



# Enteroendocrine cells in the Echinodermata

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Received: 9 January 2019 / Accepted: 24 May 2019 / Published online: 20 June 2019  
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## Abstract

Enteroendocrine cells are endocrine-like cells found in the luminal epithelia of the digestive tract. These cells have been described in most animal phyla. In echinoderms, the cells have been described mainly in organisms of the class Asterozoa (sea stars) and Holothurozoa (sea cucumbers). Here, we describe what is known about the enteroendocrine cells of the Echinodermata, including the cell types, their distribution in the digestive tract, their neuropeptide content and their regeneration and compare them to what has been found in other animal species, mainly in vertebrates. We also discuss the newly described view of enteroendocrine cells as chemical sensors of the intestinal lumen and provide some histological evidence that similar functions might be found within the echinoderms. Finally, we describe the temporal regeneration of the enteroendocrine cells in the holothurian intestine.

**Keywords** Echinoderm · Digestive tract · Endocrine · Neuropeptides · Regeneration

## Introduction

Enteroendocrine cells (EECs) are endocrine cells found within the digestive tract of animal species. These cells have been described in most, if not all, metazoans, where a digestive tract is present. They comprise what has been called a “diffuse endocrine system” that, in mammals, is thought to be the largest endocrine system of the body. The cells are thought to be involved in multiple functions that include the detection and response to environmental, microbial, nutrient and other factors. Their key localization in the luminal mucosa, at the interface between the intestinal lumen and the internal body system, reinforces their importance in transducing information from the environment that lies outside the body epithelial boundaries to the tissues of the organism.

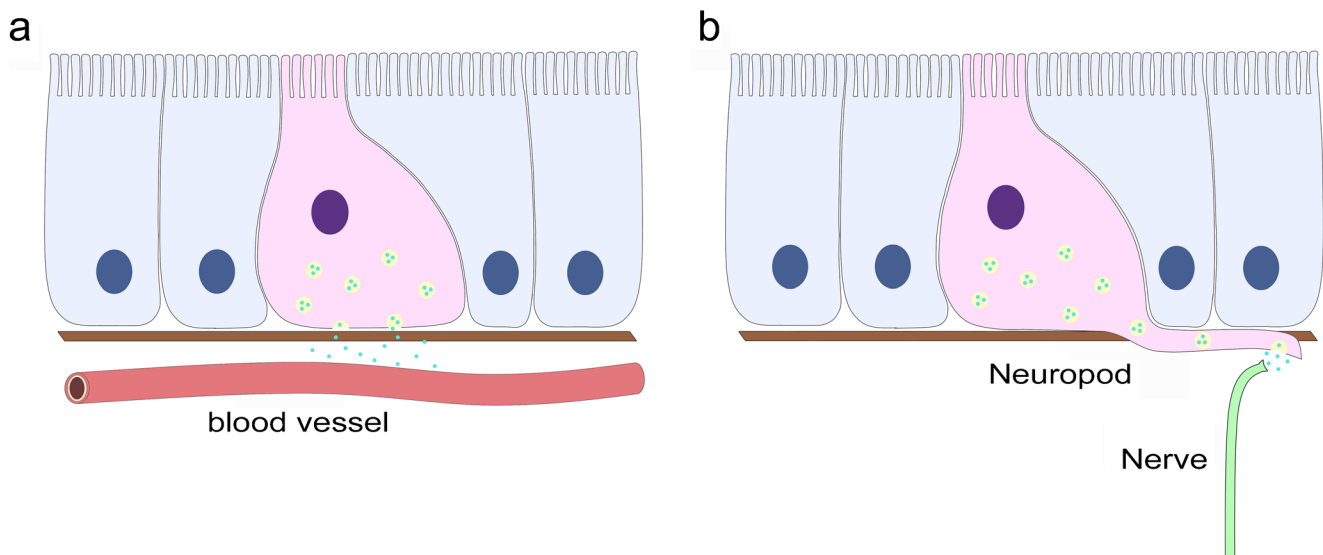
EECs have been best studied in vertebrate animals (Fig. 1a). They have been described as small granulated cells that are scattered among the cells of the luminal epithelium of the gastrointestinal tract (Fawcett 1994; Furness 2006; Latorre et al. 2016). They have an oval or pyramidal structure, lying on the basal lamina and many of them extending to the

luminal surface where they have microvilli that come into contact with the luminal content. Their cell bodies contain numerous secretory granules that have been found to contain multiple types of amine and peptide hormones. These granules are usually localized toward the basal end of the cells and, when released, their contents are thought to enter the adjacent circulatory vessels and carried to their target cells in different body organs.

There have been various attempts to classify EECs into subtypes. For example, a subtype of EEC that contained biological amines usually found in chromaffin cells (catecholamines and/or serotonin) was classified as enterochromaffin cells. Although they were initially considered to be different from peptide-containing EEC, further experimental work has shown that a large number of enterochromaffin cells also expressed peptide hormones (Diwakarla et al. 2017; Fothergill and Furness 2018). Other classification attempts relied on the type of neuropeptide hormone that the cells expressed but recent research has shown that multiple peptides could be expressed by single EECs. This was made evident by the application of single-cell sequencing to the EECs, showing that there are at least eight clusters of mature EECs that can be identified in the mammalian small intestine and that key hormones were expressed by cells in different clusters (Haber et al. 2017). These new findings have made the EEC classification systems defined by a single hormone or by a letter code obsolete.

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**Fig. 1** Models of enteroendocrine cells (EECs). EECs are portrayed as polarized cells in the mucosal membrane of the digestive tract that have their apical end projecting into the luminal space while basally they are attached to the basal lamina. **a** The “classical” view of these cells is that

they are endocrine-like cells that release their contents into the adjacent vascular vessels. **b** The “emergent” model is that they have basal extensions or “neuropods” that contact directly the nervous system acting as neurosensors

In the last few years, investigators’ perception of EECs has taken a new twist. Traditionally, EECs have been perceived as being “classical endocrine” cells releasing their peptide or neuroactive contents in a paracrine or endocrine mode. However, recent experiments in mammals have shown that, at least some EECs have axon-like basal processes, named neuropods, that form synapse-like connections with nerves (Bohórquez et al. 2011, 2015; Liddle 2019) (Fig. 1b). This has led to a new view of, at least some EECs, being epithelial sensory transducers equivalent to the olfactory cells in the nose or to taste cells in taste buds. In this way, these EECs contact directly the fiber of a sensory neuron that takes the information directly to the brainstem (Bohórquez et al. 2015). Hence, the use of the terminology “gut epithelial sensors” to highlight the sensory/electrical transducer nature of the EECs (Kaelberer and Bohórquez 2018).

### Peptide-like immunoreactivity in echinoderm EECs

As for other animal species, EECs have been described within the digestive tract of echinoderms. (Table 1 lists the various EECs that have been described, the method of detection, the specie, and the reference). The vast majority of EEC descriptions have been done using immunohistological techniques, mainly with antibodies against specific neuropeptides. This implies that most of the earlier reports identify what has been named as “peptide-like immunoreactivity” in view that the antibody could be recognizing a similar but not identical antigen and that there is no certainty that the peptide being recognized is the “real” peptide.

To our knowledge, the first EEC descriptions in the phylum Echinodermata were done by Martínez and colleagues (Martínez et al. 1989). The cells were found in the pyloric caeca of a sea star (*Marthasterias glacialis*) using electron microscopy and immunohistochemistry. EECs were described within the luminal epithelium of this organ using histological stains (Grimelius silver impregnation). They were shown to have the typical traits of EECs (Figs. 2, 3, 4 and 5 show typical echinoderm EECs): a thin oval morphology that extends from the basal lamina to the lumen, an elongated nucleus in a central position and large numbers of secretory granules that are usually electron-dense. A long apical cytoplasmic process extended to the lumen and the basal region was in direct contact with nerve fibers. Some of these cells were found to express neuropeptide-like immunoreactivity to somatostatin, glucagon and pancreatic polypeptide.

The pyloric caeca is an organ specific to the digestive tract of asteroids but in later studies, Martínez and other investigators showed the presence of EECs in other regions of the digestive system (Martínez et al. 1993, 1994, 1996). In addition, they expanded the number of neuropeptides that could be found within the neuroendocrine cells and fine-tuned the description of the asteroid EEC system. Thus, EECs were found throughout the digestive system of the sea star, including the cardiac and pyloric stomach, the intestine and the pyloric and rectal caeca (Martínez et al. 1993, 1996). Immunoreactivity to alpha-MSH, PYY, adrenomedullin and FMRFamide was also detected in EECs (Martínez et al. 1993, 1996). However, the definitive result was the finding of S1 immunoreactivity in EECs, since this was the first neuropeptide isolated from echinoderms (in what will be named the SALMFamide peptide family) and as such the first neuropeptide where the observed

**Table 1** The various EECs that have been described, the method of detection, the species and the reference

Neuropeptide	Species	Class	In situ	Immuno	Distribution	Reference
Somatostatin	<i>M. glacialis</i>	Asteroidea		X	PC	Martínez et al. (1989)
Glucagon	<i>M. glacialis</i>	Asteroidea		X	PC	Martínez et al. (1989)
Pancreatic polypeptide	<i>M. glacialis</i>	Asteroidea		X	PC, CS	Martínez et al. (1989)
a-MSH	<i>M. glacialis</i>	Asteroidea		X	PC, PS RC	Martínez et al. (1993)
S1 and FMRFamide	<i>M. glacialis</i>	Asteroidea		X	CS, PS, PC, RC	Martínez et al. (1993)
PYY	<i>M. glacialis</i>	Asteroidea		X	PC, RC, CS	Martínez et al. (1993)
NO synthase	<i>M. glacialis</i>	Asteroidea		X	CS, PS, PC	Martínez et al. (1994)
Adrenomedullin	<i>M. glacialis</i>	Asteroidea		X	CS, PS, PC	Martínez et al. (1996)
GFSKLYFa	<i>H. glaberrima</i>	Holothuroidea		X	E, SI, LI, C	Díaz-Miranda et al. (1995)
Calbindin	<i>H. glaberrima</i>	Holothuroidea		X	I	Díaz-Balzac et al. (2016)
NGIWFamide	<i>A. japonicus</i>	Holothuroidea		X	I	Inoue et al. (1999)
SALMFamide 1 (S1) SALMFamide 2 (S2)	<i>A. rubens</i>	Asteroidea		X	E, CS, PS, PC, RC	Moore and Thorndyke (1993) Newman et al. (1995)
ArPPLNP1 (pedal peptide/orcokinin)	<i>A. rubens</i>	Asteroidea	X	X	CS, PS, PD	Lin et al. (2017)
ArPPLNP2 (pedal peptide/orcokinin)	<i>A. rubens</i>	Asteroidea		X	CS, PS	Lin et al. (2018)
GnRH	<i>A. rubens</i>	Asteroidea	X	X	CS, PS	Tian et al. (2017)
Corazonin	<i>A. rubens</i>	Asteroidea	X		CS, PS, PD	Tian et al. (2017)
Luqin/ACEP-1	<i>A. rubens</i>	Asteroidea	X		CS, PS	Yañez-Guerra et al. (2018)
NGFFYamide	<i>A. rubens</i>	Asteroidea	X		CS, PS, PD	Tinoco et al. (2018)
ArCT	<i>A. rubens</i>	Asteroidea	X	X	E, CS, PC, PD, PS, R	Cai et al. (2018)
Other markers						
Pax6	<i>H. glaberrima</i>	Holothuroidea		X	I	Díaz-Balzac et al. (2014, 2016)
Nurr1	<i>H. glaberrima</i>	Holothuroidea		X	I	Díaz-Balzac et al. (2014, 2016)
PAM (C-terminal amidating enzymes)	<i>M. glacialis</i>	Asteroidea		X	CS, PS, RC, I	Martínez et al. (1993)

C cloaca, CS cardiac stomach, E esophagus, I intestine, LI large intestine, PC pyloric caeca, PD pyloric ducts, PS pyloric stomach, RC rectal caeca, SI small intestine

immunoreactivity in the EEC could be directly correlated to the peptide sequence (Elphick et al. 1991). Moreover, these findings were confirmed by immunocytochemical studies in the digestive system of a second starfish species, *Asterias rubens* (Moore and Thorndyke 1993; Newman et al. 1995).

Neuropeptide-containing EECs have also been described within the luminal epithelia of other echinoderm classes. In the sea cucumber *Holothuria glaberrima*, our group has shown that cells immunoreactive to GFSKLYFamide, a holothurian neuropeptide of the SALMFamide family, are found in the small and large intestine (Díaz-Miranda et al. 1995) (Fig. 3). Immunoreactivity to a different peptide, calbindin, has also been documented in the intestine of *H. glaberrima* (Díaz-Balzac et al. 2016). Another neuropeptide that has been isolated and characterized from holothurian tissues NGIWFamide has also been documented in the EECs of the sea cucumber *Apostichopus japonicus* (Inoue et al. 1999). Thus, in holothurians, similar to sea stars, EECs have been described that expressed holothurian-isolated neuropeptides and these cells can be found along most of the digestive tract

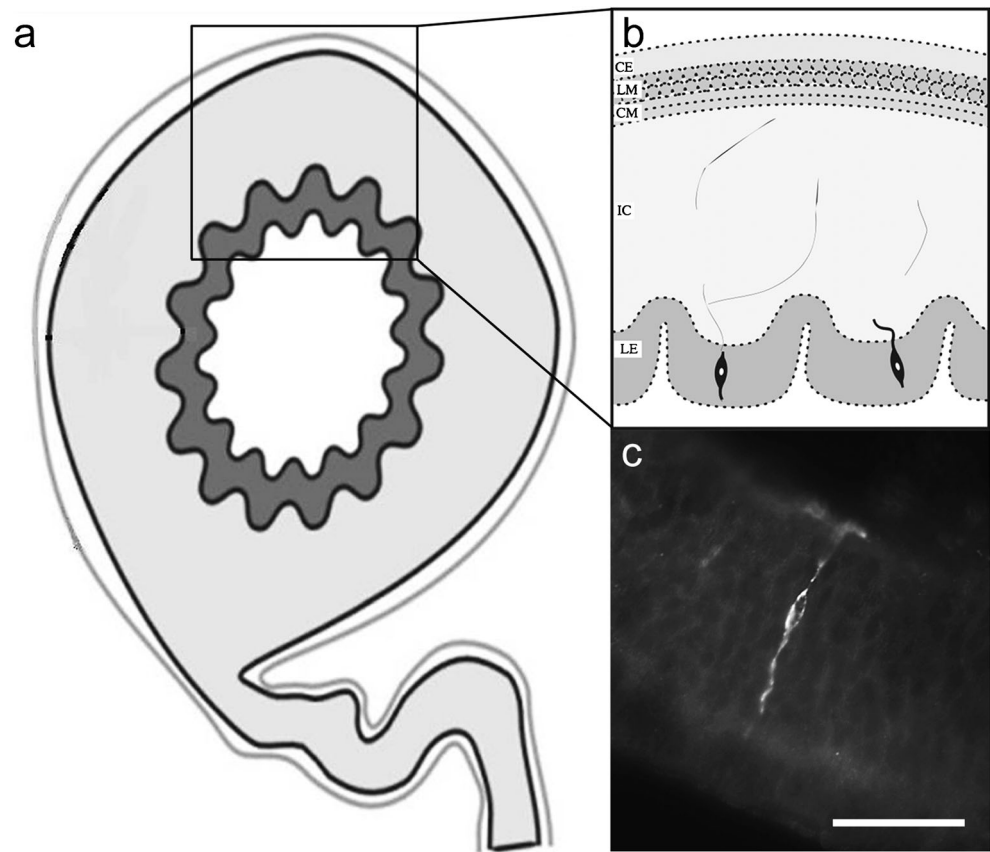
including esophagus, stomach, small and large intestines and rectum (García-Arrarás et al. 2001).

Echinoderm EECs have been immune-labeled with other non-peptide markers. In this respect, they have been shown to express immunoreactivity to C-terminal amidating enzymes (PAM) and the nitric oxide synthetase enzyme in the digestive system of asteroids (Martínez et al. 1993, 1994) and to Pax6 and Nurr1 transcription factors in the digestive system of holothurians (Díaz-Balzac et al. 2014).

## The genomic era of neuropeptide characterization

In the not so distant past, EECs were recognized by the expression of peptide-like immunoreactivity; however, in most of these cases, the specific nature of the immunoreactivity (the peptide sequence) remained uncertain. In recent years, the situation has reversed. The sequencing of the sea urchin genome and the study of mRNA transcriptomes caused a major

**Fig. 2** Anatomy of echinoderm intestinal system showing the localization of enteroendocrine cells. **a** Diagram of transverse section of intestinal system. **b** Higher magnification showing the localization of two enteroendocrine cells within the luminal epithelium. **c** Typical morphology of an enteroendocrine cell showing the elongated body extending from the lumen to the basal lamina. ce coelomic epithelium, cm circular muscle, ic internal connective tissue, le luminal epithelium, lm longitudinal muscle. Bar = 25  $\mu$ m

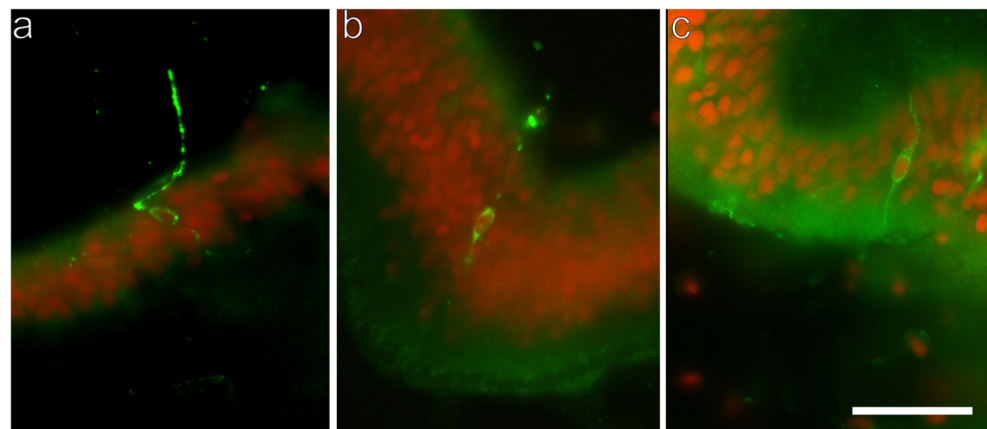


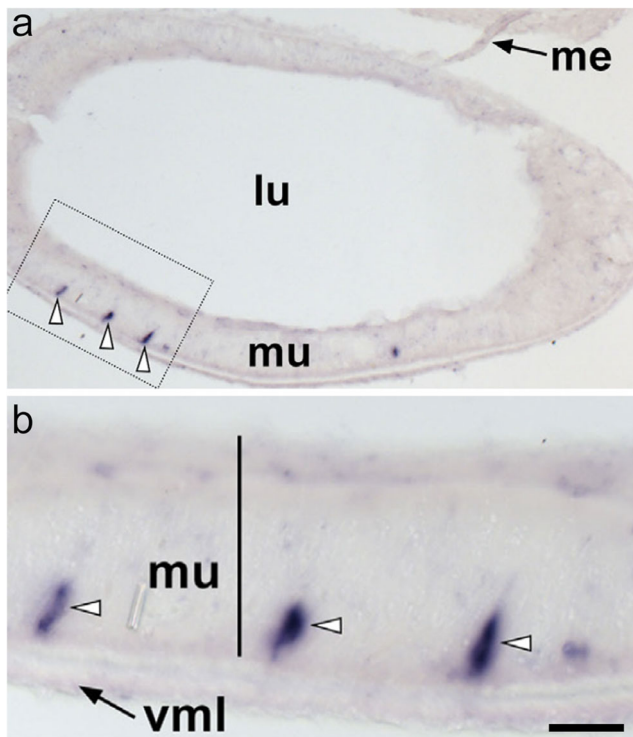
shift in the study of the echinoderm neuroendocrine system. While previously most of the studies depended on the purification and characterization of neuropeptides using protein isolation technologies or on the identification of peptide-like immunoreactivity using anti-sera, the genomic and transcriptomic studies suddenly provided the sequence to multiple putative neuropeptides that now needed to be characterized in relation to their cellular/tissue localization (Burke et al. 2006; Rowe and Elphick 2012). Thus, in situ hybridization was used to determine the expression of putative neuroactive peptides in EECs (see Fig. 4 as an example). In addition, a few

studies managed to bridge the gap in knowledge and used immunohistochemistry and mRNA in situ hybridization to reveal the expression by EECs of putative peptides encoded by the genes characterized from the genomic data. These studies mainly led by Elphick and colleagues are summarized in Table 1.

Nonetheless, while genomic or transcriptomic studies have been done in members of most echinoderm classes (Rowe and Elphick 2012; Rowe et al. 2014; Semmens et al. 2016; Zandawala et al. 2017; Kim et al. 2018; Suwansa-ard et al. 2018) and new putative neuropeptides have been described,

**Fig. 3** Immunohistochemistry of enteroendocrine cells of *Holothuria glaberrima*. Enteroendocrine cells expressing **a** GFSKLFamide, **b** Pax 6 and **c** calbindin in the intestinal luminal epithelium. Note the axon-like extension of the GFSKLYFamide expressing cell that extends into the connective tissue. Bar = 25  $\mu$ m



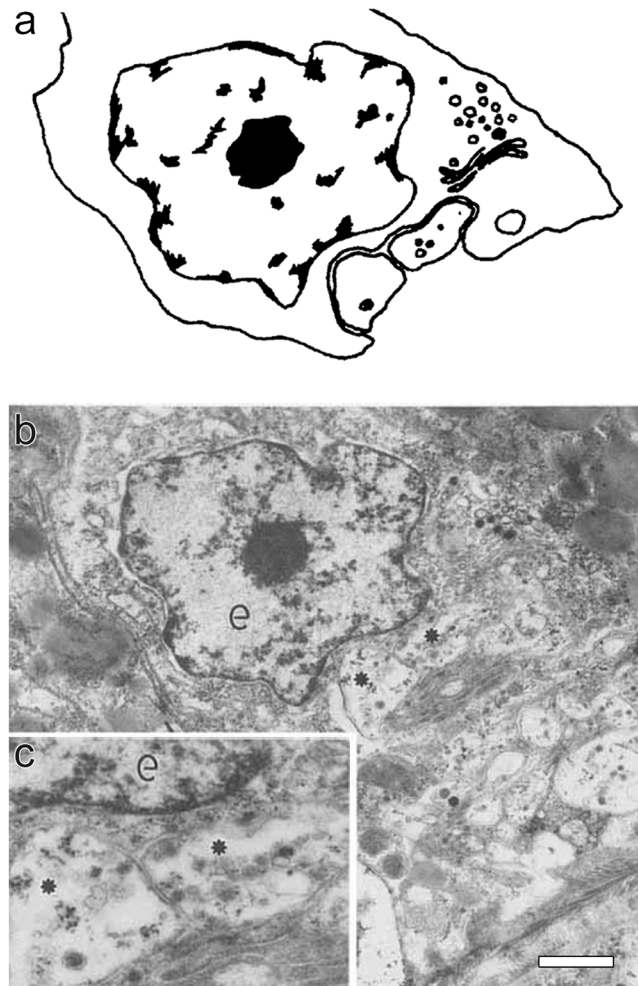


**Fig. 4** In situ hybridization for corazonin neuropeptide. **a, b** Transverse section of the pyloric duct of the sea star *Asterias rubens* showing the mRNA in some enteroendocrine cells in the mucosal layer. Cells expressing the mRNA for the peptide are marked with arrowheads in low (**a**) and high (**b**) magnification. lu lumen, me mesentery, mu mucosal layer, vml visceral muscle layer. From Tian et al. (2017). Bar = **a** 90  $\mu\text{m}$  and **b** 20  $\mu\text{m}$

the characterization of EECs expressing these peptides has been limited to members of the Asterozoa and the Holothurozoa. To the best of our knowledge, no neuropeptide-expressing cell has been described in sea urchins, ophiuroids, or crinoids.

### Echinoderm EECs share many characteristics of vertebrate EECs

It is important to highlight that in addition to their elongated morphology that extends from the basal lamina to the lumen and their vesicular content, echinoderm EECs share many other similarities with vertebrate EECs. First, similar to vertebrates the peptides expressed within the EECs are the same peptides expressed by neurons of the central and peripheral nervous system. For example, in vertebrates, EECs might express PYY or CCK, while in echinoderms, EECs express the same peptides that are expressed in the echinoderm neurons, such as SALMFamides or corazonin-type peptides. This makes the functional characterization of the EECs very difficult because sometimes neuronal fibers containing the neuropeptide can be found within the digestive tract and even within the adjacent subepithelial plexus that underlies the luminal



**Fig. 5** Electron microscopy of enteroendocrine cells in the sea star *Marthasterias glacialis*. **a** Diagram of accompanying electron micrograph. **b** Enteroendocrine cell (e) contacting nerve terminals (asterisks). **c** Higher magnification of nerve processes. From Martínez et al. (1989). Bar = 1.5  $\mu\text{m}$

epithelium. Thus, an effect elicited by extrinsic neuropeptide addition cannot be clearly ascribed to the nervous system or to the EECs.

Second, there are different populations of EECs. In vertebrates, as explained earlier, the neuropeptide content has been used to identify different cell populations. In echinoderms, of the many peptide markers that have been identified, some show specific distributions that suggest different subpopulations. In our laboratory, we put this to the test by labeling cells with anti-GFSKLYFamide and with anti-calbindin. These markers identify two different subpopulations of EECs. They show somewhat different morphologies, where those expressing GFSKLYamide usually have an extension that enters the connective tissue. In addition, their localization within the epithelial lumen differs: the nuclei of those expressing GFSKLYamide are located at about 12  $\mu\text{m}$  from the basal lamina, similar to those of most other luminal cells, while the nuclei of those expressing calbindin are located at about

30  $\mu\text{m}$ , much closer to the cells apical end (unpublished observations) (Fig. 3). Thus, in sea cucumbers, at least two populations of EECs are present.

Third, the number of EECs is relatively small in comparison with the number of other cells in the luminal epithelium. In experiments done in our laboratory, the number of EECs labeled with either GFSKLYFamide or calbindin in a particular region of the digestive tract of a holothurian lies between 4 and 7% (unpublished observations).

Fourth, they are found along the entire length of the digestive tract. Although their numbers might differ, EECs can be found along the entire digestive tract and cells in different locations might express the same or different neuropeptides. Even within a region of the digestive tract, the EEC populations might be heterogeneously distributed. For example, cells expressing ArPPLN1-type peptides in the digestive system of *A. rubens* can be found throughout the cardiac stomach but in the pyloric stomach they are mainly found within floor region and are concentrated on the oral side of the pyloric ducts and of the pyloric caeca (Lin et al. 2017).

There are, however, some differences between echinoderm and vertebrate EECs. The main one lies in that while EECs expressing serotonin have been widely described in vertebrates (Bordi et al. 2000; Gershon 2013), no echinoderm serotonergic EEC has been described.

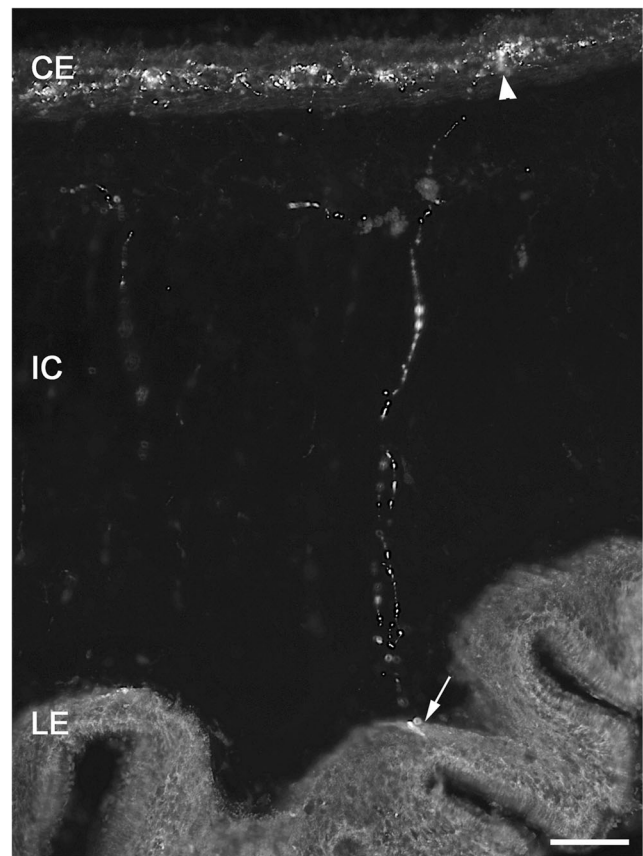
### EECs as “gut epithelial sensors”

It is interesting that some echinoderm EECs perfectly fit the description as epithelial sensory components of the nervous system (Bohórquez et al. 2015; Liddle 2019). In the initial description of echinoderm EECs, there is already the description of how EECs usually have nerve terminals closely associated with their basal end (Fig. 5). The contact between the nerve fibers and the endocrine cells is very close where both membranes are seen in close association, contrary to what had been described in mammals that the nerves do not cross the basal lamina. It took almost three decades to finally show that nerve fibers in mammals do cross the luminal epithelia basal lamina and synapse with the EECs.

One of the findings in holothurians is that some of the neuroendocrine-like cells extend long neurite-like fibers that can be followed into the submucosa or connective tissue and in some cases even to the muscle layer in the mesothelium (Fig. 6) (García-Arrarás et al. 2001).

### Neuroendocrine cell regeneration

Echinoderms are well known for their regenerative abilities. In fact, in recent years, regeneration of the sea cucumber digestive system has been well studied by our group. We described that

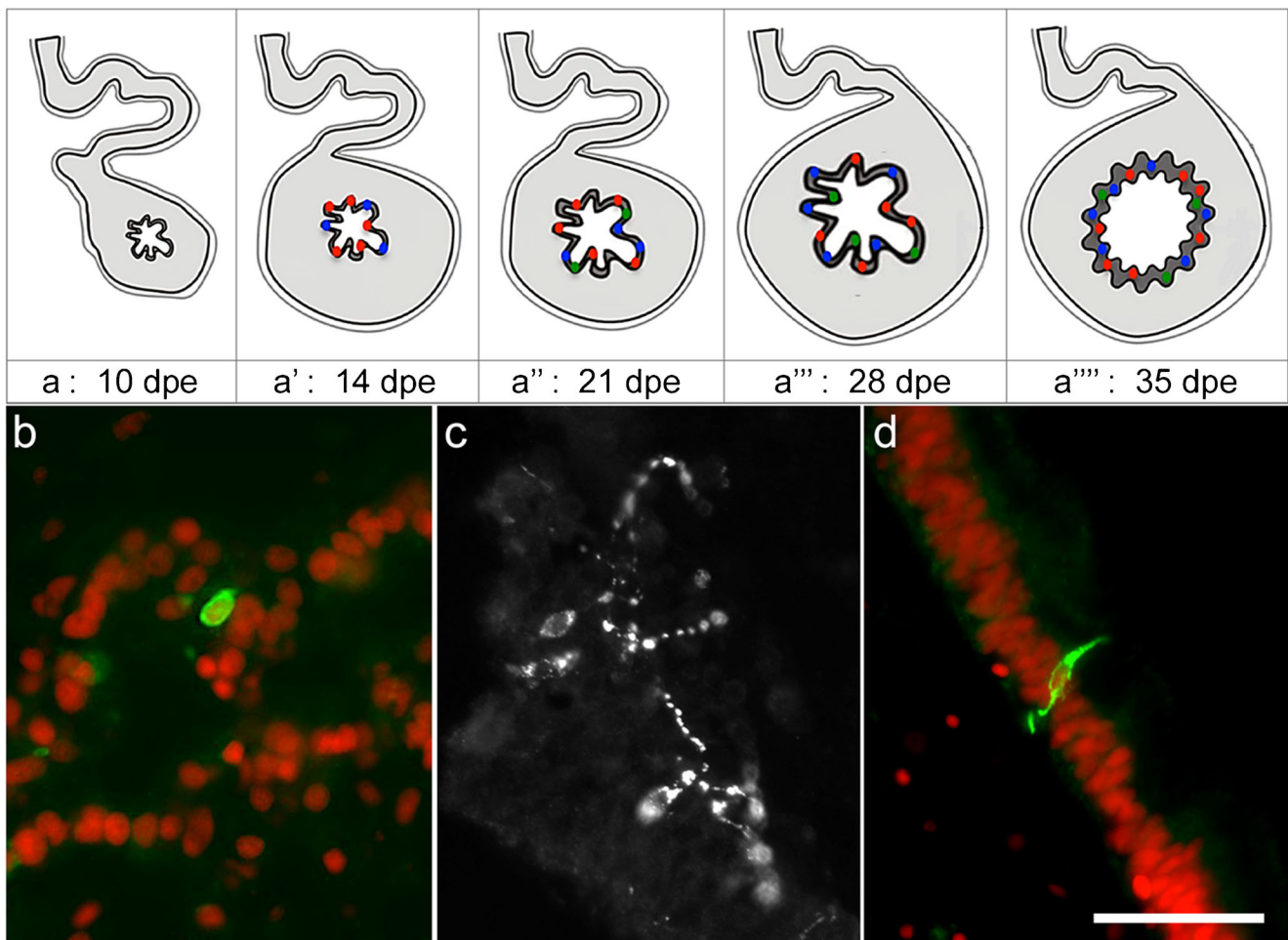


**Fig. 6** GFSKLYFamide expressing cell in the luminal epithelium of the sea cucumber *Holothuria glaberrima*. The cell body (arrow) is labeled in the cytoplasm and a long axon-like process extends from it that runs through the connective tissue and enters the muscle layer (arrowhead). CE coelomic epithelium, IC internal connective tissue, LE luminal epithelium. Bar = 50  $\mu\text{m}$

EECs are found within the regenerating intestine and the timing of their reappearance (García-Arrarás et al. 1999; Tossas et al. 2014). Our data show that not only the intestinal EECs are regenerated but that they appear very early in the formation of the luminal epithelia (Fig. 7). Soon after the luminal epithelia forms, the first EECs can be observed (Tossas et al. 2014). However, the results suggest that not all neuroendocrine cell types appear simultaneously but that different subpopulations appear at different regeneration stages, being those labeled with the anti-calbindin antibody the first to appear, followed by those expressing GFSKLYamide and the latest population to appear are those labeled with the anti-Pax6 marker.

### Concluding remarks

In summary, extensive evidence is available that describes EECs in echinoderms. At present, most of the information is restricted to a small number of species from two classes of the Echinodermata: the Holothuroidea and the Asteroidea. Future experiments will surely extend these findings to other species



**Fig. 7** Enteroendocrine cell regeneration in *H. glaberrima* intestine. **a** Diagram of regenerating intestine at different stages showing the appearance of enteroendocrine cells expressing calbindin (red), GFSKLYFamide (blue) and Pax6 (green) immunoreactivity.

Immunoreactive cells are shown at **b** 10 days of regeneration for calbindin, **c** 14 days for GFSKLYFamide and **d** 21 days for Pax6. (Modified from Tossas et al. 2014). Bar = 25  $\mu$ m

and other classes. This will provide an extended view of comparative findings that will help to discern the commonalities of EECs in echinoderms. Similarly, future studies should provide information on the number and different types of EECs found in the digestive tract and a cohesive view of how these cell types are related to those of other deuterostomes.

Future studies should also address the origin of the EECs. While it is well established that most of the echinoderm nervous system originates from the embryonic ectoderm (Mashanov et al. 2007; Hinman and Burke 2018), the origin of the EECs is assumed to be the same as in its vertebrate relatives. Thus, EECs are thought to be of endodermal origin and it has been assumed that they are formed during intestinal homeostasis and regeneration in the same way as they are in mammals, that is, from a multipotent stem cell in the luminal epithelium. In view of the phylogenetic distance of echinoderms and mammals, it might be interesting to determine if this assumption is correct.

Arguably, one of the least explored characteristics of the EECs is their function. Although the neuroactive molecules

that are found within echinoderm EECs have been well described and continue to be expanded, what are now necessary are studies on the functional aspects of these cells and their contents. The effects of neuropeptides on gut motility have been reported in sea cucumbers (CCK, GFSKLYFamide, NGIWFamide) and starfish (S1, S2, NGFFYamide, ArGnRH, ArCRZ, ArPPLN1-type peptide and ArPPLN2-type peptide). However, as mentioned above, the presence of the same neuropeptide content in both EECs and nerve fibers increases the difficulty in studying EEC function, making it impossible to ascertain a specific role to the EECs versus the enteric nerves. Additional limitations are found in the lack of genetic/molecular tools, such as those used in vertebrates (Sinagoga et al. 2018; Kaelberer et al. 2018), to identify EEC function. Nonetheless, we expect that in the not so distant future, similar tools will be made available to pursue the functional analysis of echinoderm cells. This should include EECs, endocrine cells and neurons associated with the digestive tract.

**Acknowledgments** We would like to thank Ms. Griselle Valentin for editorial help and the preparation of the figures.

**Funding** This project was funded by NIH (Grant R15NS01686). We also acknowledge partial support from NIH R21AG057974 and the University of Puerto Rico.

## Compliance with ethical standards

**Conflict of interest** The authors declare that they have no conflict of interest.

**Ethical approval** Most data have been previously published and are available in the scientific literature. In the case of unpublished data applicable, international, national and/or institutional guidelines for the care and use of animals were followed.

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