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Control of myeloid cell functions by nociceptors

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The immune system has evolved to protect the host from infectious agents, parasites, and tumor growth, and to ensure the maintenance of homeostasis. Similarly, the primary function of the somatosensory branch of the peripheral nervous system is to collect and interpret sensory information about the environment, allowing the organism to react to or avoid situations that could otherwise have deleterious effects. Consequently, a teleological argument can be made that it is of advantage for the two systems to cooperate and form an "integrated defense system" that benefits from the unique strengths of both subsystems. Indeed, nociceptors, sensory neurons that detect noxious stimuli and elicit the sensation of pain or itch, exhibit potent immunomodulatory capabilities. Depending on the context and the cellular identity of their communication partners, nociceptors can play both pro- or anti-inflammatory roles, promote tissue repair or aggravate inflammatory damage, improve resistance to pathogens or impair their clearance. In light of such variability, it is not surprising that the full extent of interactions between nociceptors and the immune system remains to be established. Nonetheless, the field of peripheral neuroimmunology is advancing at a rapid pace, and general rules that appear to govern the outcomes of such neuroimmune interactions are beginning to emerge. Thus, in this review, we summarize our current understanding of the interaction between nociceptors and, specifically, the myeloid cells of the innate immune system, while pointing out some of the outstanding questions and unresolved controversies in the field. We focus on such interactions within the densely innervated barrier tissues, which can serve as points of entry for infectious agents and, where known, highlight the molecular mechanisms underlying these interactions.

KEYWORDS

neuroimmune interactions, nociceptors, myeloid leukocytes, neuropeptides, control of immunity

1 Introduction

Innate immunity is, in some form, present in virtually all multicellular organisms (1, 2). The functions of the innate immune system include not only elimination of pathogens by direct killing and activation of the adaptive immune system (in organisms in which it is present) (3), but also the induction, modulation and resolution of inflammation, tissue

repair (4–6), and control of metabolism (7). Functionally, innate immune cells lack the receptor repertoire diversity of the adaptive immune system, but instead express a selection of germlineencoded receptors that allow recognition of conserved pathogenassociated molecular patterns (PAMPs) as well as damageassociated molecular patterns (DAMPs). Such receptors imbue the innate immune cells with the ability to rapidly respond to the presence of pathogens as well as signs of cellular distress in their environment (8). Furthermore, tissue-resident innate immune cells are strategically located throughout the body and are concentrated in barrier tissues, such as the skin or mucosal surfaces. Thus, in most cases, innate immune responses are initiated within minutes to hours after insult and constitute the body's first line of defense (9).

On a single-cell level, innate immune cells represent a heterogenous group of leukocytes, which differ in their function, tissue distribution, migratory properties, life-span, turnover, origin, and plasticity. Historically, the various cell types have been differentiated based on their morphology, physiological functions and phenotypes as well as, more recently, their ontogeny. Most innate immune cells are of myeloid origin, i.e. they arise from a common myeloid progenitor (CMP) in the bone marrow (BM) in adults, and from erythro-myeloid progenitors (EMP) in the yolk sac during development (10). Complex relationships between further developmental stages of various myeloid cell types exist, and our understanding of the plasticity of their respective progenitors remains incomplete. Nonetheless, based on phenotypical similarities, myeloid cells can be broadly divided into three families: the mononuclear phagocytes (dendritic cells, macrophages and monocytes) (11), polymorphonuclear granulocytes (neutrophils, basophils and eosinophils) (12), and mast cells (13). In addition to myeloid cells, innate lymphoid cells (ILCs) originating from the common lymphoid progenitor (CLP) have been a focus of intensive research in recent years (14). Notably, neural control of ILCs has been described and reviewed recently (15). Consequently, this review will specifically focus on the effects that nociceptors have on the myeloid immune cell compartment.

Nociceptors are specialized afferent nerve fibers that respond to noxious stimuli such as physical damage, excessive pressure, irritant chemicals, or extremes of temperature and initiate withdrawal/ avoidance behavior or irritant removal (16). Based on the modality of the stimuli that they detect, nociceptors can be unimodal or polymodal, i.e. responding to a single or several types of noxious stimuli, respectively (17). Historically, nociceptors have also been classified by the physiological properties of their axons into myelinated A fibers and nonmyelinated C fibers, with the latter being the more prevalent group (16-18). More recently, transcriptomic sequencing techniques have begun to define nociceptor subsets based on their gene expression signatures. Indeed, a recent single-cell RNA sequencing study of neurons in murine dorsal root ganglia (DRGs) (19), has proposed classification of nociceptors in into six broad subsets: three non-peptidergic (NP1-3), two peptidergic (PEP1-2) and one tyrosine hydroxylase (TH)-expressing population. Within this classification, the NP1 subset appears to be involved in inflammatory pain, NP3 in inflammatory itch, and all three NP subsets in pruritus in general. The PEP1 population corresponds to thermo-sensitive neurons, whereas PEP2 represent lightly-myelinated A δ nociceptors (19), which normally respond to mechanical or thermal stimuli (18). Somewhat confusingly, both peptidergic and non-peptidergic nociceptors within this classification express different patterns of neuropeptides (Table 1), several of which are known to modulate myeloid cell functions (see below). Of note, more recent studies have suggested dividing nociceptors into as many as 10 different subsets (20). Additionally, even nociceptors belonging within the same subset have been shown to express specific gene patterns depending on the organs they innervate (21), indicating that additional heterogeneity exists and the classification of nociceptors is anything but straightforward.

Nevertheless, nociceptors share certain molecular features that allow their identification and selective experimental manipulation. For example, the NaV1.8 voltage-gated sodium channel is expressed in approximately 90% of all nociceptors and is often considered a pan-nociceptor marker (22). Consequently, although NaV1.8 is also found in certain low-threshold mechanoreceptors (23), NaV1.8-Cre 'knock-in' mice (24) have been widely used to target and manipulate nociceptors by genetic means (25–28). Additionally, prominent transient receptor potential (TRP) cation channels have been identified, which often correlate with the specificity of nociceptors for various noxious stimuli including heat (TRPV1), chemical irritants (TRPA1), cold (TRPM8), and others (29). As a

TABLE 1 Expression of the main neuropeptides with immunoregulatory potential toward myeloid cells across nociceptor subsets at the steady state, as reported by the (19) single cell RNAseq dataset.

	PEP1	PEP2	NP1	NP2	NP3	тн
CGRPα	++++	++++	+	++++	+	+
CGRPβ	++	+++	+++	++++	_	+++
Adrenomedullin	+	+	_	-		
Intermedin	_	_	_	_	_	+
SP	++++	_	+	+	+	+
VIP	_	_	_	_	_	-
PACAP	+++	_	+	+	-	+

- no expression detected, + expression in <25% of cells, ++ expression in 25 - 50% of cells, +++ expression in 50 - 75% of cells, and ++++ expression in > 75% of cells.

result, genetic, as well as chemical means of targeting TRP channels have been developed, which allow for manipulation of nociceptors of a given specificity (30). Experimental means of targeting TRPV1⁺ nociceptors in particular have been widely used in the field, as exemplified by TRPV1-Cre 'knock-in' mouse models (31) and the use of TRPV1 agonists, such as capsaicin and resiniferatoxin (RTX), which can hyper-activate and specifically ablate the TRPV1⁺ nociceptors (32, 33).

Anatomically, nociceptor cell bodies are located in the trigeminal ganglia for nociceptors innervating the head and in the DRGs for nociceptors innervating all other parts of the body. Morphologically, nociceptors are pseudo-unipolar neurons with one axon that bifurcates into a proximal and a distal branch. The proximal process terminates in the dorsal horn of the spinal cord or in sensory nuclei of the brainstem, while the distal processes project to peripheral target tissues where they terminate in free endings (18). In addition to their afferent function, nociceptors also exhibit efferent modalities, which are believed to be mediated by several mechanisms including the axon reflex (backpropagation of action potentials through collateral branches) (34) and the antidromic activity (conduction in the reverse direction) (35) (Figure 1). Notably, as we will discuss in detail below, efferent functions of nociceptors include the peripheral release of neuropeptides, which act on cells in their proximity, including myeloid leukocytes (34, 35). Of note, the impact of nociceptors and nociceptive neuropeptides on specific target cells depends, at least in part, on the target tissue (36). For example, in the skin, myeloid immune cells are the main targets of nociceptors (26, 37, 38), while in lymph nodes (LNs), the effect on LN-resident myeloid cells is more limited. Instead, non-immune stromal cell types have been identified as the primary communication partners of nociceptors within LNs (21). Lastly, further highlighting how intimately intertwined immune and peripheral nervous systems are, nociceptors express, and are functionally impacted by many of the receptors traditionally thought of as "immune", including Fc receptors (39), PRRs (40), and checkpoint molecules such as PD-L1 (41).

2 Neuropeptides and neuropeptide signaling in immune cells

Neuropeptides are considered the main communication signals that nociceptors emit to impact immune responses, and, indeed, expression of neuropeptide receptors is widespread among myeloid immune cells (Table 2). In addition to their immune-modulatory properties, however, neuropeptides also influence numerous other cell types and, as a result, their specific impact on immune cells has often been tested in reductionist systems in vitro or ex vivo. While this strategy is suitable to identify the molecular mechanisms of action within given cell types, it ignores the tissue context and may not accurately reflect the role that neuropeptides play in physiological settings. Complicating matters further, the concentrations of neuropeptides that can be reached in target tissues due to nociceptor activation remain poorly defined. Consequently, despite a large body of literature detailing the effects of neuropeptides on specific cells in isolation, our understanding of how they impact the function of the immune system as a whole remains far from complete. Finally, it is important to stress that nociceptors are not the only source of neuropeptides (43-45). Consequently, studies in which cells/ animals are directly exposed to a neuropeptide (or its inhibitor) can only elucidate effects of the neuropeptide itself, but may not necessarily define the physiological effect of nociceptors.

Finally, it should be noted that many neuropeptides, including calcitonin gene-related peptide (CGRP), Substance P (SP) and vasoactive intestinal peptide (VIP), exhibit structural similarities to cationic antimicrobial peptides and, as a result, can exert, at least to some degree, antimicrobial activity (46, 47). The LD_{50} described for most bacterial strains, however, lies in the high micromolar range (48), arguably, well above the concentrations that nociceptor-derived neuropeptides are expected to reach within tissues. Consequently, the physiological relevance of this phenomenon remains unclear, though, a possible role in the regulation of gut microbiota has been suggested (49).



Schematic depiction of the physiological organization of and signal transmission by nociceptors. Arrows indicate the direction of action potential propagation. Black arrows correspond to the afferent transmission of signals elicited by peripheral activation of nociceptors by a noxious stimulus, terminating in the spinal cord and leading to the sensation of pain or itch. Blue and green arrows correspond to the efferent transmission by means of antidromic activity (conduction in the reverse direction) and axon reflex (backpropagation of the action potential through collateral branches) respectively, leading to the peripheral release of neuropeptides.

Neuropeptide	Receptor	DCs	Macrophages	Monocytes	Neutrophils	Eosinophils	Basophils	Mast cells
CGRP/AM/IM	Calcrl	+/++	++/+++	+/++	++	+/++	-/+	+/++/+++
CGRP	RAMP1	++	-/++/+++	++	++/+++	-/+/++	-/++	++/+++
AM/IM	RAMP2	-/+	-/+/++/+++	-/+	-/+	-/+/++	-/+	-/+
	RAMP3	-/+/++	+/++	-/+	-/+	-/+	-/+	+/++
SP	NK1R	-/+	-/+	-/+	-/+	-/+	-/+	-/+
	MRGPRD	-/+	-/+	-/+	-/+	-/++	-/++	++
	MRGPRA1	-/+	-/+	-	-	-/+	-/+	-/+
VIP/PACAP	VPAC1	+	-/+	-/+	+	+	-/+	+/++
	VPAC2	-/+	-/+	-/+	-/+	-/+	-/+	+
PACAP	PAC1	-/+/++	-/+/++	-/+	-/+	-/++	+/++	-/++

TABLE 2 Expression of neuropeptide receptors on immune cells based on the RNAseq and microarray data deposited in the Immgen database (42) (https://www.immgen.org/Databrowser19/DatabrowserPage.html).

- no expression, + how expression, +++ medium expression, +++ high expression. Multiple symbols signify uncertainty due to differences between RNAseq and microarray datasets, or variability in expression between cell subsets.

2.1 CGRP neuropeptide family

The CGRP family of neuropeptides includes CGRP α (50) and CGRP β (51), which are thought to be functionally redundant (52), adrenomedullin (AM) (53), and intermedin (54) (also known as adrenomedullin 2 (AM2)). All the CGRP-family neuropeptides signal through a heterodimeric transmembrane receptor comprising a G protein-coupled receptor (GPCR), calcitonin receptor-like receptor (Calcrl), and one of three known receptoractivity modifying proteins (RAMPs) (55). While Calcrl is the signal-transducing, shared subunit, the RAMP proteins dictate the specificity of the complex with RAMP1-Calcrl being the CGRP receptor and RAMP2/3-Calcrl the AM_{1/2} receptors (56). Of the three related neuropeptides, CGRP is the most extensively studied, and its pleiotropic effects are too numerous to be all listed here. The interested reader is referred to an excellent in-depth review (52). Within the scope of the immune system, the effects of CGRP are mostly thought of as anti-inflammatory (25, 28, 57-60), however, several recent studies have highlighted potent pro-inflammatory functions of this neuropeptide in specific contexts (37, 38). The roles of both AM and AM2 remain much less explored; however, they too have been shown to exert some broadly immunoinhibitory functions (61-65).

Several types of G α -proteins are known to couple with the CGRP receptor, leading to the activation of a variety of signaling pathways (Figure 2A), some of which appear to be cell-type specific [reviewed in (55)]. Most prominently, G α_s coupling leads to the activation of adenylyl cyclase (AC) (66, 67) and intracellular accumulation of cyclic AMP (cAMP) (68). Conversely, G α_i coupling in certain cell types has been shown to inhibit AC and, instead, to drive JNK activation (69). Finally, CGRP signaling *via* G α_q activates phospholipase C (PLC)- β and protein kinase C (PKC) (70, 71). Consequently, it is tempting to speculate that a preferential engagement of certain G α subunits by other GPCRs could decrease their overall availability and, thus, regulate the outcomes of CGRP receptor ligation. Lastly, it has been speculated that CGRP may also

signal through $G\alpha$ -independent pathways, however, the biological relevance of this mechanism is unclear (55).

The immuno-inhibitory effects of CGRP are thought to be mainly due to the activation of AC. The resultant accumulation of cAMP activates protein kinase A (PKA) and upregulates the inducible cAMP early repressor (ICER) (72) which, in turn, prevents recruitment of the transcription factor ATF-2 to, among others, the Tnfa promoter (73, 74). Additionally, PKA can phosphorylate the cAMP response element-binding protein (CREB), resulting in nuclear translocation of the CREB-regulated transcriptional cofactors (CRTC) 2 and 3, and expression of the anti-inflammatory cytokine IL-10 (75). Finally, cAMP also induces the expression of a transcriptional repressor, Jdp2, which can bind the p65 NF-KB subunit and prevent its docking onto target promoters (57). Additional, cAMP-independent mechanisms are likely also involved, as increased intracellular cAMP concentration alone is not sufficient to mimic the effects of the neuropeptide (76). Inhibition of IKB kinase β phosphorylation and subsequent inhibition of NF-KB signaling was suggested as one such mechanism (77), however, the exact underlying details remain unclear.

Mechanisms of the proinflammatory actions of CGRP in myeloid cells are unknown. In other cell types, however, they, counterintuitively, also appear to rely primarily on the activation of PKA (78, 79). The reasons why the cAMP–PKA axis could act as both pro- and anti-inflammatory are similarly poorly understood, but could be a result of a differential balance between other signaling pathways that exhibit distinct activities in different cell types and/ or states.

2.2 Substance P

Substance P (SP) is a neuropeptide of the tachykinin family (80) and mainly exerts its functions through one of three GPCRs: the broadly expressed high-affinity neurokinin 1 receptor (NK1R) (81),



and the more recently discovered MRGPRB2 and MRGPRA1 which are selectively expressed by mast cells (82-84) and DCs, respectively (85). The actions of SP in most contexts are pro-inflammatory, and several signaling pathways have been implicated (Figure 2B) (86, 87). Specifically, NK1R ligation leads to the activation of PLC, which generates the second messengers inositol trisphosphate (IP₃) and diacyl-glycerol (DAG). These, in turn, can mobilize calcium from intracellular stores and activate PKC (87). In addition, the phosphoinositide 3-kinase (PI3K)-Akt pathway (88) as well as direct activation of the p38 and ERK1/2 mitogen-activated protein kinases (MAPKs) exert proinflammatory effects by triggering NF-KB (89, 90). NK1R can also activate AC with resultant cAMP accumulation and PKA activation (91). Considerably less is known about the signaling pathway downstream of MRGPRB2. Its engagement ultimately leads to a sustained increase in intracellular calcium levels (92) through storeoperated calcium entry (SOCE) by the calcium sensor stromal interaction molecule 1 (STIM1) and activation of p38 and ERK MAPKs (93, 94). Additionally, SOCE-independent Akt activation has been reported (94). The mechanistic underpinnings of MRGPRA1 signaling in immune cells remain unknown (85).

2.3 VIP and PACAP

Vasoactive intestinal peptide (VIP) and pituitary adenylate cyclase-activating polypeptide (PACAP) are related neuropeptides most commonly upregulated by neurons, including nociceptors, following peripheral nerve injury (95, 96). In nociceptors at steady state, PACAP and VIP often colocalize with CGRP and SP (97, 98), and the tissue content of PACAP is decreased following capsaicininduced nociceptor depletion (97, 99). The actions of VIP and PACAP are considered to be broadly anti-inflammatory, as these neuropeptides inhibit the release of pro-inflammatory cytokines including TNF α , IL-6, IL-1 α and IL-1 β , and enhance expression of the anti-inflammatory cytokine IL-10 by several myeloid cell types (Figure 2C) (95, 96). These effects are thought to be, at least in part, due to the inhibition of TNFa gene expression, which VIP/PACAP control by two independent mechanisms: blocking NF-KB binding to the Tnfa promoter elements, and inhibiting JNK activity, resulting in a decreased phosphorylation of the c-Jun protein and its absence from the CREB complexes docking onto the cAMP response element (CRE) promoter of the Tnfa gene (100). Such mechanisms of action are similar to those of CGRP, as described above (see Figures 2A,C), making it likely that, in addition to *Tnfa*, they apply also to other pro-inflammatory genes. Additionally, cAMP-dependent inhibition of interferon regulatory factor-1 (IRF-1) transactivation (101) and upregulation of IL-10 have been described (102).

Several receptors for VIP and PACAP have been identified. One receptor, PAC1, is specific for PACAP, whereas VPAC1 and 2 bind indiscriminately to both VIP and PACAP (95). All three receptors participate in immune regulation. In particular, PAC1 and VPAC1 are expressed constitutively on myeloid as well as lymphoid cells, while VPAC2 appears to be inducible, especially in T-cells (103). Like other neuropeptide receptors, VPAC1/2 and PAC1 belong to

the GPCR superfamily, and their ligation by agonists leads to coupling to $G\alpha_s$ and activation of AC with downstream accumulation of cAMP (104). Additionally, all three receptors can couple to $G\alpha_{\alpha}$, and VPAC1/2 can also signal *via* $G\alpha_{i}$ to activate PLC and calcium mobilization (105). Finally, all three receptors can activate phospholipase D (PLD) (106), however, the underlying mechanism and the relevance of PLD activation for the effects of VIP and PACAP remain poorly understood (105).

3 Control of myeloid cells by nociceptors

Nociceptors employ multiple neuropeptide-dependent and -independent means to exert control over myeloid immune cells, whereby the downstream consequences often vary with the target cell type. In the following, we will discuss the most prominent mechanisms by which nociceptors act on each of the major myeloid target cell subsets.

3.1 Dendritic cells

Dendritic cells (DCs) are a group of myeloid cells derived from the common DC progenitor (CDP) that include the classical DCs (cDCs) and plasmacytoid DCs (pDCs) (107). pDCs are known for their ability to produce copious amounts of type I and III interferons (IFNs) in response to viral infections, and only exhibit a limited ability to present antigens (108). Notably, pDCs have not been reported to be under the control of nociceptors or respond to nociceptive neuropeptides and will not be further discussed here. Conversely, cDCs (henceforth referred to as DCs) are best known for their ability to take up and present antigens to naive T-cells to activate the adaptive arm of the immune system (109) as well as to induce and maintain tolerance to self and innocuous non-self antigens (110, 111). Additionally, DCs play important roles as sentinel cells within tissues and are involved in pathogen surveillance as well as orchestration of local immune responses by secretion of cytokines, chemokines and other mediators (112). Two subsets of DCs exist - cDC1 and cDC2 - which differ in their phenotypic as well as functional properties. Specifically, cDC1s, identified by their expression of XCR-1 and DNGR-1 (a.k.a. Clec9a), exhibit better ability to present exogenous antigens to CD8⁺ T-lymphocytes, owing to their superior ability to crosspresent (i.e. to process and present antigenic peptides from exogenous proteins on MHC class-I complexes). Conversely, cDC2s, identified by their expression of SIRPa and/or CD11b, are generally thought to be superior in their ability to present antigens to CD4⁺ T-lymphocytes, and they comprise the majority of DCs in most tissues (109, 111). Additionally, significant heterogeneity exists among DCs that reside in different anatomic locations (113). This underappreciated diversity could potentially explain some of the seemingly inconsistent observations discussed below.

Importantly, both DCs and nociceptors are abundant in peripheral barrier tissues such as the skin and mucosal surfaces, which places them within close proximity of each other (114). For example, DCs and nociceptors engage in direct physical interactions in the murine skin (26, 85). Indeed, DCs in both the skin (26, 38, 85) and the airways (115) are important targets of nociceptor derived communication signals. Similarly, Langerhans cells (LCs) - a subset of specialized dermal phagocytes that are of macrophage lineage but display many functional properties of DCs (116) - are also known to associate with nociceptors, and respond to neuropeptides (117) (Figures 3-5). DCs are also abundant in secondary lymphoid tissues, including lymph nodes, which are innervated by a unique subset of nociceptors located in the outermost capsular and subcapsular regions (21). However, most lymph node resident DCs are concentrated in the paracortical T cell zone, which is



Known in vivo effects of substance P on myeloid cells. Upward pointing arrows signify upregulation/activation, downward facing arrows signify downregulation/inhibition



downregulation/inhibition

largely devoid of nociceptors, suggesting that lymph node DCs may be relatively less impacted by nociceptors than their counterparts in peripheral barrier tissues.

CGRP effects on DCs: Nociceptor-derived CGRP exerts critical effects on dermal DCs in some, but not all, settings of experimental skin inflammation: One study demonstrated that nociceptorderived CGRP induced IL-23 production by CD301b⁺ cDC2s upon cutaneous Candida albicans infection (38). By contrast, CGRP was dispensable for nociceptor-induced IL-12 and IL-23 production by dermal DCs upon topical treatment with a TLR-7 agonist, Imiquimod (IMQ), in a model of psoriasiform skin inflammation (26). Both studies utilized similar methods to show that nociceptor ablation diminished the cytokine response of dermal DCs. However, in the context of the C. albicans infection, local injection of CGRP was sufficient to drive the IL-23 accumulation even in nociceptor-depleted mice, and application of a CGRP antagonist reduced the levels of IL-23 in animals with intact nociceptors (38). Conversely, in the IMQ model, CGRP antagonists had no effect (26). The reasons underlying this discrepancy remain unexplained but could, conceivably, be a result of different subsets of nociceptors responding to a cutaneous fungal infection compared to a defined TLR agonist, or different modalities of activation elicited by the two stimuli in DCs and/or nociceptors. Alternatively, microbiota colonizing the skin could potentially have differentially modulated neuroimmune interactions, as has been recently reported in the gut (118). Lastly, it is pertinent that exposure of mice to stress can synergize with IMQ effects and enhance the accumulation of pro-inflammatory cytokines and tissue inflammation. This effect may be mediated, at least in part, by an increase in the expression of SP (119), however the exact molecular underpinnings have not been identified. Nonetheless, these observations underscore the complexity and context-dependency of neuroimmune interactions. Regardless of the initiation events, in both the IMQ and C. albicans models, DCderived IL-23 activated skin-resident yoT-cells and led to an enhanced local T_H17 response and increased neutrophil influx (26, 38). Similarly, in the KC-Tie2 model, in which psoriasiform skin inflammation is driven by overexpression of the angiopoietin



receptor Tie2 in keratinocytes (120), surgical denervation reduced skin pathology as well as DC numbers and IL-23 expression (121). Exogenous administration of CGRP and SP was sufficient to reverse this effect, and pharmacological blockade of CGRP and SP receptors mimicked the effects of denervation (121). Also in a wound healing model, IMQ-activated TRPA1⁺ nociceptors enhanced IL-23 upregulation in DCs, which, in turn, promoted tissue regeneration (122), however, the role of CGRP or other neuropeptides was not investigated. Further support for CGRP having immuno-stimulatory properties comes from a recent study in which pro-inflammatory cytokines were observed to accumulate in murine skin after optogenetic activation of cutaneous nociceptors in a CGRP-dependent manner (37). Similarly, repeated activation of TRPV1⁺ nociceptors by means of a circumneural sciatic nerve implant induced local inflammation and enhanced the inflammatory response to local adjuvant injection (123). Although neither study determined the exact cellular mechanism, it appears likely that CGRP action on DCs was involved.

Several other studies have demonstrated that CGRP can also have anti-inflammatory and immuno-modulatory effects on DCs. For example, CGRP-deficient mice showed an increase in DC infiltration into the skin after UV-B irradiation overexposure, though whether this was due to a direct effect on DCs or involvement of other cells was not established (124). In vitro, LCs exposed to CGRP showed a decreased ability to present antigens to T cells (125, 126), as did classical DCs, in which CGRP downregulated MHC-II and CD86 expression, resulting in decreased T-cell proliferation (127). Furthermore, CGRP treatment was found to suppress T_H1 differentiation in an in vitro DC - T cell co-culture model, and RAMP1-/- mice showed an enhanced T_H1 response in a model of delayed-type hypersensitivity (DTH) (128). Additionally, CGRP-pretreatment of antigen-pulsed bone marrow-derived DCs (BM-DCs) in vitro prior to transfer into naïve mice alleviated the allergic airway inflammation and expansion of allergen-specific T-cells after a subsequent antigenic challenge (129). The underlying mechanism remains unclear, however, significantly increased levels of IL-10 were observed in the tissue (129). Similarly, in the context of 2,4,6trinitrochlorobenzene-induced contact hypersensitivity (CHS), CGRP injected intradermally during the sensitization phase inhibited the migration of Langerin⁺ dermal DCs to lymph nodes by preventing upregulation of the chemokine receptor CCR7 (130). On the other hand, T-helper 2 (T_H2)-type responses induced by LCs pretreated with CGRP in vitro were enhanced (117). Finally, CGRP was reported to have chemo-attractant properties toward immature but not mature monocyte-derived DCs in vitro (131), and to alter the motility of airway mucosal DCs in living lung slices ex vivo (132).

In summary, CGRP appears to have the ability to modify functions of DCs in multiple ways, enhancing local immune response within barrier tissues, while downmodulating the DCs' ability to migrate to lymph nodes and to present antigens to T-cells. Many observations, especially pertaining to the latter, however, have only been made using exogenous administration of CGRP, and it remains unclear whether release of CGRP from nociceptors would be sufficient to induce comparable effects. AM and intermedin effects on DCs: In contrast to CGRP, much less is known about the effects of AM and intermedin on DCs. However, given the shared signal-transducing element of the CGRP and AM receptors, it is reasonable to speculate that their effects may be largely similar to those of CGRP. Indeed, *in vitro*, AM inhibited LPS-induced maturation of BM-DCs, but it also induced a "semimature tolerogenic" phenotype in unstimulated cells, characterized by intermediate upregulation of CD80, CD86, and indoleamine 2,3dioxygenase (IDO) expression (133). Of note, AM-treated BM-DCs were found to express AM themselves (133), however, the relevance of this apparent feed-forward loop remains unclear.

SP effects on DCs: The effects of SP on DCs are generally considered to be pro-inflammatory. In vitro, in GM-CSF-induced BM-DCs, SP prevented apoptosis upon withdrawal of GM-CSF through the PI3K-Akt signaling pathway (134), and enhanced Tcell proliferation in a DC – T cell coculture (135). Additionally, SPtreated in vitro generated DCs showed decreased IL-10 production and induced an enhanced T_H1 response when transferred in vivo (136). Local administration of a synthetic analog of SP during a gene-gun immunization also resulted in enhanced T_H1 and cytotoxic T-cell responses (137). Furthermore, pulmonary DCs displayed increased motility when exposed to SP in vitro and localized in proximity of SP⁺ nociceptive fibers in vivo. Ablation of nociceptors in the lung resulted in a decreased number of DCs and diminished infiltrates after pulmonary antigen challenge (115). Whether this effect was mediated by SP or other nociceptor-derived signals remains to be established. Finally, repeated stress exposureinduced SP accumulation enhanced LC migration out of the skin to the draining lymph-nodes, blocked production of the $T_{\rm H}2$ cytokines IL-4 and IL-5, and enhanced the levels of TNF α and IFN γ , alleviating allergic skin inflammation (138). Nevertheless, there is scant direct evidence for nociceptor-derived SP-mediated control of DC functions under physiological conditions. One recent study reported that the CD301b⁺ subset of skin DCs is activated by TRPV-1⁺ nociceptor-derived SP in an allergen challenge model (85). Such SP-activated DCs showed altered migratory properties and enhanced priming of allergen-specific T_H2 responses (85). Interestingly, in this study CD301b⁺ DCs recognized SP exclusively through the MRGPRA1 receptor and, in contrast to previous studies (134, 136), the authors were unable to detect expression of any SP receptors on other DC subsets in vivo (85). In light of these observations, regulation of the SP receptor expression in DCs emerges as an outstanding question important for our understanding of the interaction between nociceptors and DCs and their outcomes.

VIP and PACAP effects on DCs: The effects of VIP and PACAP on DCs are context-dependent. On one hand, in *in vitro* generated BM-DCs, VIP synergized with TNF α in inducing IL-12 and CD83 expression (139), and VIP/PACAP could also induce CD86 upregulation in immature BM-DCs and allow them to stimulate T-cell proliferation and differentiation into T_H2 effector cells (140, 141). On the other hand, LPS-activated BM-DCs treated with VIP/ PACAP showed impaired upregulation of CD80 and CD86, and decreased ability to stimulate T-cell responses (140). Additionally, BM-DCs that had been differentiated in the presence of VIP/ PACAP showed a tolerogenic phenotype, failed to upregulate CD80, CD86, and CD40, expressed high levels of IL-10, and induced the expansion of regulatory T-cells in vitro and after transfer in vivo (142). Similarly, in vitro treatment of antigenpulsed LCs with VIP prior to their adoptive transfer into previously immunized mice ameliorated LC-dependent DTH responses, possibly due to downregulation of IL-12 and IL-1β, and upregulation of IL-10 (143). Indeed, when administered exogenously in vivo, PACAP also suppressed the induction of CHS by modulating Langerhans cell functions (144). However, a more recent study has argued the opposite and showed that denervated mice exhibited an attenuated CHS response, which was improved by repeated intradermal injections of PACAP. Mechanistically, PACAP injections increased the number of dermal DCs that migrated to the draining lymph node and, in vitro, the neuropeptide induced upregulation of CCR7 and CXCR4 on immature BM-DCs and improved their ability to migrate toward CCL21 and CXCL12 (145). While it remains unclear whether these in vitro and in vivo phenomena are connected, the findings suggest that PACAP can control dynamics of DC migration.

Other modes of nociceptor-DC communication: TRPV1⁺ nociceptors have also been shown to be involved in anti-viral responses. Through a yet-to-be defined mechanism, activation of cutaneous nociceptors was sufficient to induce IL-27 expression by dermal CD301b⁺ cells - a heterogenous group of myeloid cells that includes cDC2s as well as monocyte-derived cells - which, in turn, induced expression of anti-viral peptides in keratinocytes. Indeed, skin explants from TRPV-1 deficient mice were more susceptible to HSV infection than those from WT controls (146). Furthermore, DCs isolated from skin-draining lymph nodes of HSV-infected nociceptor-deficient animals were unable to efficiently prime cognate T-cells. Importantly, addition of exogenous antigen rescued the phenotype, indicating that there was no inherent defect in the ability of DCs from nociceptor-depleted mice to activate T-cells but rather that nociceptors control the ability of DCs to acquire and/or process and present antigens to T cells (147).

In summary, nociceptors and nociceptor-derived neuropeptides have the potential to control the trafficking and functions of DCs in multiple ways and, in several cases, a single neuropeptide can exhibit both pro- and anti-inflammatory properties. Many of these observations, however, were only made in vitro or in the context of exogenously administered synthetic neuropeptides. Consequently, the extent to which they are relevant for the interaction between nociceptors and DCs under physiological settings often remains unclear. Nevertheless, a picture is beginning to emerge, which suggests that nociceptors may control DC functions in a context-dependent manner through a controlled release of distinct neuropeptides. In particular, the recent in vivo data argue in favor of a model in which nociceptors, depending on the type of stimulus encountered, can skew immune responses toward local T_H17-like (26, 37, 38, 122) or adaptive T_H2 type responses (85). Whether the diverging responses are mediated by different subsets of nociceptors or the same subset that can itself respond differently to distinct stimuli is currently unclear.

Finally, we note that DCs engage in physical interactions with nociceptors (26, 85), yet the effects of nociceptors that have been described to date are, almost universally, attributed to soluble

neuropeptides. Whether there is a role for the physical association of the two cell types *per se* beyond ensuring that DCs are exposed to high concentrations of the locally released neuropeptides remains to be established.

3.2 Macrophages

Macrophages comprise a heterogeneous population of tissueresident myeloid cells with complex ontogeny. In adults, depending on the tissue, macrophage subsets may have a variety of origins: some are derived from yolk sac progenitors and maintained by selfrenewal, while others arise from migratory monocytes or other bone marrow derived hematopoietic precursors (148). The immune functions of macrophages include phagocytosis and degradation of cellular debris and foreign objects as well as cytokine production, wound healing and, to a limited degree, antigen presentation (149). Additionally, macrophages have non-immune roles that contribute to the homeostatic functions of various organs including the brain (150), heart (151), lung (152), and liver (153). Macrophages exhibit significant plasticity and, based on phenotypic and functional criteria, are often categorized as M1 or M2 type cells (154). M1 macrophages are characterized by the expression of the inducible nitric oxide synthase (iNOS) as well as the costimulatory molecules CD80 and CD86, and they exhibit pro-inflammatory properties. Conversely, M2 macrophages express the mannose receptor, CD206, and are mostly associated with anti-inflammatory, tissue repair-promoting functions. While the M1/2 nomenclature is often too simplistic to accurately capture the variability observed in macrophages in vivo (155), it is, nevertheless, often used as a convenient shorthand. Similar to DCs, macrophages are known to associate with and be impacted by sensory (Figures 3-5) as well as other types of neurons (156) in a variety of tissues including the gut (157), eye (158), and skin (159).

CGRP effects on macrophages: The effect of nociceptive neuropeptides on macrophages is among the earliest recognized examples of nociceptor-immune cell communication. Indeed, CGRP-mediated inhibition of the ability of macrophages to produce H₂O₂ and to act as antigen-presenting cells in response to IFNy was first reported in the late 1980s (160). Subsequent in vitro studies have shown that CGRP decreased the expression of other cytokines, including IL-12 and IL-1β, and upregulated IL-10 (161), LIGHT, and SPHK1 through a CREB-dependent mechanism (75). Moreover, CGRP can regulate macrophage polarization in vitro (162, 163), by inhibiting LPS-induced degradation of I-KB and promoting IL-4-induced STAT6 phosphorylation, thereby favoring the acquisition of the M2 phenotype (163). Accordingly, in recent in vivo experiments CGRP promoted M2 accumulation in two models of post-operative tissue regeneration (162, 164). Additionally, in a β-glucan osteoinflammation model, CGRP released from NaV1.8⁺ nociceptors inhibited osteoclast-mediated bone resorption and decreased TNFa and IL-6 levels (57). Ex vivo, CGRP also decreased TNFa production by peritoneal macrophages after LPS stimulation, and CGRP-treated mice were protected from lethal endotoxemia after systemic LPS injection (60). In contrast to these anti-inflammatory actions and similar to its pleiotropic effects on

DCs (discussed above), CGRP can also elicit pro-inflammatory responses in macrophages, at least in some settings. For example, in the context of acute postoperative intestinal inflammation, endogenously released CGRP potentiated the expression of the proinflammatory cytokines IL-6 and IL-1 β by peritoneal macrophages (165). CGRP also enhanced phagocytic activity in cultured peritoneal macrophages through a cAMP-dependent mechanism (166) and improved their capacity to kill *Leishmania* parasites (167). Interestingly, LPS-activated RAW264.7 macrophage cells *in vitro* (168), as well as macrophages invading an injured nerve site *in vivo* (169), have been shown to produce CGRP themselves, suggesting a regulatory mechanism to promote tissue repair *via* an auto- or paracrine negative feedback loop.

AM and intermedin effects on macrophages: The effects of AM and intermedin on macrophages are not well understood, however, they appear similar to those of CGRP. AM has been shown to decrease the production of TNF α by macrophages *in vitro* (170) and of IL-6 and IL-8 by fallopian tube macrophages *in vivo* (171). Similarly, intermedin can skew macrophage differentiation in white adipose tissue toward the M2 phenotype (172). Conversely, in the NR838 macrophage cell line, AM enhanced the secretion of IL-1 β and IL-6, while also reducing the expression of TNF α (173). Just like CGRP, expression of AM has also been shown in the RAW264.7 macrophage cell line as well as in peritoneal macrophages (174) and in macrophages in atherosclerotic plaques (175).

SP effects on macrophages: SP has long been known to induce an oxidative burst in macrophages (176) and to enhance the production of nitric oxide (NO) (177), TNFa, IL-1β, and IL-6 after LPS stimulation in vitro (178). Additionally, SP also decreased the production of TGF β , a cytokine with anti-inflammatory properties (179). SP has been implicated in the mediation of immunological changes induced by stress. For example, in a murine model of sound stress, the percentage of SP⁺ and CGRP⁺ sensory neurons innervating skin was increased (180) and this was accompanied by an SP-dependent increase in MHC-II⁺ macrophage clusters and neurogenic inflammation (181). Likewise, in a model of cold-water stress, SP released from TRPV-1⁺ nerve fibers accumulated in the peritoneal cavity and enhanced IL-6 production by peritoneal macrophages (182). Conversely, in a model of spinal cord injury, intravenous injections of SP decreased the abundance of M1 and increased the abundance of CD206⁺ M2 macrophages at the site of the injury, resulting in a decreased accumulation of IL-6 and TNFa, and an increase in IL-10 (183). Similarly, in vitro, SP prevented IFNYinduced M1 differentiation and activated the PI3K/Akt/mTOR pathway, which promoted differentiation into M2-like tissue repair-promoting macrophages. After adoptive transfer, SPinduced M2 macrophages migrated to sites of tissue injury and improved functional recovery (184).

VIP and PACAP effects on macrophages: VIP and PACAP have been dubbed "macrophage deactivating factors" owing to their ability to inhibit the expression of iNOS (101), proinflammatory cytokines [TNF α (100), IL-6 (185), IL-12 (186)] and chemokines [CCL2, CCL3 (187), CCL4, CCL5, CXCL2 and CXCL8 (IL-8) (188)], and to increase the production of IL-10 (102) by macrophages in vitro. Furthermore, VIP and PACAP induced rapid shedding of CD14 - an LPS-coreceptor - thus dampening macrophage responses to bacterial endotoxin (189). Similar to their effect on DCs, VIP/PACAP can, paradoxically, induce upregulation of CD86 on immature macrophages and allow them to stimulate $T_{\rm H}^2$ -biased adaptive immune responses (190) even though they inhibit CD80 and CD86 upregulation in activated macrophages (191). In vivo, exogenous administration of PACAP decreased inflammatory responses by macrophages in CNS and ocular inflammation models, and promoted neuroprotection (192, 193). Conversely, in rat peritoneal macrophages in vitro, PACAP has been reported to enhance phagocytosis and production of superoxide anions (194) as well as their adherence and mobility (195), while VIP inhibited TGF β production (196). Interestingly, pre-treatment of macrophages with VIP and PACAP increased their resistance to HIV infection in a PKA and PKC-dependent manner (197, 198). Of note, while the in-vitro observations of immunosuppressive qualities of VIP and PACAP have largely been recapitulated in in vivo models in which exogenous neuropeptides were introduced, whether and to what extent nociceptors can and, indeed, do use these neuropeptides to impact macrophage functions in vivo is less clear.

TAFA4 effect on macrophages: Because TAFA4 is mostly expressed within the central nervous system (199), its potential role in the periphery had previously been largely overlooked. Recently, however, a $G\alpha_i$ -interacting protein (GINIP)-expressing subset of NaV1.8⁺ neurons was shown to express TAFA4 neuropeptide upon UV-irradiation (200). The nociceptor-derived TAFA4 promoted IL-10 expression by macrophages and, consequently, supported the survival of skin-resident antiinflammatory TIM4⁺ macrophages, resulting in reduced levels of pro-inflammatory cytokines and improved tissue repair (200).

Other modes of nociceptor-macrophage communication: While neuropeptide-dependent communication pathways have been at the forefront of neuroimmunology research in general, nonneuropeptide-dependent modes of communication between nociceptors and macrophages have been described. One such mechanism is the release of HMGB-1 from activated nociceptors. In models of sciatic nerve injury and arthritis, nociceptor-restricted HMGB-1 ablation resulted in decreased inflammation and ameliorated pathology (201). While the HMGB-1-responsive cells were not identified, involvement of macrophages, which express HMGB-1 receptors (202) and play important roles in the development of arthritis (203) appears likely. TRPV1⁺ nociceptors were also implicated in macrophage accumulation, activation, and ROS production in IMQ-induced psoriasiform skin inflammation. In this model, CGRP and SP were responsible for the nocifensive behavior, which was inhibited by treatment with neuropeptide antagonists, but had no effect on the inflammatory response (204), similar to the observations made for DCs in the same model (26). Additionally, in the context of peripheral nerve injury, DRG neurons have been shown to release exosomes containing miR-21. Following phagocytosis of such vesicles, miR-21 induced a pro-inflammatory phenotype in DRG-resident macrophages, characterized by enhanced expression of iNOS, TNFa, and IL-6, and downregulation of CD206 and Arginase

(205). Finally, signaling through Toll-like receptors (TLR) and myeloid differentiation factor 88 (MyD88) licenses nociceptors to secrete the chemokine CCL2, resulting in enhanced macrophage infiltration into the DRG (206). The role of such macrophage infiltration in neuropathic pain and neuroinflammation is an important topic of ongoing investigation and has recently been reviewed elsewhere (207).

In summary, nociceptors can utilize multiple neuropeptidedependent and -independent means to exert control over macrophages, and, *in vivo*, the macrophage response will likely result from a combination of these signals. It is interesting to note that while DCs and macrophages are developmentally and functionally related and their responses to nociceptive neuropeptides bear certain similarities, there appears to be a stark dichotomy in the type of control that nociceptors exhibit over these two cell types. In particular, currently available *in vivo* data suggests that in the case of DCs, nociceptors impact the type of proinflammatory immune response, as discussed in the previous section. This is in contrast to macrophages, where the effect of nociceptors appears to focus on controlling the balance between pro- and anti-inflammatory, tissue repair-promoting phenotypes.

3.3 Monocytes

Monocytes are circulating phagocytes derived from the common monocyte precursor (CMoP) that are broadly divided into two distinct subsets - classical (also known as inflammatory) and non-classical (a.k.a. patrolling) monocytes - which differ in their phenotype, function and migratory properties. Under inflammatory conditions, monocytes rapidly migrate into the affected tissues where they can undergo diverse differentiation pathways to acquire functional as well as transcriptional properties of DCs or macrophages. Under steady-state conditions, monocytes only emigrate into tissues in small numbers and either help repopulate local macrophage niches or remain in an undifferentiated state to fulfill homeostatic roles and serve as local monocyte reservoirs (208, 209). Interestingly, CGRP, SP, and VIP all exhibit chemotactic properties toward monocytes in vitro (210) indicating that at least some monocytes are equipped to sense sensory neuropeptides. Nevertheless, owing to their migratory nature and paucity in uninflamed tissues, direct effects of nociceptors on monocytes in vivo have not been studied and our current understanding is limited to in vitro effects of neuropeptides (Figures 3-5). Additionally, DCs and macrophages derived from monocytes are rarely differentiated from their bona fide counterparts during analyses and, consequently, whether the effects that nociceptors have on DCs and macrophages in vivo are also applicable to monocytederived cells remains to be established.

Neuropeptide effects on monocytes: Similar to macrophages, CGRP exerts anti-inflammatory effect on monocytes, including inhibition of proliferation, antigen presentation, upregulation of CD86 (but not CD80), and secretion of IL-12 p40, while at the same time enhancing production of IL-10 in response to *Staphylococcus aureus in vitro* (211). Similarly, human peripheral blood monocytes

treated with VIP showed a reduction in the production of TNF α (212) and IL-8 (213) with no effect observed on IL-10 production (212). By contrast, SP exerts primarily pro-inflammatory effects on monocytes. Indeed, several studies have noted that SP can induce the release of IL-1 β , TNF α , and IL-6 from monocytes (214, 215) and TNF α from monocyte-derived macrophages (216). A subsequent study, however, reported that human peripheral blood monocytes are unable to respond to SP alone but rather that the neuropeptide synergized with low doses of LPS and enhanced LPS-induced IL-6 expression (217). Consequently, the authors speculated that undetected low levels of LPS in tissue culture media could have been responsible for the proposed cytokine-inducing effects of SP and that SP, in fact, does not act on unstimulated monocytes (217). The controversy has not been fully resolved to date.

In summary, the *in vitro* effects of neuropeptides on monocytes appear similar to their effects on other cells of the mononuclear phagocyte system; however, the absence of *in vivo* data makes it difficult to assess the pathophysiological relevance of this putative communication pathway. One intriguing observation in that context is the reported chemoattractant activity of neuropeptides toward monocytes, which could suggest the possibility of nociceptors controlling monocyte migration and/or localization in tissues.

3.4 Polymorphonuclear granulocytes

Polymorphonuclear granulocytes derive from the shared granulocyte-monocyte progenitor (GMP) in the bone marrow and are characterized by the presence of granules in their cytoplasm, which contain a variety of biologically active molecules released upon cellular activation. Granulocytes are often considered the first line of immune defense due to their rapid recruitment to tissues in response to inflammatory stimuli and their potent anti-microbial functions (12).

3.4.1 Eosinophils

Eosinophils play an important role in the pathology of allergic and parasitic diseases as key effectors of T_H2 -type inflammation (218). They respond to cytokines such as IL-5 and IL-13 by proliferating and releasing a variety of cytokines (IL-13, IL-4, IL-25, TNF α , GM-CSF), leukotrienes (C4, D4), prostaglandins, and the contents of their cytotoxic granules, particularly, enzymatic and nonenzymatic cationic proteins, as well as reactive oxygen species (ROS) (218–220).

Although eosinophils also populate the intestinal (221) and uterine mucosa at homeostasis (219, 222), they have been mostly studied in the context of their recruitment in response to allergens and parasites, especially in the airways and the skin. They localize close to sensory as well as parasympathetic neurons both in animal models of asthma and in biopsied lungs of asthma patients (223). In prurigo nodularis, a chronic skin disease characterized by nodules and intense itch, eosinophils are closely associated with CGRP⁺ nociceptive fibers (224) and in atopic dermatitis patients eosinophils are more closely associated with nerve fibers than in healthy controls (225). Finally, although the identity of the neurons was not determined in this context, close nerve-eosinophil contact was also observed in the colon and Peyer's patches of rats infected with the parasite *Fasciola hepatica* (226). One way eosinophils are postulated to interact with nociceptors is through the CCR3-Eotaxin 1 (CCL11) chemokine pathway. In addition to Eotaxin-1, VCAM-1 expression was also shown on DRG neurons in the presence of nerve growth factor (NGF) *in vitro* (227), which could support adhesion of eosinophils through VLA-4 (integrin $\alpha 4\beta$ 1) (228).

Neuropeptide effects on eosinophils: Like other myeloid leukocytes, eosinophils are also responsive to neuropeptides (Figures 3, 4, 6) such as CGRP, SP, and VIP, each of which have been shown to decrease IL-16 production by eosinophils in vitro (229). When isolated and cultured with SP, Guinea pig eosinophils released eosinophil peroxidase (EPO) (230), a cytotoxic protein contained within "specific" cationic granules (231). The C-terminal fragment of SP also triggers additional eosinophil responses in vitro, including the release of another component of the specific granules, the eosinophil cationic protein (ECP) (232), generation of superoxide (232), and inhibition of apoptosis (233). Additionally, equine eosinophils were shown to respond to SP by increased adherence, migration, and superoxide production, although the authors noted that only relatively high SP concentrations were able to elicit these effects (234). SP is also considered an eosinophil chemoattractant (233, 235-237) and intradermal injection of the neuropeptide led to eosinophil recruitment in human volunteers, while CGRP and VIP injections did not (235). Similarly, intranasal SP administration following allergen exposure led to increased eosinophil recruitment in patients with allergic rhinitis (238). Finally, in rat trachea venules, administration of SP increased the numbers of adherent eosinophils in an NK1R-dependent manner (236). However, it is unclear in these experiments whether SP acted directly on eosinophils or rather on endothelial cells to upregulate adhesion molecules [for a review see (239)].

In vitro studies with eosinophils from allergic individuals suggest that both CGRP and SP can enhance eosinophil migration toward other chemoattractants, such as leukotriene B4 (237), platelet-activating factor (237, 240), or IL-5 (240). CGRP by itself can induce human eosinophil chemotaxis – directed migration along a concentration gradient, while VIP stimulates chemokinesis

- random, directionless motility - by signaling *via* VPAC1 but not VPAC2 (241).

Only a few studies have addressed the effects of nociceptors on eosinophil recruitment and functions in vivo: TRPA1-KO mice, which exhibited lower levels of SP, CGRP, and neurokinin A had decreased levels of IL-13, Eotaxin-1, CCL2, RANTES, and IL-5 after an allergic challenge. This phenotype was correlated with a IL-5 decrease in the numbers of eosinophils in the bronchioalveolar lavage fluid (BALF) (242). Similar findings were made in a model of airway inflammation in mice genetically lacking all NaV1.8-expressing nociceptors and mice that were administered QX-314 - a charged derivative of lidocaine and a potent sodium channel blocker, which silences the electrical activity of nociceptors. These mice had fewer eosinophils in their airways and lower levels of IL-5, IL-4, Eotaxin, and TNFa in the BALF (27). Reciprocally, capsaicin treatment, which induces the activation of TRPV1⁺ nociceptors, led to increased eosinophil recruitment (27). Similarly, in a model of allergic asthma, ablation of TRPV1⁺ neurons prevented development of airway hyperreactivity and bronchoconstriction phenotypes (243), which are known to be in part mediated by eosinophil major basic protein (MBP) (219, 244). Notably, unlike the observations made in mice lacking NaV1.8⁺ nociceptors (27), no overall changes in the immune cell infiltrate were observed in mice devoid of TRPV1⁺ nociceptors (243). Conceivably, these disparate results might be due to different populations of nociceptors being targeted (NaV1.8⁺ vs TRPV1⁺) or differences in the experimental models themselves. Such differences notwithstanding, a TRPA1 antagonist has shown efficacy in preclinical models of asthma, and has recently entered phase 1 human clinical trials (245), providing an important proof of concept for targeting peripheral neuroimmune interactions in clinical settings.

Taken together, nociceptive neuropeptides have the ability to enhance recruitment and chemotaxis of eosinophils under inflammatory settings and SP in particular can induce eosinophil degranulation and ROS generation. In light of the fact that many neuropeptides can be produced by cells other than nociceptors, the extent to which neuropeptides released specifically by activated nociceptors impact eosinophil functions *in vivo*, however, remains largely unclear.

3.4.2 Basophils

Basophils, which account for ~1% of the blood leukocytes, are recruited to sites of $T_{\rm H}2$ -mediated inflammation and are often



regarded as the "blood mast cells" (246, 247). It is becoming increasingly clear, however, that basophils have non-redundant functions, especially in their response to haptens and peptide antigens (248). *In vitro*, nociceptor neuropeptides secretoneurin and SP, signaling through NK1R, showed chemoattractant properties comparable to the N-formyl peptide, fMLP, – a known granulocyte chemoattractant – and LPS (249). In mice, intraperitoneal injection of SP led to a marked increase in blood basophil numbers (250). In addition, basophils from patients with chronic spontaneous urticaria had higher expression of NK1R and of SP itself (250). Importantly, the threshold for SP-induced histamine release from such basophils was decreased compared to basophils from healthy subjects (250).

In summary, while it appears that SP enhances the proinflammatory functions of basophils (Figures 3, 4, 6) similarly to what has been described for the other granulocytes (251), our understanding of the neuronal control of their functions is in its infancy.

3.4.3 Neutrophils

Neutrophils are short-lived, sentinel immune cells that comprise 50-70% of all circulating leukocytes in humans and 10-25% in mice (252). As the first responders to pathogen entry, these myeloid cells are equipped with a plethora of effector functions aimed at eliminating pathogens. Specifically, neutrophils phagocytose bacteria, release proteases and oxidants, form neutrophil extracellular traps (NETs), and communicate with other immune cells through cytokine release (253). Importantly, neutrophil activation also has the potential to cause significant collateral damage to the host tissues and, as such, their functions are tightly regulated. Nociceptors have been reported to form close contacts with neutrophils that have emigrated into tissues (28), and nociceptive neuropeptides in particular have been shown to profoundly influence neutrophil migration and functions both *in vitro* and *in vivo* (Figures 3, 4, 6).

Multiple studies have utilized various depletion or activation techniques to test the impact of nociceptors on neutrophils in vivo. These studies usually focus on whether neutrophil recruitment and activity are affected by assessing the expression of adhesion molecules on blood neutrophils, determining neutrophil numbers in the tissue, as well as evaluating the activity of whole tissue myeloperoxidase (MPO), a peroxide-degrading enzyme released during neutrophil degranulation. Although often presented as such, whole tissue MPO activity as a proxy of neutrophil-mediated effector functions has important limitations as it does not account for the production of MPO, albeit at lower levels, by cells other than neutrophils (254). Only a few studies used ex-vivo assays to measure neutrophil-specific MPO and direct antimicrobial properties by coincubating neutrophils with bacteria in the presence of relevant neuropeptides or nociceptors. Overall, an immunosuppressive effect of nociceptors on neutrophils has been observed. Notably, in rats, electrical stimulation at intensities that activate noxious C fiber afferents led to a reduced accumulation of neutrophils in a model of bradykinin-induced knee joint inflammation (255). While total blood neutrophil numbers remained similar, a marked decrease in L-selectin (CD62L) positive neutrophils in the nociceptor-activated

group was reported. Additionally, in a laminar flow assay using blood neutrophils isolated from rats after noxious stimulation, a significant reduction in the number of rolling and tethering cells was observed (255), suggesting that nociceptive fiber stimulation might downmodulate the expression of neutrophil adhesion molecules (256) in parallel with its effects on endothelial cells (239). Building on these original observations and using complementary loss-of-function models, further studies have arrived at similar conclusions: In an LPS-induced subacute airway inflammation model, depletion of TRPV1⁺ nociceptors resulted in a higher lung MPO activity and IL-1 β secretion (257). Likewise, in a Staphylococcus aureus mouse model of lethal pneumonia, neutrophil recruitment and functions were suppressed by TRPV1⁺ nociceptors, as mice lacking TRPV1⁺ fibers exhibited a higher percentage of neutrophils in the lungs 6h post infection, a lower bacterial burden and better survival (25). Accordingly, an increase in crawling of neutrophils in the subpleural vascular bed was observed using intravital microscopy in the TRPV1⁺ nociceptor-ablated mice (25).

Further studies have also explored the role of nociceptorneutrophil communication within the gastrointestinal tract in dextran sodium sulfate (DSS) and trinitrobenzene sulfonic acid (TNBS) induced colitis models (258–261). Strikingly, ablation of TRPV1⁺ fibers (258), as well as TRPV1 antagonism (259, 260) resulted in a lower disease score, accompanied by lower tissue MPO activity (258–260). Conversely, however, another study has reported that TRPV1 agonism through daily administration of low doses of capsaicin during DSS colitis in WT rats led to a lower tissue MPO activity (261). While it is possible that such treatment led to the desensitization of the TRPV1⁺ fibers (262), these results indicate that the effects of nociceptors on neutrophils might not be straightforward.

CGRP effects on neutrophils: CGRP acting through the RAMP1-Calcrl receptor is known to have an immunosuppressive effect on neutrophils (25, 263-266). In vitro, CGRP prevented LPSinduced upregulation of CD11b on human neutrophils (263) while, in vivo, mice lacking RAMP1 exhibited increased infiltration of CD11b^{Hi} neutrophils into the peritoneal cavity and improved antibacterial defense in the early stage of septic peritonitis (265). After myocardial infarction, TRPV1-KO mice showed increased number of recruited neutrophils and higher pro-inflammatory cytokine concentrations (IL-6, $TNF\alpha$) in the heart, which correlated with lower levels of CGRP and were reversed by exogenous CGRP administration (266). Similarly, in a mouse model of Streptococcus pyogenes skin infection, TRPV1⁺ nociceptor-derived CGRP suppressed neutrophil recruitment and functions (58). More recently, analogous observations have also been made in a model of urinary tract infection with uropathogenic E.coli, where nociceptor depletion as well as direct CGRP antagonism improved recruitment and functions of neutrophils to the urinary bladder, and resulted in improved bacteria clearance (267). Also in vitro, DRG neurons decreased clearance of S. pyogenes by bone marrow neutrophils, as did CGRP treatment alone, albeit to a lower degree, suggesting a possible involvement of other nociceptor-derived factors - possibly AM or Intermedin - in suppressing neutrophil antimicrobial activity in vivo. Finally, the

MPO activity of neutrophils incubated with *S. pyogenes* was also decreased by CGRP in a concentration-dependent manner (58).

SP effects on neutrophils: Consistent with the effects of SP on other myeloid cells, it also amplifies the pro-inflammatory activities of neutrophils. For example, intravenous injection of SP led to increased neutrophil adhesion in rat trachea venules in an NK1R dependent manner (236). Similarly, after intradermal injection of SP in human volunteers, increased numbers of neutrophils in the lumen of dermal venules and the interstitium were observed (235). This phenomenon, however, is likely due to the effects of SP on endothelial cells, which are known to respond to SP by transcriptional upregulation of E-selectin and translocation of Pselectin from Weibel-Palade bodies to the cell surface (235), both adhesion molecules that are critical for the recruitment of bloodborne neutrophils. Nonetheless, direct effects of SP on neutrophils have also been reported, albeit only in vitro (232, 268-270). Indeed, SP, through NK1R signaling, enhances neutrophil survival by inhibiting caspase 3-mediated apoptosis (269) and induces superoxide generation (232, 268). Additionally, incubation of human neutrophils with SP resulted in phosphoinositide hydrolysis, an increase in intracellular calcium, activation of NADPH oxidase, and production of reactive oxygen species as well as enhanced exocytosis of azurophilic and specific granules after cytochalasin B treatment (268). Finally, SP has been shown to enhance fMLP-mediated production of arachidonic acid metabolites LTB4 and 5-HETE by human neutrophils, as well as to increase antibody-dependent cellular cytotoxicity (271).

VIP and PACAP effects on neutrophils: The neuropeptides of the VIP/PACAP family seem to have an immunosuppressive effect on neutrophils. In a model of LPS-induced septic shock, VIP/PACAP administration led to lower levels of liver and intestinal MPO activity in a PAC1-dependent manner (272). Furthermore, intratracheal administration of VIP and PACAP analogs prior to IL-1 β treatment resulted in a decreased neutrophil infiltrate in the BALF (273), and intraperitoneal administration of recombinant VIP decreased MPO activity in *Aspergillus fumigatus*-infected cornea (274).

In summary, the *in vivo* studies imply that the principal effect of nociceptor activation is attenuation of neutrophil-mediated

inflammation, at least in the limited number of experimental systems employed to date. Indeed, neutrophil exposure to several neuropeptides that can be released by nociceptors, particularly CGRP and VIP/PACAP, results in suppression of effector activity. However, at least one nociceptor-derived neuropeptide, SP, can exert potent pro-inflammatory nociceptor-derived activity on neutrophils. While this has only been described *in vitro* and evidence for such effect *in vivo* is currently lacking, it nevertheless hints at an unappreciated degree of complexity in the interaction between nociceptors and neutrophils.

3.5 Mast cells

Mast cells (MCs) densely populate barrier tissues and often localize within close proximity of nerve endings (275) and are well known to engage in bi-directional communication with nociceptors (276, 277). MCs play a central role in allergic reactions where, following IgE-mediated crosslinking of Fcc receptor I (FccRI), they rapidly exocytose storage granules containing mediators such as heparin, histamine, proteases, and cytokines. Granule exocytosis initiates further downstream signaling events that lead to vascular leakage, recruitment of various immune cells, and other components of allergic inflammation (278). Additionally, certain mast cell mediators, in particular histamine and serotonin are potent pruritogens and activate nociceptive fibers that transmit the sensation of itch (279). Neuropeptide-mediated activation of MCs (Figures 3, 4, 7) is independent of IgE/FcR signaling and results in spatially and temporally distinct patterns of degranulation (280) and release of unique combinations of mediators. Specifically, upon IgE-induced activation, MCs secrete larger and more heterogeneously shaped granules that are able to drain to lymph nodes and influence adaptive immune responses (281) whereas MC activation by neuropeptides induces rapid secretion of small secretory granules that are either not transported to or fail to be retained in draining lymph (280). Additionally, neuropeptide stimulation leads to higher production of diverse proinflammatory chemokines, including CCL2 (MCP-1), CCL5



Effects of select nociceptive neuropeptides on mast cells observed *in vitro*. Upward pointing arrows signify upregulation/activation, downward facing arrows signify downregulation/inhibition.

(RANTES), CXCL10 (IP-10), and CXCL8 (IL-8) (282) but limited release of prostaglandin E2 and VEGF (280). These distinct patterns of degranulation and mediator release, potentially along with other, yet to be described molecular features of neuropeptide-mediated MC activation, lead to a more rapid development of vascular leakage but more transient local inflammatory reactions in comparison to anti-IgE-induced activation (280).

MCs in different tissues display marked differences in their phenotype and function, including their neuropeptide receptor expression, granule composition, and cytokine production (283). Mucosal MCs (MMCs) are predominantly found in the gut and contain granules that consist of chondroitin sulfate and either tryptase (in human) or chymase only (in mouse). In contrast, connective tissue MCs (CTMCs) are found in the skin, intestinal submucosa, myocardium (284), nasal epithelium (285), and peritoneum and contain granules that mainly consist of heparin proteoglycans and both tryptase and chymase. Notably, these phenotypic differences have been utilized to generate mouse lines that allow for selective deletion and functional perturbation of either MMCs or CTMCs (286, 287). Although CTMCs and MMCs are ontogenetically distinct (288), they can be induced by the tissue microenvironment to take on phenotypic and functional features of the other subtype (289, 290). Thus, in light of the growing evidence of tissue microenvironment playing a role in regulating MC-nociceptor communication, we will discuss the effect of nociceptor activation on MCs according to their tissue localization:

Nociceptor interactions with CTMCs: SP appears to be the main nociceptive mediator that activates CTMCs, resulting in the release of histamine, proteases, and various chemokines that lead to further innate immune cell recruitment. Although there is some evidence for NK1R expression on human nasal mucosal MCs (291), rat cardiac MCs (292), and dermal MCs within eczematic skin lesions (293), MRGPRB2/X2 appears to be the main receptor for SP in murine and human CTMCs (84). Among the CTMCs, dermal MCs are perhaps the best-characterized population. In the context of tissue damage, SP signals through MRGPRB2/X2 to activate dermal MCs, resulting in their degranulation and release of proinflammatory cytokines (TNFa, GM-CSF) and chemokines (CXCL8, CCL2, 3, and 4) leading to neutrophil recruitment to the site of injury and development of pain hypersensitivity (83). Indeed, injection of staphylococcal enterotoxin B in combination with Dermatophagoides farinae extract (house dust mite allergen) activates TRPV1⁺ nociceptors to the release of SP, leading to MC degranulation, cytokine production, influx of eosinophils and neutrophils, and $T_{H}2$ -type skin inflammation (82).

Interestingly, unlike activation through the canonical IgE/FccRI pathway, SP/MRGPRB2-mediated degranulation results in the release of more tryptase and less histamine and, subsequently, excites the non-histaminergic itch sensory neurons (294). The mechanistic regulation and long-term functional consequence of favoring tryptase release remain to be characterized. Nevertheless, given the ability of tryptase to degrade substrates such as cytokines and neuropeptides (295), it is tempting to speculate that favoring tryptase over histamine could allow for a more controlled pattern of immune activation and rapid return to tissue homeostasis.

In addition to neurogenic inflammation, MRGPRB2/X2mediated MC activation also facilitates protection against pathogens. In mice lacking functional MRGPRB2, neutrophil recruitment to and clearance of subcutaneously inoculated S. pyogenes were impaired (285). Similarly, in a model of S. aureus skin infection, MRGPRB2-mediated CTMC degranulation resulted in the release of TNF α , GM-CSF, CXCL8, CCL2, and CCL3 and subsequent recruitment of bacteria-clearing neutrophils and wound healing-promoting CD301b⁺ DCs (296). Instead of SP, however, mastoparan - a different MRGPRB2 agonist - was used to induce CTMC activation. Therefore, whether S. aureus infection can physiologically result in the release of SP sufficient to induce CTMC activation in a similar fashion remains to be directly demonstrated. Lastly, in addition to mediating degranulation, SP can also modulate TLR2 expression on MCs (297), which may alter the MC response to subsequent bacterial exposure.

Interestingly, a recent study has shown that a non-peptidergic subset of MrgprD⁺ (Mas-related G-protein-coupled receptor D)-expressing cutaneous sensory afferents modulates gene expression in CTMCs by the release of glutamate (298). Notably, MRGPRB2 was among the downregulated genes, and activation of MrgprD⁺ neurons, which are themselves maintained by skin-resident Langerhans cells, was sufficient to suppress dermal MC responsiveness. Conversely, depletion of the MrgprD⁺ neurons resulted in increased susceptibility to MrgprD⁺ neurons may be responsible for setting an overall tone of CTMC responses in the skin.

In comparison to the skin, less is known about the effects of nociceptive neuropeptides on CTMCs residing in other tissues. In vitro studies have demonstrated that SP induces IL-6, TNFa (299) and histamine release from rat peritoneal MCs (300) and histamine from cardiac MCs (292), as well as serotonin and TNF α release from murine peritoneal MCs without a concomitant release of histamine (301, 302). In vivo, intraperitoneal inoculation of vancomycin-resistant E. faecium into mice with dysfunctional MRGPRB2 resulted in increased bacterial loads compared to controls (285). Based on what has been described for dermal MCs, loss of SP-mediated MC-activation and subsequent defect in the recruitment of bacteria-clearing neutrophils could be the mechanistic underpinning of this phenotype, however, direct experimental evidence is currently lacking. Similarly, murine nasal epithelial MCs also express MRGPRB2, and its dysfunction during nasopharyngeal infection with S. pneumoniae resulted in decreased TNFa levels in the nasal lavage fluid (NLF), decreased recruitment of neutrophils and MCs to the nasopharynx, and impaired bacterial clearance compared to control animals (285). Interestingly, in the human nasal mucosa, MCs express NK1R and NK2R but not MRGPRX2 and the functional effect of SP has not been elucidated (291). Overall, more work is necessary to better understand the effect of SP-MRGPRB/X2-mediated CTMC activation within different tissue microenvironments, especially given that SP-MC signaling is implicated in both pathological neurogenic inflammation as well as protection against pathogens.

Molecular mechanisms and functional consequences of CTMC activation by neuropeptides other than SP are poorly understood.

CGRP has been shown to induce degranulation and release of serotonin (301) and histamine (303) in peritoneal and dermal MCs and, in a model of contact hypersensitivity (CHS), intradermal injection of CGRP induced release of TNFa from MCs, which resulted in reduced LC density and, consequently, suppressed the CHS phenotype (304). Thus, contrary to other myeloid cells, the effect of CGRP on MCs appears to be preferentially pro- rather than anti-inflammatory. Similar to CGRP, AM was able to induce MC degranulation and histamine release as well as upregulation of VEGF and MCP-1, and increased MC motility in vitro (305). However, MC activation by nociceptor-derived AM has not been demonstrated in vivo. Interestingly, MCs are a prominent component of solid tumors and, in the tumor microenvironment, can be readily activated by tumor cell-derived AM, resulting in IL-17A release and enhanced tumor growth (306). Lastly, VIP has been shown to induce degranulation and histamine release in rat peritoneal MCs (300) but, paradoxically, suppresses stressmediated MC degranulation in the testes (307).

Nociceptor interactions with MMCs: Compared to CTMCs, much less is known about neuropeptide-mediated control of MMCs (308). Unlike CTMCs, MMCs do not express MRGPRB/X2 (84) and the expression of NK1R has long been controversial (309). Recent work in human and guinea pigs, however, has demonstrated that NK1R is expressed on intestinal MCs and mediates SP-induced degranulation and release of histamine and protease II (310). In addition to SP, rodent, but not human (309) MMCs also respond to CGRP and VIP by degranulation (311). Interestingly, in murine models of food allergy (312) and intestinal parasitic infection with Nipppostrongylus brasiliensis and Schistosoma mansoni (313), a significant increase in the density of CGRP+ fibers and closely associated MCs was observed in the intestinal mucosa. Similarly, patients with irritable bowel syndrome have an increased number of degranulated MCs localized in close proximity to nerve fibers, and an increase in mucosal tryptase concentrations correlates with the severity and frequency of abdominal pain/discomfort (314). Overall, these observations suggest that the neuropeptide-MC signaling axis plays important, albeit poorly understood roles in the biology of MMCs and their functions in inflammation and protection against pathogens within the intestinal mucosa. Lastly, in a model of SP-induced cystitis in the urinary bladder, MMCmediated tissue edema and neutrophil infiltration were not dependent on NK1R signaling (315), suggesting that bladder MMCs might express another, yet to be identified SP receptor.

In contrast to other tissues, the lung is uniquely populated by both MMCs and CTMCs, with MMCs being the prevailing type in bronchi, bronchioles, and alveolar parenchyma and CTMCs along the pulmonary vessels and pleura (316). Unlike their murine counterparts (84), human lung MCs do not express MRGPRX2 and, consequently, do not degranulate in response to SP at steadystate (317). However, both the number of MRGPRX2⁺ MCs and overall levels of MRGPRX2 expression were increased in lung biopsies from patients who had died from asthma (318), suggesting that MRGPRX2 may be upregulated in MCs within the inflamed airway microenvironment and potentially play a functional role in airway hyperresponsiveness. Indeed, SP concentration was also elevated in the BAL and sputum of asthmatic patients (319) and SP can induce degranulation of MCs isolated from human BAL (320). Nevertheless, further work is necessary to directly implicate SP-MRGPRB/X2 in the progression of asthma as well as to explore the effect of other neuropeptides on lung MC functions.

Finally, non-barrier, non-mucosal tissues, including the CNS, also contain MCs that are highly responsive to neuropeptides. The dural meninges are densely populated by MCs, which have been reported to form close contacts with nociceptors (321) and respond to SP, VIP, and CGRP by degranulation and histamine release (322). Nevertheless, most in vivo work has shown CGRP to be the predominant neuropeptide that mediates nociceptor-MC signaling under pathological conditions, including migraine (323) and postconcussion pain (324). Indeed, epidural infusion of CGRP in rats is sufficient to induce MC degranulation (325). Unlike the rat MCs, however, human meningeal MCs do not appear to express the CGRP receptor (326), and it is unclear whether they can respond to CGRP at all. MCs are also found within specific areas in the underlying brain parenchyma, such as the corpus striatum. Upon systemic VIP administration, these MCs become activated but do not show classic degranulation patterns (327), and their physiological role remains a mystery.

Overall, work thus far has demonstrated important functions of neuropeptide-mediated MC activation in the context of neurogenic inflammation and protection against pathogens in barrier and nonbarrier tissues. However, heterogeneity between MC populations present in various tissues remains a confounding factor and a more detailed understanding of how the nociceptor-MC signaling axis functions in tissue-specific immune surveillance is needed. In particular, understanding the function of nociceptor-MC communication in response to pathogens is of outstanding interest, given the MCs' strategic localization at the hostenvironment interface, and a growing body of work demonstrating MCs to play critical roles in protection against infection (285, 328).

4 Discussion

Although the concept of neuroimmunology in peripheral tissues is not new, progress in our understanding of the operational paradigms that dictate the functional outcomes of such interactions has long been hindered by limitations in the availability of experimental tools and our insufficient knowledge of both the immune and nervous systems *per se.* However, recent progress in multiple areas of scientific inquiry and a convergence of these hitherto separate areas have allowed at least some of these interactions to come into focus. While the field in many ways is still in its infancy, it appears likely that significant headway will be made in years to come.

It is often tempting to over-simplify the effects of nociceptors and neuropeptides on immune cells and categorize them as either pro- or anti-inflammatory. However, we must take into account the interconnectedness and context-dependency that the actions of both the immune and nervous system exhibit. A revealing case in point is the effect of nociceptor-derived CGRP on neutrophil-

mediated immune responses. When directly applied, CGRP appears to have potent anti-inflammatory properties, inhibiting neutrophil migration as well as their effector functions. However, through its actions on dendritic cells, CGRP can also enhance the local T_H17 response, resulting indirectly in an increased production and influx of neutrophils and an enhanced inflammatory response. Teleologically, it is interesting to ponder why the immediate effects of CGRP would prevent neutrophil recruitment if, ultimately, the neuropeptide orchestrates a chain of events that leads to the opposite result. This might be perceived as particularly puzzling given that pain (and concurrent release of neuropeptides) often accompanies barrier breaches, where an increased risk of pathogen invasion should favor the evolution of mechanisms that promote a preemptive recruitment of bacteria-clearing neutrophils. Nevertheless, it is important to take into account the potentially destructive effects that neutrophils can have in inflamed tissues. Indeed, if the threat from a wound is minimal, curtailing of neutrophil accumulation and activation could prevent further tissue damage and favor tissue repair and faster return to homeostasis. On the other hand, larger or infected wounds are likely to cause sustained activation of local immune cells such as the DCs and macrophages alongside the release of CGRP, which together signal a need for a strong immune response, likely overriding the effect that CGRP alone might have on neutrophils. Thus, nociceptors appear to have the ability to fine-tune the immune response in tissues, further enhancing it where needed,

is not. While it is increasingly clear that nociceptors have the ability to change the parameters of immune responses through the secretion of neuropeptides, the regulation of the processes remains enigmatic. In particular, multiple neuropeptides that often have opposing effects are known to be expressed in the same nociceptive fibers. Consequently, it appears likely that neuropeptide release patterns in vivo should be tightly controlled, similarly to what has been described for molecular communication in other branches of the nervous system (329, 330). To that end, it is conceivable that different modalities of activation could result in a spatially and/or temporally distinct pattern of release of distinct neuropeptides from the same fiber. Similar observation have, in fact, been made in rat vagal sensory neuron cultures, in which release of different ratios of CGRP and SP was observed depending on the challenge modality (331). At the same time, at a single-cell/fiber level, nociceptors are known to be non-uniform in their expression of neuropeptides (332, 333). Consequently, it is conceivable that subsets of nociceptors that are distinct in their specificity for various stimuli or their physiological location might inherently show biased patterns of neuropeptide expression to achieve a similar effect. Clearly, further work is needed to establish if and how differential neuropeptide release in vivo is controlled and how it impacts immune responses. Additionally, while volume transmission (diffusion-driven effect at a distance) is an accepted mechanism of neuropeptide-dependent communication (334), the scale at which neuropeptide gradients form and the distances at which they can act on various immune cells within tissues remain poorly defined.

while also preventing an excessive activation in situations where it

It is of note that several feed-forward loops exist in which immune cells exposed to nociceptive neuropeptides are prompted to express the neuropeptides themselves (133, 168, 169, 174, 175). While the functional relevance of such phenomenon has not been investigated, it appears to be an example of a signal-amplifying feed-forward loop. Historically, neuropeptides were thought to be synthesized exclusively in the cell bodies of nociceptors and transported through axons within vesicles to the peripheral nerve endings where they get released (335). More recent studies have, however, demonstrated intra-axonal synthesis of at least some neuropeptides (336), and it remains unclear which of the mechanisms is more important within nociceptors' axons at the steady state (337). Nonetheless, in either scenario, it appears likely that the amount of neuropeptides readily available in the termini of nociceptors is limited, as the transport along axons has been argued to take hours to days (338), and the scale of local proteosynthesis inside axons is inherently limited by the comparatively small volume of cytoplasm. On the other hand, immune cells have the ability to infiltrate tissues in large quantities, ramp up protein production when activated, and to move within the tissue, perhaps providing a raison d'être for such a feed-forward amplifier. Additionally, tissue damage will inevitably result in physical disruption of local nociceptor fibers and hence a possible transient loss of local neuronal neuropeptide sources. It is thus conceivable that myeloid cells present in the affected tissue could provide a substitute source until nociceptor innervation is restored. Experimental evidence in the form of immune cells genetically deficient in the neuropeptides of interest will be necessary to test these ideas.

As discussed throughout this review, to this day, with only a few exceptions, neuropeptide release has widely been considered the main, if not the only, mechanism by which nociceptors exert control over immune cells. Nevertheless, other modalities of interaction could exist, e.g. interactions of surface molecules, direct coupling through gap junctions or tunneling nanotubes, production of secretory vesicles, or release of non-neuropeptide mediators and neurotransmitters. Investigation of such alternative modes of communication could provide worthwhile insights into the nature of the neuroimmune interactions and shed more light on the intricate network of communication in which nociceptors and immune cells engage. Finally, while here we focused specifically on the means by which nociceptors impact on the functions of myeloid immune cells, it is important to stress that the communication between nociceptors and the immune system is bidirectional. Indeed, many of the effects that immune modulators have on nociceptor functions are known and have been reviewed recently (339, 340).

In summary, while a growing body of literature pertaining to the interaction between nociceptors and myeloid cells exists, many outstanding questions remain. In particular, a number of studies investigated the effects of nociceptive neuropeptides on immune cells in isolation or after injecting a neuropeptide into animals, while others have explored the effect of global nociceptor ablation. Although such studies have generated important insights, we are largely lacking a mechanistic understanding of how nociceptorderived neuropeptides act at the single cell or tissue level. New experimental strategies, such as inducible nociceptor-specific ablation or induction of neuropeptide expression and/or immune cell-specific ablation of neuropeptide receptors could potentially address these questions in the near future. Furthermore, while the interactions between nociceptors and certain cell-types, such as macrophages and mast cells have received much attention, others, such as monocytes or basophils remain largely unexplored.

Author contributions

PH, M-AM, and YW wrote sections of the manuscript, PH and UHvA generated and edited the final version of the manuscript. All authors contributed to the article and approved the submitted version.

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Conflict of interest

UHvA is a paid consultant with financial interest of Avenge Bio, Beam Therapeutics, Bluesphere Bio, Curon, DNAlite, Gate Biosciences, Gentibio, Intergalactic, intrECate Biotherapeutics, Interon, Mallinckrodt Pharmaceuticals, Moderna, Monopteros Biotherapeutics, Morphic Therapeutics, Rubius, Selecta and SQZ.

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