



## Review

## Tissue-resident dendritic cells and diseases involving dendritic cell malfunction☆



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## ABSTRACT

Dendritic cells (DCs) control immune responses and are central to the development of immune memory and tolerance. DCs initiate and orchestrate immune responses in a manner that depends on signals they receive from microbes and cellular environment. Although DCs consist mainly of bone marrow-derived and resident populations, a third tissue-derived population resides the spleen and lymph nodes (LNs), different subsets of tissue-derived DCs have been identified in the blood, spleen, lymph nodes, skin, lung, liver, gut and kidney to maintain the tolerance and control immune responses. Tissue-resident DCs express different receptors for microbe-associated molecular patterns (MAMPs) and damage-associated molecular patterns (DAMPs), which were activated to promote the production of pro- or anti-inflammatory cytokines. Malfunction of DCs contributes to diseases such as autoimmunity, allergy, and cancer. It is therefore important to update the knowledge about resident DC subsets and diseases associated with DC malfunction.

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## Contents

1. Introduction . . . . .	2
2. Tissue-resident dendritic cells (DCs) . . . . .	2
2.1. Skin-resident DCs . . . . .	2
2.2. Blood-resident DCs . . . . .	2
2.3. Lung-resident DCs . . . . .	3
2.4. Liver-resident DCs . . . . .	3
2.5. Gastrointestinal tract-resident DCs . . . . .	4
2.6. Kidney-resident DCs . . . . .	4
2.7. Spleen-resident DCs . . . . .	4
2.8. Lymph node (LN)-resident DCs . . . . .	4
3. Diseases involving tissue-resident DCs . . . . .	5
3.1. Allergic asthma . . . . .	5
3.2. Autoimmune diseases . . . . .	5
3.2.1. Rheumatoid arthritis (RA) . . . . .	5
3.2.2. Psoriasis . . . . .	6
3.2.3. Systemic lupus erythematosus (SLE) . . . . .	6
3.3. Sarcoidosis . . . . .	8
3.4. Inflammatory bowel diseases (IBDs) . . . . .	8
3.5. Infectious diseases . . . . .	8

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4. Cancer . . . . . 9

4.1. Tumor infiltrating DCs initiate the antitumor immune responses . . . . . 9

4.2. Apoptosis of DCs in tumor microenvironment . . . . . 9

4.3. Dysfunctional DCs in tumor microenvironment . . . . . 9

4.4. DC-based vaccines in cancer immunotherapy . . . . . 10

4.4.1. Design of DC-based vaccines . . . . . 10

4.4.2. Route of delivery for antigen-loaded DC vaccines . . . . . 10

4.4.3. Lymph node homing of DC-based vaccines . . . . . 10

4.4.4. Application of DC-based vaccines . . . . . 10

5. Translating DC biology into medicine . . . . . 11

Conflict of interest . . . . . 11

Acknowledgement . . . . . 11

References . . . . . 11

**1. Introduction**

DC development takes place in the bone marrow and is a continuous process due to the requirement for replenishment of mature DCs in peripheral tissues. DCs can be divided into at least five broad groups based on phenotypic, functional and developmental criteria (Table 1). These groups include plasmacytoid DCs (pDCs), migratory and lymphoid tissue-resident CD8<sup>+</sup> DC-like DCs, migratory and lymphoid tissue-resident CD11b<sup>+</sup> DCs, Langerhans cells and monocyte-derived DCs, which are derived from a range of progenitors [1–4]. Langerhans cells are generated from fetal liver and yolk sac during embryonic development [5,6], whereas common DC progenitors (CDPs) give rise to plasmacytoid DCs and other conventional DC subsets. Pre-DCs are downstream of CDPs and have lost the capacity to generate pDCs, but are capable of generating lymphoid tissue-resident CD8<sup>+</sup> DCs and CD11b<sup>+</sup> DCs [7] possibly with their migratory counterparts referred to as CD103<sup>+</sup> and CD11b<sup>+</sup> DCs, respectively, found in many tissues [8,9]. pDCs can also be derived from lymphoid-primed multipotent progenitors (LMPPs) of bone marrow origin [10].

The capacity of DCs to capture antigens from the environment is increased even when there is no overt infection or inflammation, probably allowing for silencing the immune system in the presence of harmless environmental antigens. In response to danger signals such as bacterial and viral PAMPs, DCs rapidly promote T cell-mediated immunity to selectively eliminate infected cells. After capturing antigens, DCs migrate through lymphatics to reach secondary lymphoid organs, where DCs present processed antigenic peptides to stimulate naïve T cells in the context of MHC molecule. There also is evidence showing steady-state migration of DCs into lymph nodes under physiological conditions, which may serve to generate tolerant T cells to self and non-dangerous antigens.

Viewing the complexity of DC system accumulated during the past decades, the aim of this review is to provide an overview of the recent developments in the understanding of the distribution of tissue resident DC subsets in different organs and the relevance of resident DCs to the pathogenesis of human diseases, including auto-immunity, inflammation, allergic diseases, and cancer.

**Table 1**  
Mouse DC subtypes.<sup>a</sup>

DC subsets	Phenotype
Plasmacytoid DCs	CD11c <sup>int</sup> , B220 <sup>+</sup>
CD8 <sup>+</sup> DC-like DCs	CD8α <sup>+</sup> , CD11b <sup>lo/-</sup> , CD103 <sup>+/+</sup> , Langerin <sup>+/+</sup> , DEC205 <sup>+</sup> , Sirpα <sup>+</sup> , Clec9A <sup>+</sup> , XCR1 <sup>+</sup>
CD11b <sup>+</sup> DCs	CD4 <sup>+/+</sup> , CD8 <sup>-</sup> , CD11b <sup>+</sup> , Sirpα <sup>+</sup> , DEC205 <sup>-</sup>
Langerhans cells	CD11b <sup>int</sup> , EpCAM <sup>hi</sup> , Langerin <sup>+</sup> , DEC205 <sup>+</sup> , CD103 <sup>-</sup>
Monocyte-derived DCs	CD11b <sup>+</sup> , DEC205 <sup>+</sup>

<sup>a</sup> Ref. [1].

**2. Tissue-resident dendritic cells (DCs)**

*2.1. Skin-resident DCs*

DCs in the epidermis and dermis recognize invading pathogens and are dedicated antigen-presenting cells (APCs) to initiate both innate and adaptive immunity. Early studies revealed two main populations of DCs present in the normal skin: epidermal Langerhans' cells (LCs) and dermal (or interstitial) DCs (DDCs). They both express CD11c, MHC-class I, MHC II, CD1b, CD45, CCR6, IFN-γR, IL-1R1, IL-1R2, TNFR-2 and CD32. LCs also are CD324<sup>-</sup>, Birbeck granule- and CD207-positive; while DDCs do not express CD324, Birbeck granule and CD207, but CD11b, CD1d, CCR2, IL-2R2a, IL-7R, IL4R, CD64, CD206, CD209 and Factor XIII. However, DDCs were recently further divided into CD11b<sup>+</sup> and CD103<sup>+</sup> subsets (Or XCR1<sup>+</sup> DCs). Mouse CD11b<sup>+</sup> DCs are further divided into monocyte-derived counterpart (CD14<sup>+</sup> DCs in humans) and pre-DC-derived counterpart (CD1c<sup>+</sup> DCs in humans). In human skin, DCs comprise up to four different subsets: Langerhans cells, CD1c<sup>+</sup> DCs, CD14<sup>+</sup> DCs and a newly identified CD141<sup>+</sup> subset [11]. Therefore, Langerhans cells are found in human and mouse: human CD141<sup>+</sup> DCs match mouse CD103<sup>+</sup> dermal DCs and human CD1c<sup>+</sup> DCs of pre-DC origin (blood borne precursor) and CD14<sup>+</sup> DCs of monocyte origin combine to correspond to mouse CD11b<sup>+</sup> dermal subset DCs. Mouse dermis additionally contains a minor population of cDCs known as double-negative DCs that are XCR1<sup>-</sup> CD207<sup>-</sup> and express very low levels of CD11b [12,13]. This population expresses a DC lineage-specific transcription factor, ZBTB46, and the development depends on IFN-regulatory factor 4 (IRF4) and FLT3L. In mouse, migratory counterpart of double-negative DCs is found in the skin-draining lymph nodes under both steady-state and inflammatory conditions with unknown specific functions. No human equivalent of mouse double-negative dermal DCs has been identified as yet (Table 2).

*2.2. Blood-resident DCs*

Two main DC precursor subtypes are identified in human blood: DCs and pDCs. They are relatively immature and express only low levels of adhesion and costimulatory molecules. Blood DCs (0.26% in leukocytes) descend from the myeloid lineage and express blood DC antigen (BDCA)-1, CD11c, and Toll-like receptors TLR2, TLR4, TLR5, and TLR3. The cells secrete mainly IL-12 in response to bacterial components. pDCs (0.2% in leukocytes) express BDCA-2 (or CD303) and CD123, and are specialized in antiviral innate immune responses by producing copious amounts of type I interferons (IFN-γ) upon exposure of intracellular TLR9 and TLR7 to DNA and RNA viruses. In addition, a small third population (0.02% of leukocytes) of blood DCs are distinguishable based on the expression of CD11c and BDCA-3 (or CD141), but not BDCA-1, CD123 and BDCA-2 [14]. Mouse blood DCs are less well characterized with a majority of circulating MHC class II<sup>+</sup> CD11c<sup>+</sup> as pDCs and low

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