

Stem Cells in Airway Smooth Muscle

State of the Art

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Very little is known regarding the function, origin, and turnover of airway smooth muscle (ASM). In this article, we discuss the embryological development of ASM, and provide information regarding candidate mesenchymal ASM progenitor cell populations specifically in relation to airway remodeling. This review also highlights the current limitations in studying ASM biology, and underscores the need for novel molecular tools and markers that will refine our understanding of this cell type in lung homeostasis and disease.

Keywords: stem cells; smooth muscle; progenitor cells; fibrocyte; myofibroblast

OVERVIEW

Very little is known regarding the role of stem cells in airway smooth muscle (ASM). While there has been considerable research on the contributions of local proliferation and circulating mesenchymal stem cells to hyperplasia of vascular smooth muscle (VSM) in both pulmonary hypertension and atherosclerosis, there is no comparable body of work for ASM in diseases such as asthma. One of the greatest difficulties in understanding the role, if any, of stem cells in ASM is the role of ASM itself. To date there is no known “beneficial” or physiologic function to ASM (1).

There are several theories as to the possible ontology of ASM muscle. (1) The rhythmic peristalsis of ASM, in the developing fetal lung generates distending pressures, which promote development and maturation of airways and alveoli (2). (2) ASM may change the resistance of airways in parallel circuits to improve \dot{V}/\dot{Q} matching and or decrease dead space (3). (3) ASM may increase the velocity of gas movement during cough to promote expulsion of foreign bodies (4). (4) ASM may serve to balance hysteresis between small airways and alveolar units (5). (5) ASM may simply be an “evolutionary oversight,” that is, a vestige of the lung’s origin from an organ, the foregut, already programmed to develop smooth muscle, with no true physiologic purpose.

When one attempts to study such a system, one usually begins teleologically, by working backward from its function, to understand the source; in this case, however, this approach is limited. This is further confounded by the lack of specific markers to distinguish ASM from VSM in the lung. While there has been considerable research into the role of stem cells in VSM, when one looks for similar progenitors in ASM, it becomes difficult to tease out whether or not what you are looking at is truly an airway or vascular cell. With these caveats in mind, we can proceed with some basic principles. (1) There is now an emerging

understanding of the source and maturation of both local and circulating mesenchymal stem/progenitor cells with the capacity and or proclivity to mature into smooth muscle *in vitro* (6–10). (2) Asthma is a disease of increased ASM mass, due to both (in varying degrees) hypertrophy and hyperplasia (11–16). (3) In disease states, circulating and local mesenchymal/progenitor cells play a role in VSM accumulation, but related data in the case of asthma is not as forthcoming (10, 17–19).

EMBRYOLOGY

The precise origin of ASM in the developing lung is not known; accumulated data suggest, however, that these cells originate from the primitive embryonic lung mesenchyme. In the mouse, smooth muscle actin-positive cells associated circumferentially with the early trachea and mainstem bronchi can be observed at the earliest stages of lung development (20). This nascent smooth muscle cell compartment is likely already enervated, consistent with observations demonstrating the presence of pulsatile airway contractions in the early fetus (2, 20, 21).

The lack of distinguishing markers has hampered efforts to absolutely clarify the ontological relationship between VSM and ASM, and to elucidate whether these cell types have distinct differentiation programs. Interestingly, the transcription factor GATA-5 is selectively expressed in ASM during late gestation; the significance of this observation has not been further explored (22). Distinct cell differentiation programs and/or cell origins are inferred by the phenotypes of mice that carry select mutations in growth factor signaling molecules. In this regard, hypomorphic FGF-10 mice display selective loss of ASM mass. On the other hand, deletion of Wnt-7b appears to selectively disrupt development of VSM in pulmonary arteries (23, 24).

Since ASM accumulation is ongoing and coupled to branching morphogenesis, it has been suggested that signals originating in the airway epithelium regulate smooth muscle cell recruitment, differentiation, and organization. Consistent with this speculation, deletion of epithelial derived sonic hedgehog is associated with reduced ASM mass (25). Several studies have shown that components of the epithelial basement membrane, particularly laminin-2, may play a role in controlling ASM differentiation by facilitating attachment and spreading of primitive mesenchymal cells (22, 26). These changes in cell shape induce translocation of serum-response factor (SRF) from the cytoplasm to the nucleus (27). Changes in SRF localization have important functional consequences, since SRF is one of the key transcription factors involved in up-regulating expression of smooth muscle-related genes (27). Stretch has also been shown to promote bronchial smooth muscle differentiation by activating an SRF-dependent pathway (27, 28).

ORIGIN OF SMOOTH MUSCLE DURING AIRWAY REMODELING

Asthma is a disease estimated to affect 10% of people in the United States, causing considerable morbidity and mortality. Although inflammation is undoubtedly a cornerstone of the disease, it is clear that structural changes referred to cumulatively

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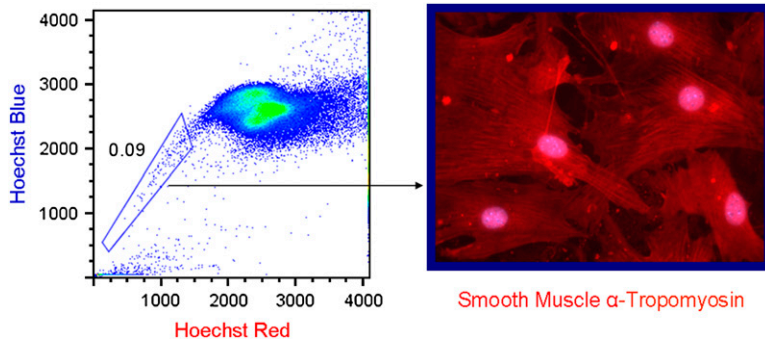


Figure 1. Lung side population (SP) cells give rise to smooth muscle. (Left) Density dot plot of Hoechst stained lung digests. The boxed area represents cells (0.09%) that efflux dye (SP cells). (Right) The CD45⁻ CD31⁻ subtype differentiates into α -tropomyosin-positive cells after collection and culture in smooth muscle media. The CD45⁺ subgroup and the CD45⁻ CD31⁺ subgroup do not display this capacity.

as “airway remodeling” contribute to the asthmatic diathesis. This airway remodeling consists of thickening of the airway wall, subepithelial fibrosis, smooth muscle myocyte hyperplasia and hypertrophy, and myofibroblast hyperplasia (29). These processes contribute to airway hyperresponsiveness and are associated with disease severity and decline of lung function (FEV₁) (29). The origin of smooth muscle and fibrogenic cells responsible for this pathology is not currently known. In 2001 Johnson and coworkers showed that ASM obtained from endobronchial biopsies in individuals with asthma proliferated much more quickly (~300%) than did those from healthy control subjects; these findings could be accounted for by expansion of local ASM progenitors or hyperplasia of already differentiated cells (16).

One potential progenitor for ASM is a circulating mesenchymal stem cell (MSC), first identified in outgrowths of adult murine bone marrow in 1976 (30). These cells have historically been isolated by *in vitro* culturing and expansion on plastic. They are characterized by expression of select surface markers such as Sca-1 (stem cell antigen 1) and by the lack of hematopoietic markers. When placed in appropriate media MSCs can differentiate into cartilage, bone, fat, or smooth muscle (6, 8, 31). In models of skeletal and cardiac injury, such marrow-derived cells have been found to contribute to the reconstitution of damaged muscle (32–34). Myofibroblasts, a cell type that actively produces connective tissue, have also been shown to be marrow derived, in part, in organ injury (35–37). Further, a circulating cell may contribute to smooth muscle cells during the healing phase of vascular wall injury (38).

To date, the study of circulating mesenchymal progenitors has been hampered by the lack of an effective surface marker for their isolation and characterization in the blood and marrow directly, forcing investigators to rely on less reliable techniques such as staining for internal markers and selection of cell types after culturing. A recent publication by Gang and colleagues has shown that the stage-specific early embryonic antigen 4 (SSEA-4) may serve as an effective and easily applied surface marker for MSCs obtained from both human and mouse bone marrow (8). Clonal analysis of CD45⁻ SSEA-4⁺ cells showed that the majority (~70%) of these cells had the capacity to differentiate into bone, fat, and cartilage cell types (8). While the authors did not specifically look at the ability of these cells to differentiate into smooth muscle, or characterize them in the peripheral circulation, this begs the question of their potential contribution to ASM accumulation in asthma.

Another potential source of ASM is the circulating fibrocyte, first described in 1994 in an article by Bucala and coworkers wherein they identified a circulating mesenchymal cell that could be cultured from the blood of mice and humans (39). These cells are isolated by plastic-adherent culture of peripheral blood mononuclear cells, express the panhematopoietic marker CD45 and the hematopoietic stem cell antigen CD34, and

synthesize collagen-I (39–44). Fibrocytes enter sites of injury localizing to areas of matrix formation *in vivo* (44). A progenitor phenotype is suggested by their ability to proliferate in culture and to differentiate after injection, or under specific culturing conditions, into α -smooth muscle actin⁺ (SMA) cells that secrete matrix proteins (40, 45, 46). Whether or not the fibrocyte is the definitive circulating progenitor cell or is merely a subtype of a broader category of mesenchymal progenitor cells is not yet clear. It is important to note, however, that classical mesenchymal progenitor cells do not express hematopoietic markers.

The fibrogenic potential of fibrocytes and the mechanisms responsible for their recruitment to lung tissue are of interest. One study showed that circulating fibrocytes from humans and mice express the chemokine receptor CXCR4 and migrated in response to its cognate ligand, CXCL12 (47). Notably, CXCR4⁺ fibrocytes traffic into murine lungs during bleomycin challenge; maximal recruitment directly correlated with increased collagen deposition. Furthermore, a CXCL12-neutralizing antibody inhibited recruitment of fibrocytes and attenuated pulmonary fibrosis, suggesting a pivotal role in the fibrotic response. Further, in an ovalbumin murine model of asthma, infused fibrocytes differentiated into lung myofibroblasts and smooth muscle cells beneath the bronchial epithelium (45). These cells were rapidly recruited to bronchial tissue after allergen exposure and could be re-isolated. Although freshly purified circulating fibrocytes do not express SMA, expression of SMA was induced in cells that engrafted in the bronchial wall. Interestingly, SMA is induced in human fibrocytes when cultured with endothelin-1 (ET-1) and transforming growth factor- β (TGF- β)—two fibrogenic cytokines up-regulated in the airways of patients with asthma (45).

In our own lab we have made use of the vital dye Hoechst 33342, which is preferentially effluxed out of stem cells by the action of the ABC transporter BCRP-1 (breast cancer resistance protein 1) (6, 48–50). When performing flow cytometry on cells stained with Hoechst, one finds a population with little or no dye that has a characteristic fluorescence pattern off to the side (so called side population or SP cells). We found that the lung SP is composed of a heterogeneous population that can be distinguished by the presence or absence of the panhematopoietic marker CD45. The CD45⁺ lung SP population behaves like hematopoietic stem cells (HSCs), possessing the ability to reconstitute the bone marrow of radio-ablated hosts (49). The CD45⁻ SP has at least two subpopulations: a CD31⁺ and -negative population (49). Like marrow MSCs, the lung CD45⁻ CD31⁻ SP cells can give rise to a variety of differentiated mesenchymal cell types *in vitro*, including smooth muscle (Figure 1) (6). The rarity of these cells, along with the lack of specific markers, has hampered efforts to specifically identify their site(s) of localization in the adult lung. Our data indicate that these cells are resident in the embryonic and adult

lung and not likely derived from the blood. Whether these cells contribute to airway remodeling is uncertain.

CONCLUSIONS AND FUTURE DIRECTIONS

Overall, there is a fundamental lack of knowledge regarding the function of ASM in homeostasis, and the source and mechanism of increased ASM mass in disease states. This state of affairs relates directly to the lack of definitive markers that distinguish ASM from VSM in the lung. The identification of such markers would facilitate our understanding of the basic biology and ontogeny of ASM, and the development of meaningful genetic models. Clarifying whether ASM cells originate from a defined progenitor or through expansion of differentiated cells may also become possible if specific markers are available; armed with this understanding, this knowledge would inform and guide new treatment strategies for airway remodeling.

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