



Tuft Cells and Their Role in Intestinal Diseases

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Hendel SK, Kellermann L, Hausmann A, Bindslev N, Jensen KB and Nielsen OH (2022) Tuft Cells and Their Role in Intestinal Diseases. Front. Immunol. 13:822867. doi: 10.3389/fimmu.2022.822867 The interests in intestinal epithelial tuft cells, their basic physiology, involvement in immune responses and relevance for gut diseases, have increased dramatically over the last fifteen years. A key discovery in 2016 of their close connection to helminthic and protozoan infection has further spurred the exploration of these rare chemosensory epithelial cells. Although very sparse in number, tuft cells are now known as important sentinels in the gastrointestinal tract as they monitor intestinal content using succinate as well as sweet and bitter taste receptors. Upon stimulation, tuft cells secrete a broad palette of effector molecules, including interleukin-25, prostaglandin E₂ and D₂, cysteinyl leukotriene C₄, acetylcholine, thymic stromal lymphopoietin, and β -endorphins, some of which with immunomodulatory functions. Tuft cells have proven indispensable in anti-helminthic and anti-protozoan immunity. Most studies on tuft cells are based on murine experiments using double cortin-like kinase 1 (DCLK1) as a marker, while human intestinal tuft cells can be identified by their expression of the cyclooxygenase-1 enzyme. So far, only few studies have examined tuft cells in humans and their relation to gut disease. Here, we present an updated view on intestinal epithelial tuft cells, their physiology, immunological hub function, and their involvement in human disease. We close with a discussion on how tuft cells may have potential therapeutic value in a clinical context.

Keywords: chemosensing, colorectal neoplasia, Crohn's disease, inflammatory bowel disease, inflammation, intestine, tuft cells, ulcerative colitis

Abbreviations: ACh, Acetylcholine; ALOX5, Arachidonate 5-lipoxygenase; ATOH1, Atonal bHLH transcription factor 1; AVIL, Advillin; C. difficile, Clostridioides difficile; CD, Crohn's disease; ChAT, Choline acetyltransferase; CK18, Cytokeratin-18; COX, cyclooxygenase enzyme; CRN, Colorectal neoplasia; DCLK1, Double cortin-like kinase 1; GABA, Gamma aminobutyric acid; GI, Gastrointestinal; IBD, Inflammatory bowel disease; ISC, Intestinal stem cell; IL, Interleukin; ILC2, Type 2 innate lymphoid cell; LTC4, Cysteinyl leukotriene C4; PGD2, Prostaglandin D2; PGE2, Prostaglandin E2; scRNA-seq, Single-cell RNA sequencing; SOX9, SRY-box transcription factor 9; SUCNR1, Succinate receptor 1; TasR, taste receptor; TC, Tuft cell; TSLP, Thymic stromal lymphopoietin; UC, Ulcerative colitis; VAChT, Vesicular acetylcholine transporter.

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HIGHLIGHTS

1) Intestinal TCs form a central hub, involved in immune and regulatory metabolic networks, monitoring luminal intestinal content using chemosensory taste and succinate receptors, thus responding to a broad palette of substances and pathogens.

2) Upon stimulation TCs produce IL-25, ACh, TSLP, β endorphins, and prostaglandins such as PGE₂, PGD₂ and LTC₄ which are potent paracrine and endocrine signaling molecules, making them attractive potential therapeutic targets.

3) Identifying human intestinal TCs may require different markers from those used in murine experiments. These markers are still being investigated and evaluated. Suggested markers include COX-1, p-EGFR, SOX9, ALOX5, AVIL, girdin and ChAT, although none are TC specific.

4) Mouse studies have highlighted the importance of TCs in intestinal function, particularly in anti-helminthic and antiprotozoan immune responses, as well as in obesity models.

5) Certain human diseases such as inflammatory bowel disease, coeliac disease, and duodenal ulcer, are associated with alterations of intestinal TC populations and TC associated cytokines.

INTRODUCTION

The intestinal epithelium serves a crucial role in maintaining gut mucosal homeostasis (1, 2). It forms a specialized physical and chemical barrier between self and non-self and controls activation of the host's largest immune apparatus by interacting with triggers in the gut luminal content. At the same time, the epithelium is central for absorption of water, ions, and nutrients as well as secretion of ion-containing fluids with waste products especially in the colon.

The intestinal epithelial barrier is maintained by proliferating intestinal stem cells (ISCs) located at the bottom of the intestinal crypts. ISCs give rise to an array of differentiated cells scattered within the epithelial lining, including tuft cells (TCs) (3). While functions of most intestinal epithelial cells are well established, far less is known about the lately rediscovered TCs.

Despite their discovery more than sixty five years ago (4), a functional characterization of TCs was delayed into the 1990s due to lack of specific markers. Meanwhile, progress in TCmarker identification over the last 15 years has provided functional insights and identified more detailed hallmarks of TCs. These include their signaling pathways via luminal cues, apical receptors, second messengers and secretory mediators. Only recently a proximate consensus on detection of human TCs in the gut emerged, which now also enables clinical research on these cells. With this overview we aim to present A) what is currently known about the molecular basis of TC fate and differentiation as well as the identification of human intestinal TCs, and B) the available knowledge of involvement of TCs in diseases of the gastrointestinal (GI) tract, including a discussion of TCs in a clinical context and how elucidating TC functions and abundance might inspire novel therapeutic strategies.

TUFT CELLS

All segments of the GI tract, including esophagus, stomach, intestines, and associated organs, cooperate for nutrient uptake, and at the same time, they form the largest immune apparatus in the body and a physical barrier to the outside world containing a huge variety of potentially harmful agents. This special localization requires a tightly balanced immune surveillance achieved by specialized immune cells cooperating with non-immune cells such as epithelial cells, including TCs.

TCs are found in many tissues, including airways (4), gallbladder (5), thymus (6), pancreas (7), urethra (8), and GI tract (9, 10), and have been described under various names such as brush-, solitary chemoreceptor-, microvillous-, fibrillovesicularand caveolated cells. The name "tuft" is the latest addition to designators for this cell type and originates from its unique brush border morphology, Figure 1. TCs have a flask-shaped body with a narrow neck from which a tuft of microvilli extends from the apical membrane into the intestinal lumen (11). The microvilli are longer and more compact than those of neighboring non-TCs and are connected to a rich apical-basal oriented network of microfilaments and microtubules (5, 12). In TCs these microtubules extend deep down towards the nucleus connecting to the perinuclear endoplasmic reticulum. Membrane bound vesicles (glycocalceal bodies) have been identified both at the microvillus base and between microtubules, suggesting a function as a macromolecular-exchange route between the intestinal lumen and the endoplasmic reticulum, Figure 1 (5, 11, 12). The cvtoskeletal make-up further consists of intermediate filaments such as cytokeratin-18 (CK18) and neurofilaments, which are otherwise a distinctive feature of mature neurons (13). Some TCs have cytoplasmic spinules, which are projections from the lateral TC border that enter adjacent cells reaching their nuclei, and might function as a transport route for molecular cargo to and from adjacent cells. These projections appear on TCs ranging from just below the apical junctional complex to the nucleus level and are often associated with a pair of desmosomes at their base, Figure 1 (5, 11). Also, the basal segments form neuropod-like protrusions extending along the basal lamina indicating paracrine signaling (11, 14, 15), although no secretory vesicles are observed within the TC neuropods. This is unlike the basal aspect of TCs close to adjacent neuronal fibers of the enteric nerve system, where vesicles are present presumably containing acetylcholine (ACh) for paracrine signaling (16).

Most of available data on TCs are based on studies in mice. The main part of this review therefore covers findings from mice, unless otherwise specified in the text.

INTESTINAL TCs AND ORIGIN

The intestinal epithelium is particularly dynamic, as vigorously proliferating ISCs constantly renew the epithelium every 3-5 days (17). ISCs positive for leucine rich repeat containing G protein-coupled receptor 5 (Lgr5) are found at the crypt-bottom



in both the small and large intestine and differentiate towards all the different epithelial lineages, including TCs (18, 19). Intestinal TCs are relatively rare as compared with other epithelial cell types. They are usually distributed in a sporadic and solitary manner, accounting for just 0.4-2.3% of the total epithelial cell population in murine intestinal epithelium (9, 20–22), while the only study on TC density of the human sigmoid colon report ~100 TCs per square millimeter tissue (23). Upon leaving the crypts, migrating along the villus axis, the function of TCs might change with increasing differentiation similar to what has recently been described for enteroendocrine cells (24-26). While the average TC has a turnover rate of 1-2 weeks (27), a small subpopulation of cells (~5%) expressing double cortin-like kinase 1 (DCLK1), a TC marker in mice, is remarkably longlived, surviving for up to 18 months. These long-lived TCs may contribute to the epithelial regeneration following damage as they retain stem cell potential. Furthermore, they can give rise to organoids in vitro (small and large intestine), and might serve as

cancer-initiating cells upon injury and deletion of adenomatous polyposis coli (APC) (28). Meanwhile, it remains uncertain whether these long-lived TCs represent a unique cell population, or a common secretory progenitor given that other secretory cells are shown to have similar properties (29).

TC Differentiation

The identification of DCLK1 as a marker for TCs in mice facilitated their further characterization. It was quickly established that differentiation of ISCs to TCs is independent of the transcription factor neurogenin-3 (essential for differentiation of ISCs to intestinal enteroendocrine cells), and thus not associated with the enteroendocrine cell lineages (9). Instead, a common differential marker for all TCs across organs and regions is the taste-cell specific transcription factor, Pou2f3. Pou2f3 is a master regulator of and an absolute requisite for differentiation into TCs, indicating lineage similarity for lingual taste buds and TCs along the entire GI tract (30–33). Despite

sharing this common transcription factor and being morphologically similar across tissues, transcriptional profiling using population-based and single-cell RNA sequencing (scRNA-seq) uncovered TC heterogeneity both across (34) and within tissues (22, 35-37). Haber et al. (22) highlighted two different clusters of TC progenitors, as well as two distinct clusters of mature TCs in the murine small intestine. One of them (denoted Tuft-1) expresses a neural development gene signature, whereas the other (denoted Tuft-2) is enriched with genes associated with immunological functions (22). Despite a major overlap in expression patterns between Tuft-1 and Tuft-2, solely Tuft-2 expresses thymic stromal lymphopoietin (TSLP) (22, 38), an epithelial derived cytokine promoting Th2 immunity. This suggests functional, and potentially spatial, heterogeneity of TCs (22). TC heterogeneity is further illustrated as an additional transcription factor associated with TC differentiation, Atonal bHLH transcription factor 1 (ATOH1), is region specific. ATOH1 is essential for the differentiation into secretory cell lineage (goblet cells, Paneth cells and enteroendocrine cells), as well as in the differentiation into TCs within the large intestine, while small intestinal TCs are ATOH1-independent (30, 39, 40). Krüppel-like factors such as Klf3 and Klf6 were also identified as transcription factors in small-intestinal TCs, while their roles in colonic TC development were not investigated (22). A recent study pinpointed the role of another intracellular signaling regulator in TC differentiation, namely sprouty2, which is mostly expressed in colonic epithelium compared to the remaining GI tract (41). It was demonstrated, both in vitro and in vivo, that acute colonic inflammation reduces sprouty2, leading to TC and goblet cell hyperplasia (41).

Identification of Human Intestinal TCs

Identification of new reliable markers has provided insights into the function of TCs, including their involvement in diseases of the human body. However, the identification and characterization of TCs in the human intestinal tract based on marker expression is still awaiting consensus. Two studies (42, 43) describe a population of human colonic DCLK1⁺ TCs, however, Leppänen et al. (44) reported that human colonic epithelial DCLK⁺ cells have a morphology similar to absorptive enterocytes rather than a classical TC shape, thus questioning whether DCLK1 marks human TCs. In this context we recently tested a commercially available antibody (ab31704) on human colonic material, and failed to obtain convincing immunolabeling for DCLK1 (23), a conclusion also reached by Banerjee et al. (40). Differences in DCLK1 immunoreactivity in human GI epithelium could be due to differences in protocol and fixation methods or that human TCs simply do not express DCLK1. By contrast, cyclooxygenase enzyme (COX)-1 has been identified as a reliable marker for human TCs (9, 21, 23). Gerbe et al. (9) were first to validate COX-1 as a marker for human TCs by co-staining for other markers such as SRY-box transcription factor 9 (SOX9) and hematopoietic prostaglandin D synthase. Other useful markers include p-EGFR, arachidonate 5lipoxygenase (ALOX5), advillin (AVIL), girdin and choline acetyltransferase (ChAT) (20, 21, 40, 45-47). Notably, none of these markers are TC specific. However, within the epithelium, they are often restricted to TCs, and a combination of one of these

markers with a marker for epithelial cells, such as EpCAM or E-Cadherin, therefore faithfully identifies TCs (21).

Tuft Cell Input

TCs monitor the intestinal lumen using a variety of apical receptors making them proficient sentinels of the GI tract as they respond to a broad palette of substances. Apical signaling includes taste receptors similar to those of taste buds of the tongue and soft palate. These receptors are usually divided into three types: Type 1 transduces signals for sweet and umami substances; type 2 for bitter substances, and type 3 registers sour substances (48). TCs express type 1 (Tas1Rs) and type 2 (Tas2Rs) (49-53). Another important receptor is succinate receptor 1, SUCNR1, which respond to succinate, a metabolite secreted by certain symbiotic bacteria, protists and helminths (54, 55). Receptor tyrosine kinases have also been identified in TCs, although the mechanism of activation is unclear. Basolateral signaling include other receptors, although not fully elucidated, such as gamma aminobutyric acid (GABA)receptors in small intestinal TCs (55, 56), dopamine receptor Drd3 (22, 55), and the orphan adhesion G protein receptor ADGRG2 (26). Lastly, a recent study report a pathway for IgG activation of intestinal TCs, although only in a minor subpopulation of about 2.75% of TCs in the mouse small intestine (57).

Intestinal Chemosensing by Taste Receptors

Chemosensory taste receptors in the GI tract comprise G proteincoupled receptors of the two taste receptor families Tas1R and Tas2R. Several subtype receptors are expressed by different cell types all along the murine GI tract (58), and a varying expression pattern of these bitter taste subtype receptors are also identified along the human GI tract (48, 59–68). However, their region-specific expression distribution is not systematically corroborated by immunohistochemical studies.

The first indication of chemosensory functions of TCs was reported in 1996. Here, Höfer et al. (69) identified brush cells of the gut expressing α -gustducin (a G-protein subunit involved in sweet, umami, and bitter taste signal transduction), which was thought to be specific for taste receptor cells of the tongue. Recent reports indicate the presence of Tas1R and Tas2R taste receptors in TCs of the GI tract (49-53) as well as of the complete set of downstream entities in the taste receptor signaling cascade. Out of these, the monovalent cation conductive and calcium-regulated Trpm5 channel is specific for the TCs (although not specific for taste signaling transduction) in the GI epithelium, Figure 2 (32, 38, 48, 50, 68–70). Although the mechanism has not been fully established, TCs use taste receptors to identify helminths and protozoa. For instance, the Tas1R3 in mice is central in TC activation by the protozoa Tritrichomonas muris, but not by the helminth Heligmosomoides polygyrus (49). On the contrary, Heligmosomoides polygyrus appear to actually inhibit tuft and goblet cell gene expression and expansion (71). Furthermore, in an experimental model with small intestinal organoids from mice, Luo et al. (51) observed calcium responses and interleukin (IL)-25 secretion from TCs when stimulating with excretory/secretory products, and extract from the helminth Trichinella spiralis as well as the bitter compound salicin - a response mediated through Tas2Rs (sensing bitter substances) expressed by TCs.



lipoxygenase (ALOX5) into prostaglandin E_2 (PGE₂), prostaglandin D_2 (PGD₂), and cysteinyl leukotriene C_4 (LTC₄), respectively. Basolateral secretion of IL-25 and ACh occurs through yet unknown mechanisms, whereas secretion of PGE₂ and PGD₂ is facilitated through the prostaglandin transporter (PGT), and LTC₄ possibly through the multidrug resistance protein 1 (MRP1). Basolateral signaling involves GABAA (ligand-gated chloride channel) and GABAB (G-protein coupled) receptors. Stimulation initiates cell repolarization mediated by GABA receptors.

These responses were blocked by the bitter-taste receptor inhibitor allyl isothiocyanate. Similarly, denatonium, another bitter substance, elicits a strong increase in intracellular calcium levels in *ex vivo* stimulated colonic TCs (72). However, more studies are required to determine the role of taste receptors in TC activation.

Succinate Receptor 1, SUCNR1

Succinate, a microbial metabolite, stimulates IL-25 secretion from TCs *via* SUCNR1, thus triggering a type 2 immune response, including activation of type 2 innate lymphoid cells (ILC2), eosinophilia and differentiation of tuft- and goblet cells (54). The highest TC expression of SUCNR1 has been observed in ileal TCs, although some SUCNR1 expression was also observed in colonic samples (73).

Gamma Aminobutyric Acid (GABA)-Receptors

The function of GABA is best characterized in the central nervous system but is also expressed in other tissues such as the GI tract. Although the role of GABA signaling in GI function (e.g., motility, blood flow, secretion, and immune cell activation) is not fully understood, it is well established that neurons of the myenteric plexus and mucosal endocrine-like cells synthesize

and secrete GABA (74). Studies examining the role of GABA in modulating immune function indicates both protective and aggravating properties of GABA signaling, suggesting a complex signaling pattern (75, 76). TCs were shown to express both GABA_A and GABA_B receptors (55, 56), although the significance of this has yet to be elucidated.

Tuft Cell Network Signaling

TCs are first responders to a range of luminal stimuli and diligently communicate with the mucosal immune apparatus and neuronal network. Most TCs express an enzymatic apparatus enabling them to synthesize a variety of paracrine and endocrine effector molecules, including IL-25 (32, 38, 50), ACh (21), eicosanoids (55, 77), β -endorphins (9), and TSLP (22) (**Figure 2**).

IL-25

TCs represent a major epithelial source of IL-25, a critical cytokine for the initiation of type 2 immune responses against helminths and protozoan parasites - and potentially other enteric pathogens - as part of the small intestinal immune system (32, 38, 40, 50, 54, 78). During homeostasis, TCs secrete small amounts of IL-25, while secretion is substantially increased upon activation, e.g., during helminth infection. A subsequent rapid IL-25 mediated expansion of ILC2s provides a crucial early source of IL-13, which in turn stimulates differentiation of intestinal stem cells toward a tuft- and goblet cell fate, and is key to the intestinal "weep and sweep" response expelling intruding parasites (32, 38). ILC2s also secrete IL-4, IL-5 and ACh, all contributing to parasite clearance (32, 79, 80). Several other immune cells of the lamina propria, such as natural killer T cells (81) and nuocytes (an innate type 2 immune effector leukocyte) (82), also play a role in the IL-25 anti-helminth response circuit. Their contributions remain to be fully elucidated, unlike TCs, that are proven indispensable for mounting a proper anti-helminth response (32).

Eicosanoids

Expression of COX-1 and -2 as well as 5-lipoxygenase enables TCs to synthesize prostaglandin D_2 (PGD₂) and $-E_2$ (PGE₂) and cysteinyl leukotriene C_4 (LTC₄) (55), with important physiological roles both during homeostasis (e.g., GI motility, secretion, and mucosal protection) as well as under pathological conditions (e.g., inflammatory bowel disease (IBD), and colorectal cancer) (83, 84). Specifically, intestinal inflammation has been associated with increased COX-2 enzyme activity and PGE₂ production (85, 86), although PGE₂ can exert either proor anti-inflammatory effects depending on the context (87). LTC₄ is a further potent inflammatory mediator, synthesized via the lipoxygenase pathway when stimulated by helminths or protozoa through a yet unknown receptor mechanism (77, 88, 89). While LTC₄ works synergistically with IL-25, it is strikingly dispensable for anti-helminth immunity. This is in surprising contrast to IL-25, which is essential to worm-expulsion. Interestingly, both the synthesis and degradation of LTC₄ are more rapid than that of IL-25 (77, 90).

Acetylcholine

ACh, the main neurotransmitter in cholinergic neurons, is also an important modulator of epithelial proliferation and intestinal physiology (91). In the human GI tract, TCs are the only epithelial cell type expressing ChAT, the enzyme required for ACh production (21). It is still unclear what stimulates ACh synthesis in TCs, but it is likely a result of activation of the canonical taste receptor signaling pathway (92, 93). Recently, an important and thorough study was published on possible debris products of bacterial origin as stimuli for ACh-dependent autoand paracrine pathways in chemosensory cells specific for activation of tracheal clearance in mice. Conclusions in the study are certainly relevant for TC function in the intestine (94). Their study shows that a cluster of three subtype bitter taste receptors (-126, -135 and -143), when eliminated, results in a significant 44% reduction of an ACh-induced peak effect for tracheal clearance. The same was true for contraction induced in urinary bladder strips by a significant 30% reduction. The tracheal chemosensory cells express 18 subtypes of which some, but not all, could be discarded as involved in the remaining ACh-signaling. Hence, more studies on this subject, also for intestinal TCs, are needed before a firm conclusion can assess additional involvement of bitter taste receptors on downstream ACh-induction as well as in general.

In neurons, vesicular acetylcholine transporter (VAChT) is responsible for loading ACh into secretory organelles for secretion. According to Schütz et al. (72), murine intestinal TCs do not express VAChT except for few TCs in the ascending (proximal) colon. VAChT immunoreactive epithelial cells of the human sigmoid (distal) colon, however, have previously been described (95). As such, the release of ACh from TCs is still not clarified and may involve different secretion mechanisms compared to those of neurons. For instance, transporters that can export ACh from colonic epithelium cells in a non-vesicular fashion have been described in human sigmoidal epithelium as well as choline importers and several enzymes needed for ACh synthesis. Future work should address potential co-localization of these transporters with TCs (96).

β-Endorphin and Thymic Stromal Lymphopoietin

Intestinal TCs also secrete β -endorphins and thymic stromal lymphopoietin, although the importance of these TC derived effector molecules is unclear (22). For instance, opioids, such as β -endorphin, are usually associated with analgesic effects, but are also involved in basic GI physiology including intestinal motility, secretion, and absorption (97). In the GI tract, expression of β endorphins is restricted to TCs (9, 98). Notably, all intestinal TCs produce β -endorphins (9), and the secretion hereof depends on the calcium-regulated Trpm5 channel, which is specific for TCs within the intestinal epithelium. TSLP, on the other hand, is a cytokine promoting Th2 immunity similar to IL-25 (99). Some basolateral TSLP secretion derives from TCs (38), and secretion is increased upon tissue damage and exposure to pathogens (100, 101). Of note, TSLP appears to be involved in various immunological conditions including IBD (102).

TUFT CELLS AND GASTROINTESTINAL DISEASES

Mouse Obesity as a Chronic Low-Level Inflammation Model

Obesity is a condition with gradually increasing prevalence worldwide, representing a risk factor for developing a variety of diseases including several GI disorders and changes in GI motility. Recently, the effects of high fat diet and induced obesity on small intestinal TCs was examined in a mouse model. Although the ratio of number of TCs to total epithelial cells was unaltered, TC specific expression of IL-25 and TLSP was decreased (56). Altered GABA (A) and GABA(B) receptor activation pathways correlated positively with the altered expression of TC signature genes IL-25 and TSLP, suggesting an association between TC numbers and GABA receptor signaling (56).

Infections of the GI Tract

An essential role of TCs in anti-helminthic and anti-protozoan immunity has been established. TCs respond to such parasites by secreting IL-25 and LTC₄ initiating a type 2 immune response ultimately expelling the parasitic agent (32, 38, 50, 77). Generally, immunomodulatory effects and potentially beneficial effects of parasitic worms on intestinal function, inflammation and mucosal healing have been studied in detail in mice and humans (103). Considering the role of TCs in anti-helminthic/ protozoan immunity it is tempting to speculate that TCs could mediate these effects. However, only sparse data exists on humans, and so far, helminth therapy appears to have no obvious effect compared to placebo treatment in the management of human disorders like IBD (104, 105). Furthermore, helminth therapy has its drawbacks given that prolonged exposure may cause complications as the host is exposed to a wide spectrum of helminth-derived products, including potent antigens and inflammatory stimuli, in addition to desired helminth immunomodulators. Although short-term administration of Trichuis suis ovae (pig worm eggs) appears to be safe and tolerable in humans (104, 106, 107), individuals exposed to chronic helminth infections appear to develop an immune dysregulation. This dysregulation results in hypo-responsiveness and difficulties in mounting proper immune responses, e.g. towards neurotropic flaviviruses, an effect attributed to the persistent immune activation (108, 109). The effect is thought to be mediated by TCs via an IL-4 signaling pathway resulting in a compromised $CD8^+$ T cell response (109).

TCs also appear to be a target of norovirus, which is the leading cause of gastroenteritis worldwide. In mice, TCs express high levels of murine norovirus receptors, CD300lf, enabling infection, while also serving as viral reservoirs causing chronic infections and viral shedding for weeks following the acute phase of infection (110, 111).

Yet another candidate driver of TC induced immune responses is succinate, which indirectly activates the host immune system through SUCNR1 receptors also expressed on TCs (54). Succinate, a metabolite produced by commensal bacteria and helminths, has been shown to induce ATOH1-independent (small intestinal) TC expansion. In line with the protective TC functions described above, administration of succinate in two mouse models suppressed ileal inflammation and restored epithelial architecture, accompanied by an enhanced anti-parasite-, and a reduced type Th17 response (40). However, no increase in TC numbers following oral succinate administration was observed in the cecum and colon of mice, limiting the succinate response via TCs to the small intestine (54). How increased luminal succinate levels link to cytokine expression and downstream responses remains to be shown. Interestingly, vancomycin-induced dysbiosis (potentially leading to colonization of Clostridioides (C). difficile) has been associated with TC hyperplasia. Although a direct involvement of TCs in *C. difficile* infections has so far not been studied, neither in mice nor in humans, some studies indicate a correlation (112). In C. difficile pathophysiology IL-23 is considered mainly responsible for the destructive inflammatory response leading to severe tissue damage (113, 114), whereas IL-25, the main TC derived cytokine, demonstrates protective properties (115). The effect is thought to be mediated through IL-25 stimulated IL-13 release and recruitment of eosinophils; a hallmark of type 2 immunity and a TC characteristic (116).

Inflammatory Bowel Disease

IBD is a group of chronic idiopathic inflammatory disorders characterized by latent symptom-free periods and recurrent flares with varying degrees of inflammation of the GI tract (117–119). The two most prevailing entities are ulcerative colitis (UC) and Crohn's disease (CD). Symptoms such as abdominal discomfort, urge to defecate, (bloody) diarrhea and fatigue may be present in both disorders, they differ greatly in terms of pathophysiology although the pathogenesis is still not fully elucidated (119).

Accumulating evidence has demonstrated that T helper cell (T_h) 1-related cytokines (e.g., tumor necrosis factor- α (TNF- α), interferon- γ and IL-12) as well as T_h 17-associated cytokines like IL-17A, IL-21 and IL-23 are markedly increased in the inflamed mucosa of patients with CD, whereas the cytokine profiles in inflamed mucosal areas in patients with UC seem to exhibit increased production of T_h 2 associated cytokines, including IL-4, IL-5, and IL-13 (119, 120). By contrast, Su et al. (121) found IL-25 to be significantly decreased in both serum and inflamed intestinal mucosa of patients with flaring IBD compared to healthy controls. The same trend was observed in non-inflamed tissue and serum from patients with quiescent UC and CD. After treatment with infliximab (a TNF- α inhibitor) for flaring of IBD, serum IL-25 levels were, however, restored (121).

Thus, a critical role of IL-25 has been suggested in the pathogenesis of IBD, and accordingly increasing IL-25 levels could be a potential therapeutic option for IBD. As TCs are potent producers of IL-25 it is tempting to speculate about a potential role in IBD pathophysiology. However, the abovementioned study by Su et al. (121) did not specifically investigate the role of TCs, nor did it differentiate between epithelial and subepithelial IL-25 expression. Several immune cells of the lamina propria produce and secrete IL-25, and TCs alone might not fully account for the observed changes (121, 122). Banerjee et al. (40) did observe a reduced TC population in inflamed ileal tissue in CD, which is similar to what was demonstrated in the non-inflamed

colon of patients with UC in clinical, endoscopic and histologic remission (23). Whether this reduction in TC numbers in both UC and CD is a consequence of the disease or an actual contributing pathogenic factor remains unclear.

Coeliac Disease, Duodenal Ulcer and Acute Duodenitis

A recent study by Huh et al. (47) described the TC abundance in biopsy specimens from pediatric patients with acute and chronic duodenitis. TC numbers were lower in patients diagnosed with coeliac disease and duodenal ulcers when compared to healthy specimens. In acute duodenitis, TC numbers decreased as the severity of inflammation increased, indicating a loss of TC differentiation or death during active inflammation. Whether inflammation precedes TC decimation or is a consequence hereof is still unclear.

Gastric and Colorectal Neoplasia

TCs are massively expressed in locations of the ventricle, including epithelial areas of the cardia-near esophagus, where they seem to protect and repair the epithelium, for instance after acidic injury by gastro-esophageal reflux ultimately leading to Barrett's adenocarcinoma (123). Recent insights kindle speculations about the genesis of sporadic intestinal tumors where TCs might be involved in the tumor etiology, although a direct involvement of TCs in the development of colorectal neoplasia (CRN) has not been reported. In this context, Roulis et al. (124) found rare pericryptal fibroblasts expressing COX-2, located in close proximity of the stem cell zone, that stimulated the expression of markers associated with regeneration (Sca-1+ cells) and thus demonstrated possible tumorinitiation properties via paracrine PGE₂-signaling (124, 125). Similarly to TCs, these mesenchymal cells expressed both COX-1 and COX-2. In human colonic mucosa, COX-1 positive TCs are identified located within the epithelium together with subepithelial COX-1 expressing cells that may represent the pericryptal fibroblasts, Figure 3 (126). Both the TCs and subepithelial cell types are located near the stem cell zone. As TCs seem to be a major source of prostaglandins they may also be a contributing cell type to CRN development (9). Indeed, an imbalance of arachidonic acid metabolism, including prostaglandin synthesis as well as production of free oxygen radicals, is suspected to be central to the pathogenesis of CRN (83, 127).

DISCUSSION

The number of studies examining TCs is increasing rapidly. Several breakthroughs elucidating TC function and significance have recently been made, including the identification of TCs as first responders to intestinal parasitic infections. Despite their relatively low numbers, TCs are important sentinels of the GI tract monitoring the luminal content with an array of luminal receptors responding to a broad palette of substances such as succinate, canonical tastes (sweet, bitter, umami) and secretory agents from parasitic helminths and protozoa. A close physical and functional relation to enteroendocrine cells and the



FIGURE 3 | Cyclooxygenase 1 (COX-1) positive tuft cells (TC) of the human colon. Illustration of colonic TCs from healthy human sigmoid biopsies (controls). Orange arrows indicate TCs, while white arrows indicate what could be interpreted as subepithelial/pericryptal fibroblasts. Adapted from "Possible predisposition for colorectal carcinogenesis due to altered gene expressions in normal appearing mucosa from patients with colorectal neoplasia" by Petersen et al. (126).

underlying immune cells and nervous apparatus enable TCs to mediate responses to metabolic shifts in the intestinal lumen expelling potentially harmful agents such as parasitic worms.

However, our knowledge of TCs is still limited, especially concerning their involvement in human diseases. Observations from murine studies suggest an involvement of TCs in multiple diseases, e.g., neoplasia and colitis, but only few studies deal with human TCs, partly due to a late identification of relevant markers of human TCs. These are still being investigated, while possible markers, although not TC specific, include COX-1, p-EGFR, SOX9, ALOX5, AVIL, girdin and ChAT.

It appears that the number of intestinal TCs change during human pathologies. Whether this is merely a consequence of the disease or an actual contributing pathogenic factor, is a question that requires further exploration. With the help of new, specific markers and an improved understanding of TC heterogeneity, it is of importance to outline their potential as future diagnostic/ predictive markers and to understand potential therapeutic benefits of modulating TC abundancy.

It is tempting to speculate that stimulating TC expansion in parallel with other cell populations, such as goblet cells, ILC2s and eosinophils, could promote epithelial barrier integrity and homeostasis and have a preventive effect on disease progression and risk of/severity at a new flare. TC specific activation may be achieved by substances activating canonical taste receptor signaling pathways, e.g., certain sweet, bitter, and umami substances, or even parasitic derived products. The isolation and administration of certain helminth derived products could have a beneficial effect and circumvent the negative consequences of chronic infections with helminths or other parasites. An illustration of the hypothetical beneficial effects of TC stimulation is shown in **Figure 4**.

Accordingly, a better understanding of TC biology may ultimately contribute to the efforts of prevention and development of novel treatment strategies against a wide range of human disorders of the GI tract.



FIGURE 4 | Tuft cells (TC) as potential therapeutic targets. Illustration of hypothetical beneficial effects on mucosal integrity when stimulating TCs. Left and right side show intestinal mucosa before and after TC stimulation, respectively. Upon stimulation, TCs increase secretion of interleukin (IL)-25 and cysteinyl leukotriene C_4 (LTC₄) which in turn stimulate lamina propria immune cells such as, type 2 innate lymphoid cells (ILC2), natural killer T cells and nuccytes to produce and release type 2 inflammatory cytokines (e.g. IL-4, IL-5 and IL-13). This initiates differentiation of T helper type 2 cells (T_h2), recruitment of eosinophils and stimulation of intestinal stem cells leading to tuft cells and goblet cell differentiation and hyperplasia.

AUTHOR CONTRIBUTIONS

This study was conceived by SH and ON. SH wrote 1st draft. LK, AH, NB, KJ, and ON critically edited and approved the manuscript. All authors contributed to the article and approved the submitted version.

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