The known molecules involved in MSC homing and migration

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Abstract
Mesenchymal stem cells (MSCs) are multipotent adult stem cells and known as promising candidates for use in cell-based therapy. Following systemic administration, MSCs have been shown to accumulate at sites of inflammation and injured tissue. There are several factors regarding the MSC trafficking and homing in MSC therapy. This review describes the known molecules in MSC migration and homing.

Introduction
Mesenchymal stem cells are non-hematopoietic stem cell population that was first characterized by Friedenstein and colleagues [1]. MSCs population approximately were estimated 1 in 10,000 nucleated cells in bone marrow [2]. MSCs have a supportive function for the haematopoiesis in the bone marrow and prepare a niche for maturation of hematopoietic progenitor cells [3]. MSCs are identified by the expression of a number of surface markers including, CD14, CD11b, CD79a or CD19, and HLA-DR; (c) Differentiate into osteoblasts, adipocytes, and chondrocytes in vitro [13]. They also do not express hematopoietic surface markers CD45, CD34, CD14, CD11b, CD79a, and CD19. They also do not express the costimulatory molecules CD80, CD86, or CD40 [4]. Stro-1 [5], CD271(low-affinity nerve growth factor receptor) [6], stage specific embryonic antigen-4 (SSEA-4) [7], CD146 [8], CD49a [9], CD106, and leptin receptor [10] are cell surface antigens that have been reported to enrich homogeneous MSCs. It is shown that MSC could be enriched through specific markers, for example, they are enriched from peripheral and the UC blood through selecting CD133, or from BM through selecting stage-specific embryonic antigen (SSEA)-1, SSEA-4 [11]. Analysis of cell surface biomarkers revealed that MSC with strong self-renewal capability preferentially express Stro-1 and PDGF-α (platelet-derived growth factor receptor alpha), and they also expressed the highest level of TWIST-1 and DERM-1 mRNA transcripts [12]. The Mesenchymal and Tissue Stem Cell Committee of the International Society for Cellular Therapy (ISCT) provided a set of standards to define MSCs include: (a) Remain plastic-adherent under standard culture conditions; (b) Express CD105, CD73, and CD90, and lack expression of CD45, CD34, CD14 or CD11b, CD79a or CD19, and HLA-DR; (c) Differentiate into osteoblasts, adipocytes, and chondrocytes in vitro [13].

MSCs migration and homing
MSCs have been shown to mobilize into peripheral blood and migrate to other tissues in response to injuries and various types of diseases such as acute burns, skeletal muscle injury, chronic hypoxia, inflammation, heart diseases, neurological disorders, rheumatoid arthritis, graft-versus-host disease and etc [14,15]. Although there are few studies about the mechanisms of MSC homing, several agents have suggested the homing mechanisms (Table 1).

Mechanism of MSC migration and homing
Role of Adhesion molecule interactions in MSC homing
MSCs movement in the bloodstream and transendothelial migration is one of the most important stages of homing. It is assumed that MSCs utilize the same mechanism as leukocyte migration to move into the bloodstream. Adhesion molecules are important for MSC trafficking. Integrin blocking, and knockout studies have supported that MSCs migration is related to their interactions with integrins and selectins. MSCs express adhesion molecules, including: integrins, selectins, and

Abbreviations: VLA-4: Very Late Antigen-4; MSCs: Mesenchymal Stem Cells; SSEA: Stage Specific Embryonic Antigen; PDGF-α: Platelet-Derived Growth Factor Receptor- Alpha; TWIST: Twist-Related Protein 1; HLA-DR: Human Leukocyte Antigen Class II; ICAM: Intercellular Adhesion Molecule 1; VCAM: Vascular Cell Adhesion Molecule 1; ALCAM: Activated Leukocyte Cell Adhesion Molecule; MCP: Monocyte Chemotactreactant Protein-1; MIP-1a: Macrophage Inflammatory Protein-1 Alpha; RT-PCR: Real-Time Polymerase Chain Reaction (Real-Time PCR); BM: Bone Marrow; UC: Umbilical Cord; SDF-1: Stromal Cell-Derived Factor-1; TNF-α: Tumor Necrosis Factor- Alpha; IGF: Insulin-Like Growth Factor-1; PI3K/Akt: Phosphatidylinositol 3-Kinase/Protein Kinase B; MMP: Matrix Metalloproteinase; MT1-MMP: Membrane-Typed-1 Matrix Metalloproteinase; sRNA: Small Interfering RNA; TIMP: Tissue Inhibitor of Metalloproteinases; PDGF-α: Platelet-Derived Growth Factor Receptor- Alpha; PlGF: Placenta Growth Factor; TGF-α: Transforming Growth Factor- Alpha; IL-1: Interleukin-1.

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chemokine receptors, which are involved in the tethering, rolling, adhesion, and transmigration of leukocytes from the bloodstream into tissues. MSCs express integrin molecules, such as α1, α2, α3, α4, α5, av, β1, β3, and β4. MSCs also express other adhesion molecules, including: VCAM-1, ICAM-1, ICAM-3, CD166 (ALCAM), endothelin/CD105. It has been estimated that VLA-4 is expressed in approximately 50% of hMSCs [1,16]. Although the study has shown that MSCs and leukocytes likely utilize some similar molecules and mechanisms for transmigration, there are some specific differences between MSCs and leukocytes in employing adhesion molecules. Studies have revealed β1 integrins possess a crucial role in MSC rolling and attachment, but many of β2 integrins are not expressed on MSC surface [16,17]. Some studies have shown attachment of MSC to endothelial layer has resulted from P-selectin interaction. On the other hand, it has been shown that MSC do not express PSGL-1 and an alternative ligand CD24, and their endothelial interaction is dependent on P-selectin. It is assumed that MSC might express new carbohydrate ligands to bind P-selectin expressed on endothelial cell. Also, data have shown E-selectin, and L-selectin, which function in the initial leukocyte rolling stage, do not or barely express on MSCs and their contributions to MSC trafficking are not remarkable in comparison with P-selectin [16]. It has been reported rolling mediate due to VLA-4/VCAM-1 interactions and they make firm adhesion of MSC to endothelial cells. To MSCs express VLA-4 and VCAM-1 adhesion molecules and it has been shown that VLA-4 and VCAM-1 are important receptors to transendothelial MSCs migration. Observations have shown anti-VCAM-1 antibody reduced attachment of rat MSC to endothelial cells [17]. Steingen has reported that MSC could transmigrate through nonactivated endothelium via VLA-4/VCAM-1 and they tend to integrate with the endothelial layer rather than undergoing complete diapedesis. Among the integrins family, α4β1 integrin, which is a cell surface heterodimer and mediates cell-cell and cell-extracellular matrix contact, plays a key role in adhesion, migration and chemotaxis. Since MSC transendothelial migration is not completely blocked by the anti-VLA4 antibody and anti-VCAM1 antibody, it is suggested other integrins are involved in the MSC migration process [18]. The aforementioned data show that different adhesion molecules are expressed by MSC and involved in MSC migration and homing.

### Role of Chemokine and Chemokine Receptors in MSC Homing

Although integrins and selectins play role in the transmigration of MSC, chemokines released from tissues and endothelial cells could promote activation of ligands that are involved in attachment, migration, chemotaxis and maintenance of MSC in target tissues. It has been shown that MSCs are able to migrate into inflamed tissues in response to chemokines and chemokine receptor signals that are induced under inflammatory conditions [19]. MSCs express a wide range of chemokine and chemokine receptors that have been shown to play a role in MSC homing. Some reports have shown expression of chemokine receptors; including, CCR1, CCR2, CCR4, CCR6, CCR7, CCR8, CCR9, CCR10, CXCR1, CXCR2, CXCR3, CXCR4, CXCR5, CXCR6 on MSCs and have reported functional roles of some of these chemokine receptors in MSC migrations [20–23]. CXCR1, CXCR2, CXCR4, CCR1, CCR2, IL-8, MIP-1α, MCP-1 are involved in MSCs migration to damaged tissue [24]. Recent studies showed that 43–70% of MSCs expressed functional CCR1, CCR2, CCR9, CXCR4, CXCR5, and CXCR6 [20,25]. Using RT-PCR the data showed that mRNA expression of CCR1, CXCR4, CCR7, CXCR6 and CXCR3 is low in adipose tissue-derived MSCs and nearly undetectable in most BM-MSCs [19]. The majority of these studies have detected expression of chemokine receptors by flow cytometry analysis, real-time polymerase chain reaction, immunohistochemistry, and chemotaxis assay. Ponte and colleagues [26] demonstrated TNF-α could increase expression of CCR2, CCR3, and CCR4 but CXCR4 expression was not affected. The expression of CXCR4 plays a critical role in migration and homing of MSCs. Wynn and colleagues have reported less than 1% of MSCs express CXCR4 on the cell membrane, although intracellular expression of CXCR4 (83–98%) was noted [27]. Pre-treatment of BM-MSC with TNF-α significantly increases CXCR4 expression and the capacity of MSC migration, especially in response to SDF-1 [28]. It is known that IGF-1 not only has individually induced MSC migration, but also it has caused CXCR4 upregulation. Moreover, IGF-1 could induce MSC migration in response to SDF-1. This study also indicated that IGF-1 increases MSC migration through CXCR4 chemokine receptor signaling in a PI3/Akt dependent manner [29]. Also, treatment of BM-MSC with some cytokines such as, FGF-2, SCF, IL-6, HGF and IL-3 has significantly upregulated intracellular and extracellular expression levels of CXCR4 [30]. Studies have shown SDF-1/CXCR4 axis performs the important role in migration of MSC to the bone marrow. CXCR4 responses to SDF-1 in a dose dependent manner [31]. Su et al. [32] also indicated that the expression level of SDF-1α controls the homing of MSCs. Therefore, it is likely that under chemokine stimulation condition intracellular CXCR4 translocates to the cell surface and contributes to the migration of MSC to target site.

The MCP-1 expression is upregulated at the inflammation site and known as homing chemokine. Previous studies indicated MCP-1 contributed to monocyte recruitment to inflammation site. MSCs express CCR2, which is a MCP-1 receptor [33]. Transwell migration assay showed MCP-1 induced migration of mouse and human bone marrow derived MSCs [34,35]. In vivo studies have shown the anti-CCR2 antibody and blockade of downstream CCR2 signalling pathway could inhibit MSCs migration [36]. On the other hand, Ring et al. [20] have reported human BM-MSC do not significantly migrate in response to MCP-1 gradient. MIP-1α is the other chemokine which is possible to play role in MSC migration. MIP-1α induces proinflammatory immune cells recruitment to the inflammation site and regulates their transendothelial migration [37]. CCR1 is known as a MIP-1α receptor and is expressed on MSC surface [25]. The transwell migration assay has shown the presence of MIP-1α induces increased migration in cultured BM-MSC [35]. It is suggested MIP1α as a soluble factor might be involved in controlling of the MSCs homing process. It is shown IL-8 upregulation in the injured tissue sites may be causing MSC migration to the injured site. CXCR1/CXCR2 are IL-8 ligands on the MSC surface and MSC chemotactic response to an IL-8 has been demonstrated

### Table 1. The molecules involved in MSC migration and homing

| Molecules                                | Migration stage                        | References |
|------------------------------------------|----------------------------------------|------------|
| Adhesion molecules                       | VLA-4, VCAM-1, ICAM-1, P-selectin      | Rolling and transendothelial migration [16,17] |
| Chemokines/chemokine receptors           | IL-8, MIP-1α, MCP-1, SDF-1             | Chemotaxis and traffic [18-21] |
| Cytokines/growth factors                 | TGF-β, TNF-α, IL-1β, IL-6, IL-3, FIT3-L, IGF-1, SCF, HGF, PDGF, PDGF | Chemotaxis and traffic [22-25] |
| Matrix metalloproteinases                 | MMP-1, MMP-2                           | Invasion [26] |
Role of Matrix metalloproteinases in MSC homing

MMPs are a zinc-dependent proteolytic enzyme family regulating ECM (extracellular matrix) degradation. Different MMPs and their signalling pathways have been shown to affect differentiation, migration, angiogenesis, and proliferation of MSCs. Migration and recruitment of expanded MSCs to damaged tissue are facilitated through expression of CXCR4, MMP-2 and MT1-MMP [38]. The MMP-2 expression is associated with MSCs transendothelial migration, although it is known the increasing MSCs culture confluency diminished MSCs transendothelial migration through the inhibition of MMP-2 siRNA and the increase in TIMP-3 expression [39].

Role of Cytokines and growth factors in MSC homing

The active role of IL-6, PDGF, PDGFR-α (platelet-derived growth factor receptor α), PDGFR-β, PIGF (placenta growth factor), vascular endothelial growth factor receptor 1 (Flt-1) and IGF-1 has been known in the migration of BM-MSCs [24]. Several studies showed that PDGF-Rs are highly expressed on the BM-MSC surface and PDGF induces BM-MSC migration. The transwell migration assay has shown, PDGF is a more potent MSC chemoattractant cytokine than SDF-1 and MCP-1 [40]. Inflammatory cytokines, including; TGF-α, IL-1β and TNF-α stimulate MMP production by the MSCs and result in induction of potent chemotactic migration through the extracellular Matrix [41].

Conclusion

It is well known that MSCs have a capacity to migrate and home in injured and inflamed tissues. Different molecules were demonstrated to play a role in MSC migration and homing, but the exact mechanism of homing is still unclear. Understanding how MSCs migrate and home into target tissues could be improve future MSC-derived therapies.

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