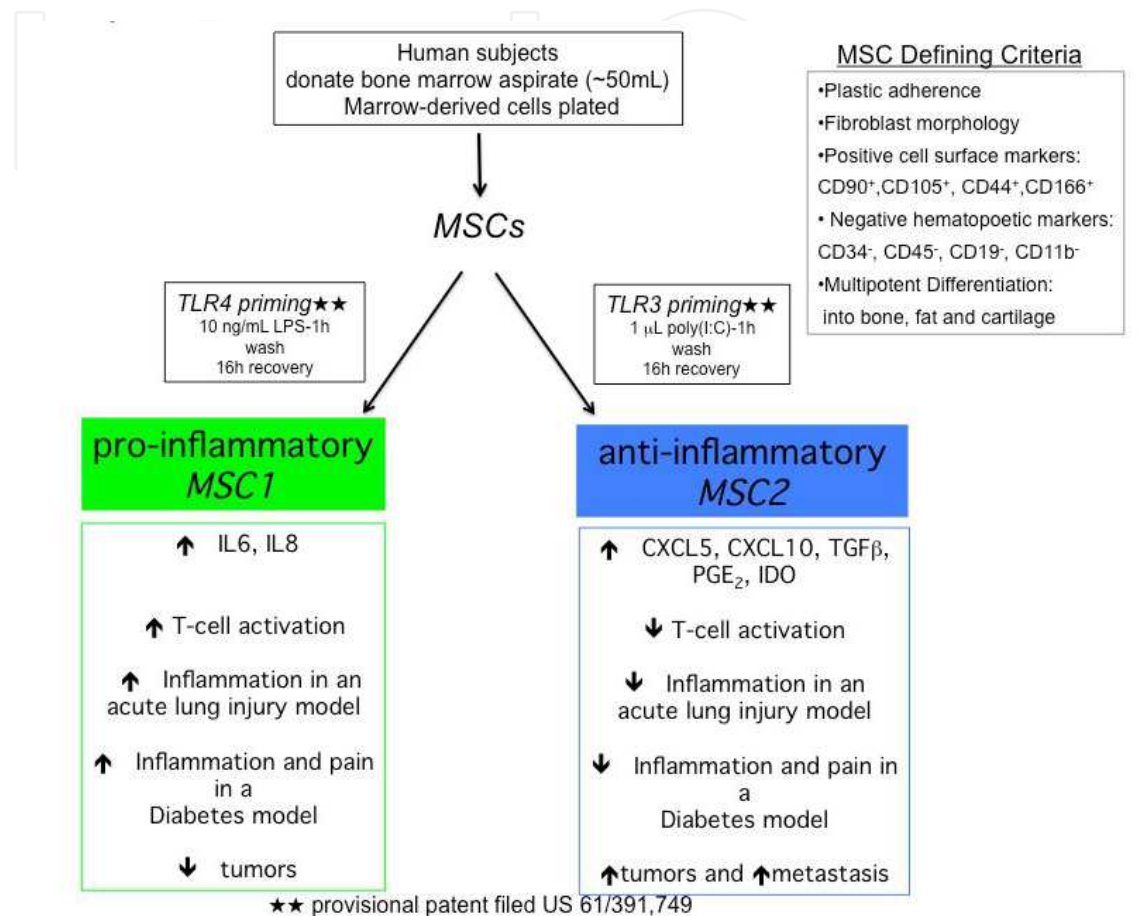


Fig. 2. Characteristics of the *MSC1* and *MSC2* Phenotypes. Short-term and low-level priming of TLR4 (left side) and TLR3 (right side) leads to the induction of heterogeneous hMSC preparations into a pro-inflammatory *MSC1* phenotype or an anti-inflammatory *MSC2* phenotype. (adapted from (Tomchuck, Zvezdaryk et al. 2008; Waterman, Tomchuck et al. 2010)).

#### 4.1 Evidence for *MSC1* and *MSC2*

Our previous work, as well as that of others, established that MSCs reside in TMEs or tumor stroma, provide structural support for the malignant cells, modulate the tumor microenvironment, and consequently promote tumor growth and spread. Therefore, gene-modified MSCs that can act as “Trojan horses” and deliver anti-cancer therapeutics into the tumor stroma are being evaluated as a promising new specific cell-based therapy for cancer. We also previously established that MSCs recruited to ovarian tumors by elevated secretion of LL-37 play a supportive role in ovarian tumor stroma. We found that specific induction of MSCs into *MSC1* causes the secretion of pro-inflammatory mediators rather than anti-

inflammatory ones, as well as promotes collagen rather than fibronectin deposition into the extracellular matrix (Figure 1)(Waterman, Tomchuck et al. 2010). Our preliminary studies support the notion that *MSC1* may be effective in new cell-based treatment of cancers. Indeed, ovarian cancer cell lines co-cultured with *MSC1* formed smaller tumor spheroids and had markedly reduced tumor colony forming potential; whereas, co-cultures with *MSC2* phenotype had the expected pro-tumor effect. Moreover, *MSC1*-treated ovarian cancer cells were less invasive than *MSC2*-treated ones in matrigel coated transwell migration assays. Pilot tests in murine ovarian cancer models were consistent with these findings. *MSC1* delivered in mice with established tumors had attenuated growth and spread. Mice treated with *MSC2* had larger and more metastatic tumors.

*MSC1* and *MSC2* therapy has been successfully tested in several animal disease models and has resulted in predictable inflammatory responses and distinct effects on tumor growth and spread (Table 5).

Animal Disease Model	MSC-based Therapy	MSC Dose (cells)	Treatment Frequency (Time of treatment)	Disease Impact	Length of study	Adverse Effects
1. LPS-induced Acute Lung Injury (ALI) (BalbC and C57BL/6J, n=12)	MSCs	0.5X10 <sup>6</sup>	1X (24hrs post-disease onset)	Mostly anti-inflammatory	1 week post-treatment	NONE
	<i>MSC1</i>	0.5X10 <sup>6</sup>	1X (24hrs post-disease onset)	Pro-inflammatory	1 week post-treatment	NONE
	<i>MSC2</i>	0.5X10 <sup>6</sup>	1X (24hrs post-disease onset)	Anti-inflammatory	1 week post-treatment	NONE
2. Streptozotocin-Induced Diabetes and neuropathic pain (C57BL/6J, n=30)	MSCs	1-3X10 <sup>6</sup>	3X (given in 10-day intervals post-disease onset)	Mostly anti-inflammatory	70 days post-treatment	NONE
	<i>MSC1</i>	1-3X10 <sup>6</sup>	3X (given in 10-day intervals post-disease onset)	Pro-inflammatory	70 days post-treatment	NONE
	<i>MSC2</i>	1-3X10 <sup>6</sup>	3X (given in 10-day intervals post-disease onset)	Anti-inflammatory	70 days post-treatment	NONE
3. Immune-incompetent human tumor xenografts (Balb scid and nude n=60)	MSCs	0.5X10 <sup>6</sup>	3X (given weekly post-disease onset)	Mostly anti-inflammatory	>120 days post-treatment	NONE
	<i>MSC1</i>	0.5X10 <sup>6</sup>	3X (given weekly post-disease onset)	Pro-inflammatory	>120 days post-treatment	NONE
	<i>MSC2</i>	0.5X10 <sup>6</sup>	3X (given weekly post-disease onset)	Anti-inflammatory	>120 days post-treatment	NONE
4. Immune-competent MOSEC (C57/BL6J n=20)	MSCs	0.5X10 <sup>6</sup>	3X (given weekly post-disease onset)	Mostly anti-inflammatory	>70 days post-treatment	NONE
	<i>MSC1</i>	0.5X10 <sup>6</sup>	3X (given weekly post-disease onset)	Pro-inflammatory	>70 days post-treatment	NONE

Table 5. Human MSC-based therapy of murine disease models.

Please NOTE that for all of the data presented MSCs represent conventionally prepared human MSCs, *MSC1* are defined as the hMSCs incubated for 1hr with 10 ng/mL LPS and washed prior to delivery. *MSC2* are defined as the hMSCs incubated for 1hr with 1 mg/mL poly(I:C) and washed prior to delivery (provisional patent filed US 61/391,749).

**Cancer models:** Pilot studies with the mouse ovarian cancer model (MOSEC) and with a xenograft model demonstrate our assertions. A single delivery of *MSC1*-based therapy resulted in slower growing tumors, whereas comparable therapy with MSCs or *MSC2* resulted in larger tumors and metastasis at the end of the study (day 65, Figure 3).

### *MSC1* do not support tumor growth whereas *MSC2* favor tumor growth and metastasis

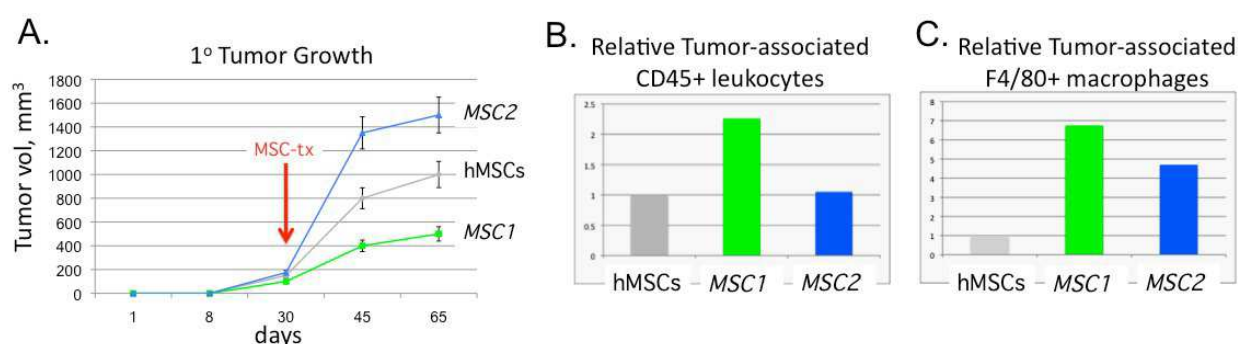


Fig. 3. *MSC1* do not support tumor growth whereas *MSC2* favor tumor growth and metastasis. The data show differences in tumor volume, CD45+leukocyte, and F4/80+ macrophage recruitment after the treatment of mice with established ovarian tumors, with human *MSC1*- and *MSC2*-based therapies. Methods The established syngeneic mouse model for epithelial ovarian cancer used is based upon a spontaneously transformed mouse ovarian surface epithelial cell (MOSEC) line ID8 that has been previously described (Roby, Taylor et al. 2000). 4-6 week-old female mice ( $n > 10$  mice/MSCTreatment) were injected subcutaneously (s.c.) in the right hind leg with  $1 \times 10^7$  MOSEC cells. At approximately 4 weeks a single dose of labeled human MSCs (hMSCs), *MSC1*, or *MSC2* ( $1 \times 10^6$ /per mouse) were injected intraperitoneally (IP) as indicated by red arrow  $\downarrow$ . (A.) Tumor growth was measured with callipers as standard at weekly intervals until day of mouse sacrifice (Day 65). Harvested tumors and metastasis were weighed, counted and processed for flow cytometry and immunohistochemical analysis (IHC, Coffelt et al., 2009). Metastasis was found only in *MSC2*-treated mice (data not shown). MSCs were detected by flow cytometry and IHC. All MSC-treated samples had similar detectable MSCs within the tumor tissue-trending towards more *MSC1* and *MSC2* measured than hMSCs: approximately 15-25 cells counted per 200X field after 24hr of MSC-treatment and 2-5 cells at time of tissue harvest (day 65, data not shown). Sectioned tumor sample slides were stained with murine CD45 (B.) or F4/80 (C.) antibodies and the number of positively stained immune cells per 200X field were scored as described previously (Coffelt et al., 2009). Data are expressed as average cells counted in 4 fields/slide relative to hMSC sample. Data indicate in vivo stability and predictably distinct effects by the *MSC1* and *MSC2*.

**ALI model:** In an established endotoxin-induced acute lung injury (ALI) mouse model, LPS, or endotoxin (0.1 mg/kg) was instilled intratracheally into adult Balb/C mice. After 24 hrs, mice were each treated with  $0.5 \times 10^6$  MSCs, *MSC1*, *MSC2*, or HBSS vehicle. To characterize inflammation, the lungs of the animals were lavaged and bronchioalveolar lavage fluid (BALF) was analyzed after 24 hr for changes in neutrophil/monocyte recruitment (myeloperoxidase activity), total cell content by flow cytometry, and lung integrity by total protein leaked into the BALF ( $n=12$ ). *MSC1*-therapy aggravated the disease and resulted in

increased neutrophil recruitment and more compromised lungs than the conventional MSC or MSC2 therapy.

**Diabetes Model:** Streptozotocin (STZ)-induced diabetic mice were procured from Jackson Laboratory (Bar Harbor, Maine). Blood glucose levels and animal weights were measured by standard methods. A month post STZ-injection, mice received intraperitoneally (IP)  $0.5 \times 10^6$  cells of MSCs, MSC1, MSC2, or HBSS vehicle for a total of 3 times in 10-day intervals. Established behavioral assays to evaluate hyperalgesia and allodynia were conducted one day prior to each MSC therapy, as well as prior to sacrifice. Inflammatory factors and immune cell changes were measured as before to characterize the treatment effects on inflammation ( $n=30$ ). Again, all indicators were consistent with enhanced inflammation by MSC1-treatment and an improvement of disease by the MSC2- or MSC-treated animals. *Manuscript in preparation.*

Additionally *in vitro* studies show divergent effects of MSC1 and MSC2 on cancer cells. Co-culture of various human cancer cell lines with MSC1 and MSC2 in Colony Forming Units (CFU) assays and 3-D tumor spheroid assays agree with the *in vivo* tumor models with different MSC1 and MSC2 treatment effects (Figure 4).

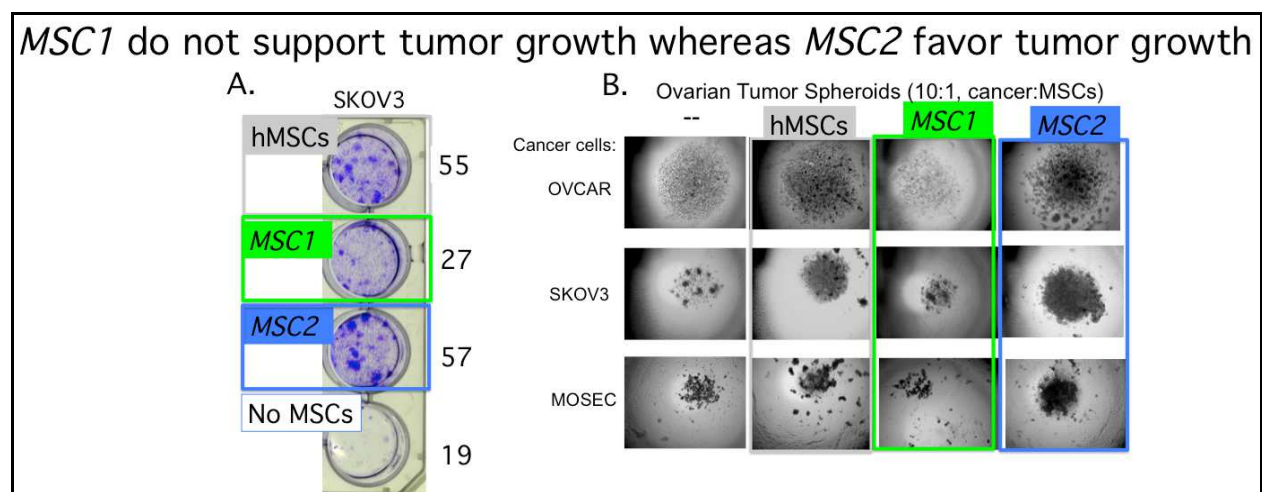


Fig. 4. MSC1 do not support tumor growth whereas MSC2 favor tumor growth: A. Data demonstrates that there are distinct effects on colony forming units (CFU) after coculture of different human cancer cell lines with untreated MSCs (hMSCs), MSC1, or MSC2. Methods: CFU assay was performed by culturing human tumor cells (200 cells/well) mixed with hMSCs, MSC1, or MSC2 (2 cells/well) at a ratio of 10 cancer cells per 1 MSC and plated in 24-well plates in growth medium supplemented with 10% FBS as indicated in figure. Cultures were grown for 14 days at 37°C in a humidified atmosphere of 5% carbon dioxide balance air. Growth medium was changed every 3-4 days. Colonies were visualized by staining with a crystal violet solution (0.5% crystal violet/10% ethanol). The resulting colonies were enumerated by the colony counting macro in ImageJ software, SKOV3-ovarian cancer cell lines. Micrographs of the stained plates are shown. Colony counts are at right. ( $n=8$ ) B. Data demonstrates that there are distinct effects on tumor spheroids after coculture of different cancer cell lines with unprimed MSCs, MSC1, or MSC2. Methods: Tumor spheroids were formed by culturing tumor cells (2000 cells/well) mixed without any other cells (--) or with hMSCs, MSC1, or MSC2 (20 cells/well) at a ratio of 10 cancer cells per

1 MSC and plated over 1.5% agarose in 96-well plates in growth medium supplemented with 10% FBS as indicated in figure. Cultures were grown for 14 days at 37°C in a humidified atmosphere of 5% carbon dioxide balance air. Growth medium was changed every 3-4 days. Micrographs shown represent 20X magnified field of the 96-well plate. Cancer cell lines used are: OVCAR-human ovarian cancer, SKOV3-human ovarian cancer cell lines, and MOSEC-murine ovarian surface epithelium carcinoma cells. Data indicate distinct effects by MSC1 and MSC2 on cancer cell growth and spread.

## 5. Conclusion

The unique pathology of individual tumors presents a huge problem for conventional mono-specific therapies. New approaches aiming at developing effective treatments against cancer include the use of MSC-based therapies. There are many features that make this new strategy attractive and feasible. First, MSC-based therapies are already in clinical use and thus far have not been associated with adverse effects. Second, MSCs can be easily expanded and stored without any impact to their capabilities—a phenomenon that has triggered the creation of many new biotech start-ups. Third, once delivered, MSCs preferentially home to tumors and affect tumor growth and spread. Fourth, MSCs from non-self (allogeneic) or autologous (self) hosts can be safely delivered since they do not elicit immunity. Lastly, pre-clinical studies have demonstrated efficacy with genetically-engineered MSCs that carry anti-cancer therapeutics that reached the tumors and prevented their growth.

MSCs targeted to cancers are expected to contribute many soluble factors such as mitogens, extracellular matrix proteins, angiogenic and inflammatory factors, as well as exosomes with as yet poorly defined potentials, once resident in the TME. MSCs are also expected to affect tumor-associated leukocytes either directly by cell-cell contact or indirectly by the secretion of trophic factors. MSCs are known to affect the proliferation and differentiation of dendritic cells, monocytes/macrophages, B and T cells, NK cells, and even mast cells. There has been a great deal of debate in the field in trying to assert whether MSCs resident in the TME contribute to tumor growth and spread or prevent it, and if so, by what mechanisms. Many reasons have been advanced to explain the contradictory MSC role in cancer including the heterogeneity of MSC preparations, the age or health of the MSC donor, and the experimental model or condition, to name a few. Our group has suggested a new paradigm for MSCs that we believe will help resolve some of the conflicting issues. The induction of MSCs into uniform and consistently acting pro-inflammatory *MSC1* or anti-inflammatory *MSC2* phenotypes should provide convenient experimental tools that dissect the potential pro- and anti-tumor contributions of MSCs. MSC-based therapies stand to revolutionize medicine with the myriad ways that they can be manipulated and guided to reach pathologic tissue sites such as tumors. The continued investigation of these cells will ensure safe and effective therapy of human disease.

## 6. Acknowledgment

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## **Tumor Microenvironment and Myelomonocytic Cells**

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Tumor microenvironment represents an extremely dynamic niche shaped by the interplay of different cell types (e.g. tumor cells, stromal cells), their soluble products (e.g. cytokines, chemokines and growth factors) and varied physico-chemical conditions (e.g. low oxygen concentration or hypoxia). Recent studies have identified myelomonocytic cells as key players in regulating the tumor microenvironment and hence, tumor progression in a variety of cancers. In view of these findings, the present book attempts to provide a comprehensive account of the diversity of tumor microenvironment across different cancers and how myelomonocytic cells have taken the center-stage in regulating this niche to direct cancer progression. A better understanding of the myelomonocytic cells and the mechanisms by which they regulate cancer progression will open new vistas in cancer therapeutics.

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