# REVIEW



# Evidence on the neural crest origin of PEComas

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# **Abstract**

The perivascular epithelioid cell (PEC) has been proposed to be the proliferating cell type in a group of tumors known as PEComas. The histogenesis of PEComas is one of the most mysterious aspects of pathology. Hypothesis on its precursor are many, including a cell from blood vessel walls or the myoblast. In the current report, we review many morphologic, clinical, ultrastructural, molecular and genetic aspects that support the hypothesis of an origin of PEComas from the neural crest.

Keywords: perivascular migration, PEComa, "sugar" tumor, perivascular epithelioid cell, angiomyolipoma, neural crest.

# ☐ Introduction

The perivascular epithelioid cell (PEC) has been proposed to be the proliferating cell type in a group of tumors known as PEComas [1]. These so-called PEComas include angiomyolipoma, lymphangiomyoma, lymphangioleiomyomatosis, renal capsuloma, clear cell "sugar" tumor and clear cell myomelanocytic tumor [2, 3].

The histogenesis of PEComas is one of the most mysterious aspects of pathology. It has been proposed that they may originate from a precursor from blood vessel walls (Perivascular Epithelioid Cell) [4], maybe from the pericyte. Others say that they might have a myoblastic origin [5].

The hypothesis that they might originate from undifferentiate cells of the neural crest has also been mentioned in literature [6], although some are contrary to this, due to the lack of expression of S-100 by PEComas [7]. One should remember, nevertheless, how some cases of immunoexpression of S-100 by PEComas have been reported [8-10]. In this context, it is also interesting to remember the capacity of the cephalic neural crest, which gives origin to pericytes and smooth muscle cells of blood vessels of the face and the forebrain [11], many of which do not express S-100 protein. Also, some other well recognized derivatives from neural crest do not express S-100 either, such as Merkell cells [12]. Therefore, such an expression is neither synonymous of "derivative from neural crest" nor a requisite to admit such an origin of a cellular lineage.

In the current report, we review many aspects that support the hypothesis of an origin of PEComas from the neural crest.

# Development of the neural crest and derivatives

Neural crests are bilaterally paired strips of cells

arising at the margins of the neural tube. They have a neuroectodermal origin and later they undergo an epithelial—mesencymal transformation, subsequently exiting from the dorsal neural tube and migrating to different organs. Shortly after the neural tube is closed, the migration of cells from the neural crest begins. This happens in a head-to-tailward (rostrocaudal) sequence. The cells of the neural crest start by "delaminating" from the neuroepithelium, a fact to which some anatomic peculiarities probably contribute: for instance, the basement membrane surrounding the neural tube is discontinuous over its dorsal aspect [13].

Summarizing, there are many aspects that distinguish the migration of cells from the cranial neural crest and the one from the trunk neural crest. Cells from the latter migrate through two main paths: a dorsal pathway underneath the ectoderm, and a ventral pathway through the somites [14]. As a result, many structures that are currently recognized as derivatives from the neural crest will develop (Table 1).

In embryology, several techniques have been used to study if a cell population has a neural crest origin. For instance, the "cell-labeling" technique allows the visualization of the cell migration in developing embryos [15]. Melanocytes would be a recognized example of a neural crest derived cell population [16].

The problem in studying the origin of PEComas with such a technique is, that we are not aware so far, of a normal histological counterpart for these tumors. Assuming that they might develop from scanty residual perivascular cells, trapped in their migration from the neural crest, it would require a lot of luck to locate them. Moreover, it would be almost an act of faith to believe that those cells actually are a counterpart of PEComas.

However, there is indirect evidence, which supports the origin of PEComas from the neural crest.

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#### Table 1 - Cells derived from neural crest

#### Neuronal cells:

- Ganglion cells of the autonomic nervous system;
- Spinal ganglia;
- Sensory ganglia of cranial nerves (V, VII, IX and X).

#### Cells of the peripheral nervous system:

- · Glial cells of the ganglia;
- Schwann cells.

#### Supportive cells of the brain:

Meninges (anterior brain)

#### Some cutaneous cells:

- Melanocytes;
- Merkel cells

#### Endocrine and paracrine system:

- Catecholaminergic adrenal cells;
- · Calcitonin-producing cells;
- Type I cells of the carotidian body

#### Mesectodermal derivatives:

- Facial dermis and cranial cartilage, bone, adipocytes and meninges;
- Wall of large arteries (from the aortic arches);
- Connective tissue of thymus and parathyroid gland;
- Adipocytes within parasympathetic ganglia of the gut;
- Peripheral nerve fibroblasts.

# Proof related to the immunohistochemical and morphological peculiarities of PEComas

PEComas typically co-express melanocytic markers (such as HMB-45 [11, 12], melan-A [MART-1] [8] and microphthalmia transcription factor [MITF] [9]) and muscular markers (such as actin and less frequently desmin [8–10]) (Figure 1).

The expression of melanocytic markers is not aberrant but due to the presence of pre-melanosomes in the cells of PEComas [17-22]. There is evidence that cells that are derived from the neural crest can express many of the markers mentioned above, under certain environmental influence. It is well-known how the chief cell from pheochromocytomas (a tumor developed from a neural crest derivative) expresses HMB-45 [23]. It has been demonstrated that under the influence of TGF-B. multipotential cells from the neural crest can differentiate to smooth muscle cells [24]. Also, studies of large series of neural crest-derived tumors have demonstrated that these can show expression of  $\alpha$ -smooth muscle actin on rare occasions [25]. Also, as a curiosity, some have published cases of malignant melanoma that showed differentiation to smooth muscle [26]. Neural crest cells can differentiate into smooth muscle cells in vitro in the presence of transforming growth factor (TGF)-β1 [27], folic acid [28], or when a specific type of culture medium is used [29].

In literature, there is a concept that the precursor of PEComas (whichever it is) has the capacity to modulate their morphology and phenotype, varying among a spindle cell (actin+, HMB-45+/-), an epithelioid cell (HMB-45+, actin+/-), or even to acquire a vacuolated morphology with an adipocyte-like appearance [6].

Admitting PEComas as derived from the neural crest could explain this morphologic variability and also its components of smooth muscle, melanocytic-like cells and adipocytes. For instance, it has been demonstrated how the multipotent, self-renewing neural crest-derived cells change intrinsic properties with time and location [30].

Some works on neural crest precursors isolated from the skin, have demonstrated that their differentiation *in vitro* resulted in the *de novo* generation of separate subpopulations of cells expressing neuronal, glial, smooth muscle and adipocyte markers [31].

Moreover, neural crest cells when cultured under the appropriate influences can differentiate into adipocytes [32]. A subset of adipocytes derives from the neural crest [32], such as the ones within the parasympathetic ganglia of the gut [33].

There is another interesting morphologic peculiarity of PEComas: their close relation to blood vessel walls. It is worth mentioning that a group has recently focused their attention on the phenomenon of the perivascular location of melanocytic nevi [34]. They proposed that such a phenomenon could be crucial in the histogenesis of melanocytic nevi [34]. If it was admitted that such a mechanism of migration was also used by the PEComa precursor when migrating from the neural crest, this would explain the intimate association of PEComas with the blood vessel wall [35]. In this way, in their migratory journey from the neural crest, the PEComa precursor might use (at least partly) the perivascular route. They would reach their final destination, acquiring their final phenotype... or not: some cells would remain trapped in the perivascular location in several organs. They would stay undifferentiated with capacity to express several markers under the environmental influence, such as smooth muscle, melanocytic or lipomatous markers. It was demonstrated years ago that similar undifferentiated precursors, derived from the neural crest, were present in many organs even in the adult life [36]. For instance, it has been discovered how the skin of adults has neural crest derived precursors that arise in skin during embryogenesis and persist into adulthood [37]. It has also been shown how these precursors have the ability to expand and differentiate into bone, cartilage, fat and muscle [38]. Curiously, there are cases of PEComas in literature, with an intense bone component, although it has been interpreted many times as metaplastic [39].

The fact that neural crest precursors could be the origin of certain tumors has already been suggested for some tumors and malformations of the skin and soft tissues, such as neurofibromas, melanocytic hamartomas, o some melanocytic nevi [40, 41].

Apart from melanocytic or muscular markers, there are other markers that link PEComas with the neural crest. Stem cell precursors have been recognized that can give rise to smooth muscle and melanocytes. These precursors express the neural stem cell markers NG2 and L1 [42–46]. Both markers have been found in angiomyolipomas [47].

From a genetic point of view, it should be mentioned that angiomyolipoma expresses a class of neural crest specifier genes, such as Snaill, Twistl, Sox9 and FoxD3 [48].

On the other hand, some effector genes regulate neural crest differentiation by activating some cell-type-specific effectors including, for instance, c-Kit [49]. c-Kit is expressed in many cases by angiomyolipomas [50–52].

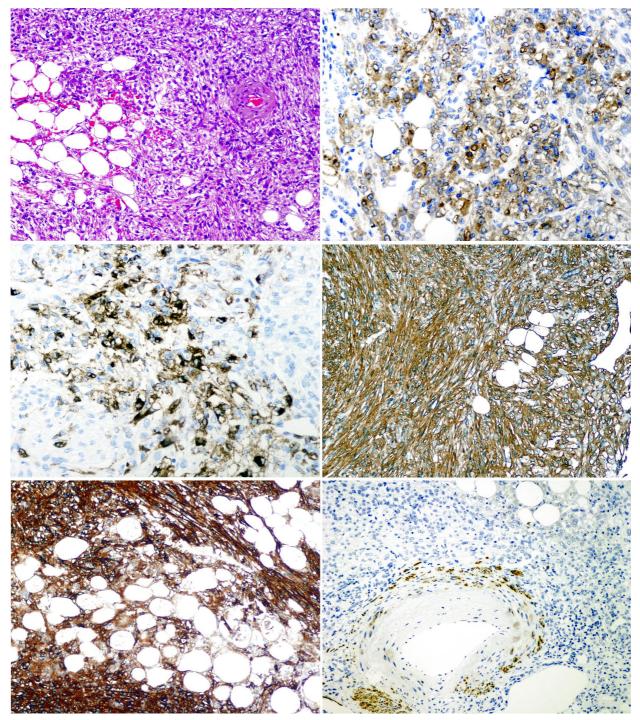


Figure 1 – Renal angiomyolipoma showing several component of epithelioid cells and adipocytes. The tumor surrounds thick-walled blood vessels (top-left). The immunohistochemistry shows immunoexpression by the tumor of melanocytic markers such as Melan-A (top-right) or HMB-45 (second row, left). It also expresses muscular marker, such as smooth muscle actin (second row, right) or actin (bottom, left). Desmin is less commonly expressed, although blood vessel walls can be use as a reliable internal control (bottom right).

# ☐ Ultrastructural evidence

Some ultrastructural features are considered as indicative of a neural crest origin, such as neurite-like cytoplasmic processes, fine filaments and microtubules that are indistinguishable from those seen in normal neurites, synaptic-like structures, and neurosecretory-like vesicles [53].

Some have demonstrated how, in tissue cultures of angiomyolipoma, the cells exhibit long neurite-like processes, which are similar to neuritis [54]. Also,

microtubular have been found in "sugar" tumor [54]. Moreover, neurosecretory-type granules have been found in cases of "sugar" cell tumor of the lung [56–59].

# ☐ Proof related to clinical peculiarities of PEComas

Another peculiarity of PEComas is their ubiquity. They have been described in many visceral and somatic locations, such as the kidney [60], urinary bladder [61], prostate [62], uterus [63], ovary [64], vagina [65], vulva

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[55], lung [66], pancreas [67], liver [68], breast [69], soft tissues [70], skull base [71], gastrointestinal tract [72], oral mucosa [73], nasal cavity [74], bone, orbit, retroperitoneum [75], inter-atrial cardiac septum [76] and, of course, the skin [3].

An origin from the neural crest would explain such ubiquity since derivatives from neural crest are found in somatic as well as in visceral locations [14].

On the other hand, the association between angiomyolipomas and tuberous sclerosis is a well-known fact [6, 76–78]. Tuberous sclerosis is related to alterations in the *TSC1* and *TSC2* genes [79–83]. *TSC1* and *TSC2* encode hamartin (also known as tuberous sclerosis protein 1) and tuberin (or tuberous sclerosis protein 2), respectively. These are two cytosolic proteins that function together as a heterodimer participating in signaling pathways that control cell growth and proliferation [84]. Chromosomal alterations in *TSC1* and *TSC2* have even been found in hamartomas of patients with tuberous sclerosis [85].

Loss of heterozigosity (LOH) of chromosome arm 16p (containing the *TSC2* locus) was demonstrated in both the inherited and sporadic forms of angiomyolipomas [85]. LOH of the *TSC1* gene was also found in angiomyolipomas [85]. Also, other chromosomic alterations that involve *TSC1* and *TSC2* have been described in PEComas [86, 87]. Similar alterations of *TSC2* have been described in cases of lymphangiomyomatosis, which is included in the group of PEComas [88–92]. *TSC1* has also occasionally been involved in focal cortical dysplasia, which is due to alterations in cell migration from the neural crest [93, 94].

## Future lines of investigation

One of the difficulties in identifying the precursor cell of PEComas could lie in a simple fact: even when cells from the neural crests have reached their final destinations, a number of them remain undifferentiated, pluripotent and even endowed with the stem cell capacity of self-renewal [95]. The techniques of cell labeling, as commented above, are difficult to use with PEComas, since the normal histological counterpart is unknown. Therefore, more evidence supporting a neural crest origin for PEComas could be found from molecular and genetic proof. It is predictable, for instance, that PEComas express neural crest effector genes, such as PAX3 (the same way melanocytes do [96]), D1x5 or Msx1/2. It is also predictable that neural crest specifiers genes are activated in PEComas. Examples are Sox10, Sox9, AP-2 and c-Myc [49]. Sox gene, for instance, has been implicated in melanocyte development and melanoma [97].

While c-myc expression was not relevant in some angiomyolipomas tested [98], in some others, a hormonal treatment has stimulated the expression of c-myc by tumoral cells in cases of lymphangioleiomyomatosis [99]. Also, some of the transcription factors mentioned above, are involved in the regulation of the MITF promoter, such as Pax3, CREB, Sox10, and Lef1 [100]. It is a well-known fact how MITF is expressed by

PEComas [101]. MITF also plays neural crest effector functions and is involved in the delamination and the regulation of adhesive properties [49], with a relevant role in the differentiation and/or the functional features of several cell types including osteoclasts, melanocytes, mast cells, and natural killer cells [102–104]. Therefore, some other similar neural crest effectors, such as P0, Connexin 32 (Cx32) o Trp, could be searched (and probably found) in PEComas.

The evidences supporting a hypothesis on the neural crest origin of PEComas are many. Such a hypothesis would clearly explain many of the peculiarities of these tumors

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