

The Role of Neuropeptide Endopeptidases in Cutaneous Immunity

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Synonyms Box: Neprilysin, neutral endopeptidases, common acute lymphoblast leukemia antigen (CALLA) CD10, enkephalinase, CD10, NEP, EC 3.4.24.11;

Key Features

- ▶ Proteolytic processing and degradation plays an important role in modulating the generation and bioactivity of neuroendocrine peptide mediators, a class of key molecules in cutaneous biology.
- ▶ Accordingly, the cellular localization and expression, and the molecular biology and structural properties of selected intracellular prohormone convertases and ectopically expressed zinc-binding metalloendopeptidases are discussed.
- ▶ A special reference will be made to the physiologic and pathophysiologic significance of these endopeptidases in cutaneous immunobiology.
- ▶ Because of the number of pathologically relevant changes in inflammation and tumor progression that can be directly attributed to neprilysin and angiotensin-converting enzyme, a particular focus will be on the role of these enzymes in modulating innate and adaptive immune responses in the skin.

angiotensin-converting enzyme, dipeptidyl carboxypeptidase, kininase II, ACE, CD143, EC 3.4.15.1; Dipeptidyl peptidase IV, CD26, EC 3.4.14.5

Abbreviations: ACE Angiotensin-converting enzyme, ACTH Adrenocorticotropin, Ag Antigen, Ang Angiotensin, BK Bradykinin, CGRP Calcitonin gene-related peptide, CTCL Cutaneous T-cell lymphomas, DC Dendritic cell(s), DPPIV Dipeptidyl peptidase IV, EAE Experimental autoimmune encephalomyelitis, ECE Endothelin-converting enzyme, EC Endothelial cells, END Endorphin, MC_x Melanocortin receptor, MHC

Major histocompatibility complex, *MSH* Melanocyte-stimulating hormone, *NEP* Neprilysin, *PACAP* Pituitary adenylate-cyclase-activating polypeptide, *PC* Prohormone convertase, *POMC* Proopiomelanocortin, *SP* Substance P, *TC* T-cell(s), *T_H* helper T-cells, *T_{eff}* Effector T-cells, *VIP* Vasoactive intestinal peptide

8.1 Introduction

Almost every aspect of cutaneous cellular and tissue function, including proliferation, differentiation, maturation, communication, antigen (Ag) presentation, and survival of cells as well as hair growths, eccrine gland function, wound healing, and tissue regeneration, is modulated by neuropeptides. It is thus quite comprehensible that a variety of mechanisms have evolved, which limit their temporal, spatial, and developmental bioactivity. These include temporally and spatially controlled mediator generation and release, the regulated expression of specific receptors on cellular targets, receptor desensitization and resensitization, and the clearance of excessive extracellular peptides. Proteases participate in several of the above mechanisms; thus taking a key regulatory role in cutaneous peptide mediator bioavailability. As such they serve to generate bioactive peptides from inactive prohormones in order to initialize inflammatory and trophic responses. Importantly, they also rapidly terminate the bioactivity of neuropeptides released from nerve terminals or endocrine cells and thus prevent the development of a neuropeptide-initialized or -augmented deleterious chronic inflammation. In addition, microbial invaders or parasites use peptidases as an evolutionarily successful strategy to manipulate the host immune defenses. Peptidases play an important role in cutaneous plasticity and wound healing by modulating trophic neuropeptide activities. Moreover, beyond a mere catalytic function, ectopeptidases trigger specific intracellular signal transduction, participate in cell–cell or cell–virus recognition, and mediate or modulate binding to extracellular matrix components. This chapter highlights some of the current knowledge on peptidase function in cutaneous immunity and outlines clinical and potential future research areas derived from key functions of these enzymes. In addition, as zinc metalloproteases are among the largest group of

proteases relevant for the extracellular cleavage of neuroendocrine mediators, a special emphasis will be made on this class of proteases.

8.2 Intracellular Endoproteases Convert Inactive Prohormones to Bioactive Mediators

8.2.1 Cellular Localization and Expression

Despite the important role that intracellular endoproteolytic processing and activation of prohormones, particularly of proopiomelanocortin (POMC), by prohormone convertase (PC) plays for cutaneous physiologic and pathophysiologic responses, this chapter's focus centers on functions of extracellular proteases. Importantly, with respect to cutaneous melanocortin generation, some extracellular proteases may also be capable of processing larger precursors, resulting in bioactive melanocortin receptor-(MC-) activating POMC peptides. Neuroendocrine hormones such as adrenocorticotrophin (ACTH) or α -melanocyte-stimulating hormone (α -MSH) are released in the skin as part of an intrinsic cutaneous hypothalamus–pituitary–adrenal axis and mediate the cutaneous response to invasive and noninvasive exogenous stress via MC receptors [11,87]. PC are an evolutionary conserved class of secretory serine proteases of the subtilisin/kexin-type that comprise PC1/3, PC2, furin/PACE, PACE4, PC4, PC5/6, PC7, and SKI-1 [81]. POMC peptides and POMC processing enzymes including PC1, PC2, PACE4, or furin have been identified in a variety of skin cells, skin appendages, cutaneous carcinoma cells, and immune cells (reviewed in [10,42,81]).

8.2.2 Molecular Biology and Structural Properties

The conserved structure of the PC catalytic domain suggests that these proteases have evolved from a common ancestral precursor gene. Subtilisin/kexin family PC are specialized for cleaving multiple hormones, growth factors, and receptor precursors by limited internal proteolysis at single or multiple basic recognition sites, within the general motive K/R-(X)_n-K/R↓ [81]. PC expression and activity is strictly regulated at the tissue, cell, or subcellular level.

Autocatalytic activation of PC zymogens is another means to control PC bioactivity within the secretory pathway. In some cases, that is, for PC2, cofactors such as the binding protein 7B2 are required for efficient zymogen activation of proPC and full functional activity [50,81].

8.2.3 Physiologic Significance

The primary function of cutaneous PC is the conversion of the POMC prohormone in various skin cells. The resulting generation of α -MSH, ACTH, or β -endorphin is highly relevant for skin immunity, stress response, and pigmentation (for more details see Chap. 6 and [42]). Studies of PC1/3- and PC2-deficient mice revealed that deletion of these enzymes impairs the processing of POMC and other prohormones, although some redundancies might exist [71]. The pathophysiologic consequences of a dysregulated PC expression for cutaneous immunity have not yet been fully explored. The simultaneous episodic expression of PC1, PC2, and POMC during the murine hair cycle suggests a regulatory function of PC for the pilosebaceous unit [49]. POMC, MC, and PC expression in some skin cells are synergistically regulated by UV light, melanocortins, and pro-inflammatory cytokines in vitro. These stimuli may simultaneously increase production and responsiveness of cutaneous cells to POMC peptides, although this has not been conclusively confirmed in vivo [73,76]. An upregulated PC1, PC2, and furin expression positively correlates with malignant neuroendocrine tumors and of several other cancers [39], suggesting that subtilisin/kex-like convertases may increase tumorigenesis and aggressiveness by augmenting processing and activation of mitogenic peptides. Thus, PC both serve as a prognostic marker for tumor progression and constitute an important pharmacologic target in cancer therapy, since PC inhibition drastically reduced the metastatic properties of certain tumor cells [39,71]. Interestingly, some bacterial toxin precursors (e.g., *Diphtheria* toxin, *Botulinum* neurotoxin), as well as viral glycoproteins of HIV-1, Ebola, and others viruses need proteolytic PC activation for their toxic or infectious capacity and/or the cell–cell spreading. This demonstrates the high relevance of PC for the cutaneous response to infectious agents and, therefore, PC inhibition could be beneficial for abrogating microbial-induced cytopathicity (reviewed in [39]).

8.3 Dipeptidylpeptidase IV/CD26

8.3.1 Cellular Localization and Expression

Dipeptidyl peptidase (DP) IV (CD26, EC 3.4.14.5) is a multifunctional homodimeric glycoprotein with functional roles in hematology, endocrinology, immunology, endothelial cell (EC), and cancer biology and metabolism. DPIV is part of a six member gene family of enzymes that, in addition to DPIV, includes fibroblast activating protein (FAP), DP-like (DPL) 1, DPL2, DP8, DP9, and prolyloligopeptidase (POP) [13,27]. Human DPIV is ubiquitously expressed by capillary EC, activated lymphocytes, DC subpopulations, and on apical surfaces of epithelial cells [27]. In addition, soluble forms of the enzyme have been described. Cutaneous DPIV is expressed on keratinocytes [60], fibroblasts [59], melanocytes [55], the axon–Schwann cell interface [20], and TCs [41].

8.3.2 Molecular Biology and Structural Properties

The structural and biochemical properties of DPIV have been described in detail in [27]. The DPIV gene product is a 766 amino acid (AA) ectoprotease with an apparent monomeric molecular weight of about 110kDa. Characteristically, full functional DPIV peptidase activity requires homodimerization between one of two extracellular hydrolase domains. This results in a rather unique post-proline dipeptidyl aminopeptidase activity of DPIV by cutting off N-terminal X-P or X-A dipeptides from polypeptides. A variety of DPIV peptide substrates have functional relevance for skin (patho) physiology. These comprise at least 9 CCL and CXC chemokines (i.e., CCL5, RANTES, or CXCL10, IFN γ -induced protein), hormones (i.e., glucagon-like peptides (GLP), prolactin, leutinizing hormone α , chorionic gonadotropin α chain), enkephalins, and neuropeptides such as neuropeptide Y, pituitary-adenylate cyclase-activating polypeptide (PACAP) 38, vasoactive intestinal peptide (VIP), and SP.

8.3.3 Functional Roles of DPIV/CD26 in Immunity and Inflammation

There is compelling evidence that DPIV has a number of important physiologic functions in endocrinology and metabolism. For instance, DPIV degrades GLP and glucose-dependent insulinotropic peptide. The

inhibition of DPIV results in accumulation of these peptides, which stimulates greater insulin production and is therefore beneficial for the treatment of insulin-independent *Diabetes mellitus* (see [27] for details). DPIV/CD26 has gained considerable interest as a T cell (TC) activation marker, and is also expressed by some dendritic cell (DC) subpopulations [24]. Accordingly, DPIV expression, together with other TC activation markers such as CD25, CD71, CD45RO, or CD29, increases significantly after antigenic and mitogenic stimulation, or treatment with the T helper (T_H)1 cytokine IL-12. Overexpression of human CD26 in TC of transgenic mice reduces thymus cellularity, impairs thymocyte proliferation, and increases the number of peripheral apoptotic CD4⁻ or CD8⁻TC, indicating the importance of CD26 for peripheral T lymphocyte homeostasis [85]. Interestingly, DPIV upregulation increases degradation of VIP and PACAP, two neuropeptides known to trigger T_H 2 immune responses via IL-4, IL-5, and IL-10 induction in CD4⁺ TC [25,27]. Up- or downregulated CD26 expression may therefore shift the T_H 1/ T_H 2 balance [72,82] with high relevance for psoriasis, atopic dermatitis [5], or rheumatoid arthritis (RA) [8]. However, studies of murine experimental autoimmune encephalomyelitis (EAE) and RA revealed surprising discrepancies between functional inhibition of DPIV by genetic knock-out and pharmacologic inhibitors. While DPIV-inhibiting drugs delayed the onset and severity of experimental EAE or RA, missing CD26 activity in knock-out mice or in human patients was inversely correlated to the severity of Ag-induced RA or EAE. Thus, DPIV inhibitors may have additional functional targets [8]. CD26 expressed in lipid rafts within the immunologic synapse also transduces intracellular signals that overlap with the TC receptor/CD3 signaling pathways. DPIV enzyme activity may therefore be dispensable for full immunologic activity of TC, since costimulatory activity of CD26 in vitro is retained in DPIV mutants that lack hydrolase activity [27].

8.3.4 DPIV/CD26 and the Development of Neoplasms

The skin is the host for a number of extranodal non-Hodgkin cutaneous TC lymphomas (CTCL) [41]. Strikingly, in the most relevant types of CTCL, mycosis fungoides (MF) and Sézary syndrome (SS), skin-homing malignant CD3⁺ CD4⁺ CD7^{variable}CLA⁺ CCR4⁺ TCs characteristically lack CD26 expression. DPIV/CD26 is an important diagnostic marker for SS or MF, and

also highly relevant for the pathophysiology of CTCL [88]. Missing CD26 compromises the TC capability to degrade the constitutively expressed cutaneous stem cell factor 1, which may promote homing of malignant TC into the skin [58]. Infiltrating CTCL-CD26⁻ TC then generate an immunosuppressive T_H 2 environment with immature DC incapable of efficiently presenting phagocytosed material derived from apoptotic malignant TC. The resulting regulatory TC (T_{reg}) then contribute to immunosuppression and CTCL-TC tolerance [41]. Evidently, DPIV may also be relevant for growth, invasiveness, and metastasis of other tumors. For instance, DPIV expression is inversely correlated to progression of melanoma and even absent in metastatic melanoma [67,101]. Conversely, overexpression of DPIV in melanoma cells suppresses tumor progression in nude mice possibly independently of a DPIV enzymatic activity. Likewise, DPIV interacts with ECM components such as collagen, fibronectin, E-cadherin, or tissue inhibitors of matrix metalloproteases [103]. Consequently, a higher DPIV expression enhances adherence of tumor cells to the ECM, which may be anti-invasive by preventing detachment of tumor cells from the solid tumor. Alternatively, DPIV may also hamper basic fibroblast growth factor (bFGF) mitogenic signaling pathways [106]. In summary, DPIV/CD26 (patho-)physiologic effects in tumorigenesis possibly depend on a direct interaction with ECM components or an indirect degradation of important inflammatory mediators.

8.4 Ectopic Zinc Metalloendopeptidases: Nephilysin and Angiotensin-Converting Enzyme

8.4.1 Cellular Localization and Expression

Nephilysin (NEP, EC 3.4.24.11, CD10) and angiotensin-converting enzyme ACE (EC 3.4.15.1, CD143), two mechanically related EC surface zinc metallopeptidases, are widely distributed in the body and highly expressed in the vascular endothelium, kidney epithelium, lung, or CNS [99,100]. NEP (“enkephalinase” or “common acute lymphoblastic leukemia antigen” (CALLA)) was initially isolated more than 30 years ago as an insulin B chain-degrading enzyme abundantly expressed in the renal brush border membrane [100]. The history of the ACE family has accomplished a journey from the original discovery and isolation of ACE 50 years ago as “hypertension-converting

enzyme” until the recent identification of the related carboxypeptidase ACE2 as vaso-peptidase and coreceptor for the severe acute respiratory syndrome corona virus (SARS-CoV) [30,31]. The somatic isoform of ACE and ACE2 are abundantly expressed in the vascular endothelium surface of the lung and in brush-border membranes of kidney, intestine, placenta, and the choroid plexus. A soluble isoform of somatic ACE is present in the plasma, and a smaller isoenzyme essential for male fertility is expressed in the testis [30,99]. Cutaneous NEP as well as components of the renin-angiotensin system (RAS) including ACE and ACE2 are expressed in basal keratinocytes, hair follicles, eccrine and sebaceous glands, the microvascular endothelium, and large nerves or nerve-ensheathing Schwann cells (reviewed in [31,75,89]). NEP and ACE, but not ACE2, are expressed in DC, macrophages, or TC and in bone marrow stromal cells, suggesting functional roles in hematopoiesis and immunity [15,31,38,45,47]. A recently identified secreted homologue NEP2 related to *D. melanogaster* NEP is expressed in the renal tube and in testis, but its relevance for skin immunity still has to be determined [96].

8.4.2 Molecular Biology and Structural Properties

Despite structural similarities between ACE and NEP, NEP is evolutionarily closer related to the bacterial Zn protease thermolysin. NEP homologues were identified in all organisms from simple prokaryotes to higher vertebrates, including men [100]. The mammalian NEP family now comprises at least seven members, with NEP and endothelin-converting enzyme (ECE, EC 3.4.24.71) as best characterized. The molecular biology of NEP is described in [69]. As a type II integral membrane protein ectopeptidase, NEP (90–110 kDa) consists of a short N-terminal intracellular domain, a transmembrane anchor, and a large C-terminal extracellular domain that contains a Zn-coordinating active site constituted by a HExxH and an ExxA/GD sequence [100]. Two closely related ACE isoenzymes (somatic and germinal) have been identified in mammalian cells. Somatic ACE (150–180 kDa) is a type I C-terminally membrane-anchored glycoprotein that contains two highly homologous extracellular domains (N-domain, C-domain), each bearing a zinc-coordinating catalytic site. By contrast, the smaller testicular ACE involved in male fertility contains a single catalytic site identical to the C-terminal domain of somatic ACE [44,99].

8.4.3 Physiologic Roles of NEP and ACE

Much help in understanding NEP and ACE function is derived from using mercaptoalkanoyl inhibitors such as captopril [99], or selective NEP inhibitors such as thiorphan and phosphoramidon [69]. The latter is a *streptomyces tanashiensis* product suggesting an evolutionary old relationship between zinc metalloproteases-expressing prokaryotes and eukaryotes [100]. In addition to its widespread role in turning off neuronal signals transmitted via SP or enkephalins, the panel of today’s known NEP substrates includes, but is not limited to, vasoactive peptides such as bradykinin (BK) or angiotensin (Ang) I, atrial natriuretic peptide (ANP), growth factors such as bombesin, chemotactic peptides such as fMLP, and most recently β -amyloid (A β) peptide, the key initiator of Alzheimer’s disease (AD) [35,86,100]. The structure of the mammalian NEP extracellular domain limits accessibility of the catalytic site for substrates. Therefore, and in contrast to the protease activity of thermolysin, NEP is an oligopeptidase cleaving peptides predominantly non-terminal before hydrophobic AA residues [69]. ACE’s active sites display endopeptidase and dipeptidyl carboxypeptidase activity, which differ in their pH/chloride dependency and substrate specificity [36,105]. A prominent physiological feature of NEP and ACE is their overlap in competitively cleaved substrates, resulting in opposing roles in renal and cardiovascular regulation (reviewed in [9,86]). The successful 30 year use of ACE inhibitors to therapeutically intervene with hypertension and cardiovascular dysfunction has encouraged attempts to additionally inhibit related enzymes such as NEP and recently ECE with a single drug. Advantage of such dual ACE/NEP- or triple ACE/NEP/ECE-specific “vasopeptidase inhibitors” is a limited generation of proinflammatory, blood pressure-rising vasoconstrictors (Ang II, ET-1), the accumulation of vasodilators (BK, Ang₁₋₇), and additionally of the diuretic ANP. Consequently, vaso-peptidase inhibition lowers blood pressure, diminishes cardiac hypertrophy and fibrosis, promotes renal natriuresis and diuresis, but also bears the risk of serious adverse effects [3,9,91].

8.4.4 Zinc Metalloproteases Terminate “Danger Signals”

NEP and ACE are ancient components of the innate and adaptive immune response of higher vertebrates. First, they participate in the immediate host defense – unfortunately with initial advances for a microbial intruder. Bacterial thermolysin-like peptidases facilitate

entry into the host by degrading peptides with antimicrobial properties, for instance nerve-derived host SP, but also α -MSH or adrenomedullin [93] [Fig. 8.1(1)]. In parallel, peptidases enhance the bacterial invasiveness by degrading ECM (collagen IV) and facilitate the entry into the host's circulation [Fig. 8.1(2,3)] [51,52]. The complex peptidase–substrate interplay is reflected by the fact that – from the defendant's point of view – limited degradation of vasoactive and neuropeptides may be advantageous in early inflammation. Neuronal and cellular-derived SP, CGRP, or BK induce a vicious cycle of releasing IL-1 or TNF α from cutaneous cells that conversely promote the release and axonal transport of SP and CGRP in sensory neurons ([65,68,74,75], and included references.). Indeed, a downregulated NEP expression by irritants or endotoxins as observed in respiratory tract or intestinal inflammation suggests that prolonged activity of SP and BK may be advantageous for kick-off and progression of (neurogenic)

inflammation and subsequent events, such as a full Ag-specific inflammatory response at the site of Ag challenge.

Numerous studies using vasopeptidase inhibitors or NEP-/ACE-deficient mice confirm the significance of NEP and ACE for cutaneous innate immunity [65,68,74,75]. The lack of NEP increased inflammatory responses and lethality in various murine inflammatory models (reviewed in [75]), and functional deletion of NEP and/or ACE markedly exacerbated murine allergic contact dermatitis (ACD) [23,77,78]. Missing NEP or ACE particularly on, or in the vicinity of, hematopoietic and immune cells modulates hematopoiesis [38], profoundly disturbs the local immune cell-activating cyto- and chemokine microenvironment, and imbalances pro- and anti-inflammatory neuropeptides. The latter fine-tune the DC:TC interface and T_H differentiation in adaptive immunity. Initialization of cutaneous delayed-type hypersensitivity requires activation and migration

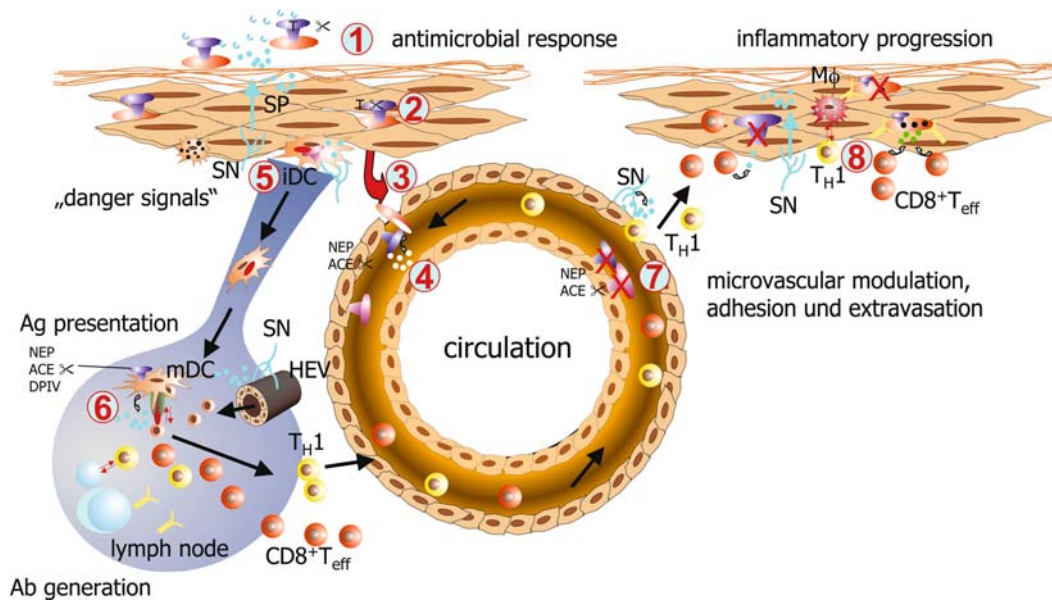


Fig. 8.1 Endopeptidases control multiple steps in inflammation and initialization of adaptive immunity. Intruding bacterial pathogens utilize thermolysin-like peptidases (T) to partly degrade antimicrobial neuropeptides, for example, substance P (SP) derived from sensory neurons (SN) or keratinocytes (1). By degrading extracellular matrix, peptidases facilitate microbial entry into the host (2) and accession to the vascular lumen (3). In some cases, host peptidases (NEP) may liberate anti-inflammatory peptides (α -MSH) from pathogen precursor molecules to compromise host's immunity (4). Released SP and other pro-inflammatory peptides serve as “danger signals” to activate residing immature dendritic cells (iDC) and induce DC maturation and migration into draining lymph node (LN) (5). In the

LN, neuropeptides (SP) released from sensory nerves contacting high endothelial venules (HEV) or from matured DC (mDC) interacting with naïve T-cells (TC) promote T-helper (T_H) 1 polarization and clonal expansion of T_H1 and CD8+ effector TC (CD8+ T_{eff}), particularly in the absence of NEP or ACE (6). Inflammation-downregulated peptidases at the site of ongoing inflammation promote vascular responses to released neuropeptides, as well as the recruitment and extravasation of inflammatory cells (7). In environment lacking functional NEP, prolonged activity of pro-inflammatory neuropeptides released from SN, macrophages (M ϕ), or TC promote multiple leukocyte effector functions, and in parallel, the temporarily increased availability of growth factors facilitates recovery of the damaged tissue (8)

[Fig. 8.1(5)] of Ag-laden dermal DC or Langerhans cells and contact to naïve CD4⁺ TC in skin-draining lymph nodes [Fig. 8.1(6)]. Vasoactive peptides such as SP, BK, and Ang II via NK₁, B₂, and AT1/AT2 receptors, respectively, profoundly modulate myeloid DC precursor differentiation [38] and boost important DC and TC functions (Fig. 8.2) [45]. APC (DC or macrophages) and TC express NK₁ and B₂ as well as RAS components (ACE, AT1/AT2 receptors) and constitute extra-neuronal sources of SP [2,17,45,46,56]. Importantly, SP and BK activate DC NF-κB, a central transcription factor involved in DC maturation marker (CD11c, MHC) and

costimulatory molecule (CD40, B7 molecules) expression, and thus NEP/ACE-deficiency may promote functional maturation of CD11c⁺ DC [2,45]. Vice versa, antagonist blocking of B₂ or NK₁ receptors impairs Ag sensitization in mice lacking functional ACE or NEP, which could be mimicked by adoptive transfer of NK₁ receptor antagonist-treated, Ag-pulsed wild type bone-marrow DC generated in vitro [23,77–79]. Exogenous or endogenous SP contributes to mitogen- or Ag-induced TC proliferation and promotes a T_H1 response characterized by reduced IL-4/IL-5 and increased IL-2/IFN γ expression [45,74,75,79]. Hence, prolonged bioavailability

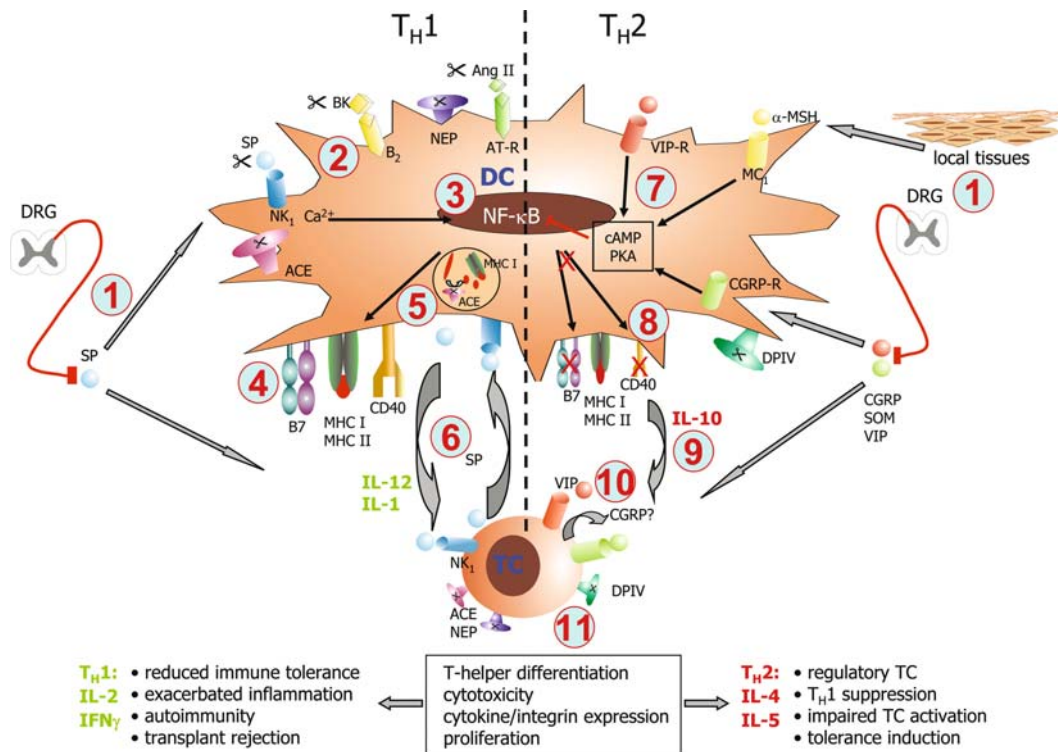


Fig. 8.2 Endopeptidases control the neuroendocrine hormone-balanced immune response transmitted by the immunologic synapse. Neuroendocrine and vasoactive peptides are released from cutaneous sensory neurons or cutaneous or lymphoid tissues (1). Ectopic (endo-) peptidase (NEP, ACE) mediator degradation controls the cellular accessibility of substance P (SP), bradykinin (BK), and angiotensin (Ang) II and activation of specific neurokinin (NK₁), bradykinin (B₂), and Ang (AT) receptors expressed by dendritic cells (DC), (2). SP, BK, and Ang II constitute endogenous danger signals that stimulate DC pro-inflammatory signal transduction pathways such as the release of Ca²⁺ and subsequent activation of NF-κB (3). NF-κB plays a central role in the induction of DC costimulatory B7 molecules, CD40 or major histocompatibility complex (MHC) expression, and pro-inflammatory T-helper 1 (T_H1) cytokine release (4). Endopeptidases (e.g., ACE) improve the efficacy of antigen presentation to T-cells (TC) by

intra- or extracellular trimming of peptides for MHC class I or II presentation (5). Endogenous SP from DC and TC may in an auto/paracrine manner drive T_H1 polarization under control of DC- or TC-expressed NEP and ACE (6). Via specific receptors expressed by DC and TC, α-MSH, calcitonin-gene related peptide (CGRP), somatostatin (SOM), and VIP trigger intracellular cAMP/protein kinase A (PKA) signal transduction (7). This inhibits endotoxin, cytokine-, or SP/BK-induced NF-κB activation and may promote a tolerogenic DC phenotype characterized by reduced TC costimulation (8), and release of anti-inflammatory cytokines (IL-10) (9). As demonstrated for VIP, such DC drive T_H2 polarization, regulatory TC (T_H2_{reg}) development, and suppression of T_H1 immune responses. T_H2-inducing neuropeptides such as VIP (10) under the control of TC-expressed dipeptidyl peptidase IV/CD26 (DPIV, (11)) may also directly trigger T_H2_{reg} development from naïve CD4⁺ CD25⁺ TC, which dampens T_H1 and CD8⁺ T_{eff} responses

of “proinflammatory” tachykinins and kinins constitutes an endogenous “danger signal” per se that generate fully matured MHC class II/I expressing DC, which drive a T_H1 polarization and efficiently prime T_H and effector cell (T_{eff}) responses (Fig. 8.2). This may be beneficial for fighting intracellular pathogens or malignant cells, but could also result in uncontrolled inflammation or even organ-specific autoimmunity. In contrast, “ T_H2 ” neuropeptides such as VIP/PACAP, CGRP, or α -MSH oppose the above by impairing Ag-presentation, and generating tolerogenic DC that attenuate TC activation. In addition, this may cause a T_H2 differentiation and give rise to regulatory CD4 and IL-10-producing CD8⁺ CD28⁻ CTLA4⁺ TC that subsequently suppress Ag-specific T_H1 mediated immune responses [25]. However, nature has predetermined additional roles for ACE and related peptidases for APC function in adaptive immunity. DC ACE is an important chaperone in the processing and MHC I-restricted presentation of viral and other peptides to CD8⁺ cytotoxic T lymphocytes (CTL) [83,108]. ACE interacts with the transporter associated with Ag presentation (TAP) complex that shuttles cytosolic peptides into the exocytic compartment for association with nascent MHC I molecules. Carboxy-terminal ACE trimming of certain peptides improves their intracellular transport, and thus rescue their otherwise inefficient presentation to CTL [108]. Cathepsins, or a recently described asparaginyl endopeptidase, play a similar role in the endocytotic lysosomal pathway of DC and B-cells, with tremendous impact on MHC II-restricted Ag presentation [104]. Importantly, glycosylation or deamidation of certain self-proteins prevents cleavage by these proteases. Thus, a loss of these initially present post-translational modifications over time may suddenly render a self-protein susceptible for APC processing and recognition by TC, with fatal consequences for the maintenance of tolerance against self-antigens [19,104]. Thus, expression and specificity of endopeptidases in the MHC I or II compartment is essential for the control of peptide antigenicity.

8.4.5 Regulated NEP and ACE: A Protective Role Against Ultraviolet Irradiation?

Recent observations nurtured the hypothesis that NEP indirectly may have UV-protecting properties. UVB irradiation has been demonstrated to downregulate the NEP activity in human melanocytes. Pharmacologic NEP inhibition increased the efficacy of α -MSH or ACTH to induce tyrosinase activity and microphthalmia

expression, suggesting that NEP plays a role in melanogenesis [1]. However, the role of UV in regulating NEP and ACE expression may depend on the cellular and functional context. UV light or cytokine-exposed microvascular EC displayed a time-dependent loss of cell surface ACE in vitro, whereas the initially low NEP expression increased [75]. This complex reciprocal regulatory system of vasoepitidases, which has also been observed in hypertensive rats [37], may be of particular significance for EC function. Recent bio-analytical studies demonstrated that endothelial NEP and ACE are highly relevant for processing stress (ACTH) and anti-inflammatory hormone (α -MSH) [42]. However, instead of a mere removal of extracellular POMC peptides, NEP and ACE peptidolysis generated bioactive novel MC_x agonists or antagonists distinct from the parental peptide. This phenomenon may have currently undefined functional roles for vascular biology and cutaneous inflammatory responses, since EC are an established source and target of POMC peptides. Of note, invertebrate parasites use vasoepitidases to generate anti-inflammatory immunosuppressive melanocortins from POMC precursors, suggesting an evolutionary conserved mechanism to compromise a host’s immune system [42].

8.4.6 Role of NEP and ACE in Cutaneous Wound Healing and Plasticity

It is now widely appreciated that NEP expression down-regulated by irritants, pro-inflammatory cytokines, endotoxins, or phorbol ester, respectively, increases the cellular accessibility of mediators and may promote inflammation and support cell growths as well as the development of neoplasms [34,95]. For instance, in nonhealing skin of *diabetes mellitus* patients or of mutant diabetic *db/db* mice, NEP expression and activity are increased ([75] and included references.). SP and CGRP may contribute to wound healing, as they promote keratinocyte, endothelial cell, and fibroblast proliferation and migration in vitro. Like bFGF, they also enhance angiogenesis and neovascularization in vivo (reviewed in [6,74]). Thus, increased NEP compromises re-epithelialization and wound healing, potentially by degrading growth factors such as bFGF-2 [26] and SP, which may facilitate development of diabetic ulcers [62]. Despite an improvement of extracutaneous wound healing after concomitantly applied SP and NEP inhibitors [7], these agents may exacerbate neuroinflammation and contribute to hypertrophic scar

formation [80]. However, in pathologic cutaneous scar tissue, an increased expression of ACE in comparison to normal or wounded skin, particularly by myofibroblasts, was detected. Potentially, by generating abundant Ang II, ACE may have profibrotic and, thus, adverse effects on cutaneous wound healing similar to remodeling in the heart after myocardial infarct [54,92].

8.4.7 Development of Neoplasms: Shutting Off Growth-Promoting Signals is the Key

Allergens or irritants such as cigarette smoke impair NEP expression and activity in the lung, which may increase respiratory distress and neurogenic inflammation. Moreover, reduced NEP activity in certain forms of prostate or lung carcinoma prevents hydrolysis of mitogens such as bombesins, ET-1, or SP, which may promote peptide-driven tumor growths [84,95]. Likewise, reduced NEP activity hampers generation of the growth inhibitory fragment SP₁₋₇ from NEP cleavage of SP [22]. One of the discussed mechanisms is an indirect phosphorylation and activation of the insulin growth factor 1 receptor (IGF-1R) that, further downstream, activates protein kinase B (Akt) anti-apoptotic cell survival pathways [95]. In line with this hypothesis, T and B cells become CD10-positive when undergoing apoptosis [16,53]. Hence, NEP expression represents a safety device that protects cells from mediators released from apoptotic cells and assures apoptotic cell-death by preventing access of cell survival-assuring mitogens. Interestingly, an apparent positive correlation of higher NEP levels with the malignancy of melanoma have been observed [4,102]. This increased NEP expression was accompanied by downregulated anti-apoptotic and upregulated pro-apoptotic proteins (Bcl-2, and Bax, respectively) [4]. Thus, NEP serves as a marker for apoptosis and progression of melanoma. However, the divergent expression pattern of NEP in metastatic melanoma, in contrast to other cancers, suggests that NEP may have diverse and tumor-specific modulating properties that require detailed future analysis. Noteworthy in this context, not all effects of vaso-peptidases can be attributed to changes in extracellular mediator levels. Very recently, an ACE inhibitor-induced dimerization, phosphorylation, and signaling of ACE has been demonstrated in EC that enhanced endothelial ACE expression [43]. Likewise, as demonstrated for ACE, NEP, and tachykinin or BK receptors, the cellular co-expression [61] or a sterically close receptor—peptidase association is important for receptor function and resensitization [18,21,48].

Therefore, direct vaso-peptidase outside-in signaling or modulation of G protein-coupled receptor (GPCR) signal transduction independent of ligand cleavage may account for some effects of NEP or ACE inhibitors that require future analysis.

8.4.8 NEP and ACE: Are There Functional Roles in Psoriasis, Alopecia Areata, and Acne?

Innervation of the pilosebaceous unit plays a pivotal role for sebocyte function and hair growth. Stress-induced SP promotes both proliferation and differentiation of sebaceous glands, resulting in an increased size and lipogenesis. Surprisingly, in skin of acne patients in contrast to normal skin, the subcellular expression of SP-induced NEP is increased in germinal sebocytes [97]. This may argue against a stress-induced inflammation, although the exact role of increased NEP awaits future investigation. Likewise, in acute and late stage chronic alopecia areata, NEP is strongly expressed in follicular structures of the affected area, whereas the perivascular expression of ACE in alopecia areata lesions is diminished [98]. This may prevent the local generation of keratinocyte/fibroblast proliferation-inducing Ang II [90]. As SP is capable of inducing significant anagen hair growth, limited bioavailability of SP, Ang II, and growth factors in alopecia areata may attenuate hair growth and increase hair follicle regression and apoptosis [64,65]. Uncontrolled cellular access to sensory neuropeptides and growth factors may also contribute to the pathogenesis of psoriasis. A reduced NEP expression in acute psoriatic lesions, but not in healthy skin, may locally increase SP levels that contribute to keratinocyte hyperproliferation, inflammation, and psoriatic pruritus [12,57,75]. In conjunction with a reportedly enhanced neuronal density and NK₁ expression, this may explain, in part, the susceptibility of psoriasis to exogenous stress and, thus, NK₁ antagonists might be beneficial in psoriasis therapy [40]. Likewise, ACE has been associated with this disease [63], although the exact role of the enzyme is somewhat obscure. Some studies proposed elevated serum ACE levels in psoriasis as a diagnostic marker and suggested that increased amounts of the pro-inflammatory Ang II may contribute to the pathogenesis of this disease [32,70]. Several other reports, however, have associated pharmacologic ACE inhibition with the induction or exacerbation of psoriasis [14,33,107]. Thus, one adverse side-effect of vaso-peptidase inhibitors may be the worsening of psoriasis due to prolonged availability of SP or BK.

Summary for the Clinician

As outlined in this chapter, endopeptidases have multiple substrates and different modes of action in the skin and elsewhere. Particularly, NEP is an excellent example that endopeptidases in addition to their metabolizing/degrading function mediate biotransformation of inert precursor peptides to active mediators, and act as convertases by converting one bioactive receptor ligand into a second peptide that activates another receptor. Finally, competitive cleavage of a given substrate generates a peptide with opposing physiological effects to that produced by a closely related enzyme (i.e., NEP: Ang I \rightarrow Ang₁₋₇, hypotensive vs. ACE: Ang I \rightarrow Ang II = Ang₁₋₈, hypertensive). Evidently, a multiplicity of functions arises from these activities, and thus endopeptidases constitute important milestones in cardiovascular regulation, in neurogenic inflammation and immune recognition, or in tumorigenesis. Although it is tempting to drug-target neprilysin and others to achieve expressional or enzymatic regulation, one may likely replace one evil with another. This dilemma is best exemplified by attempts to simultaneously inhibit multiple vaso-peptidases (ACE, NEP, and ECE) with one inhibitor. Although this is a promising strategy for the treatment of cardiovascular disorders, recent large-scale phase IV clinical studies have questioned this benefit [9]. Evidently, the ACE/NEP inhibitor-dependently increased kinin, SP, and endothelin levels elevated the risk of potentially life-threatening angioedema, with an even higher incidence particularly in African-Americans [66]. Vaso-peptidase inhibition may also worsen cutaneous disorders with a neurogenic component [91], promote Ag sensitization, or trigger elicitation of allergic inflammation in already sensitized individuals. As explained above, NEP may also have (neuro-) protective roles in cancer and Alzheimer's disease [35]. Thus, future attempts of inhibiting peptidases with A β -catabolizing properties essentially require the design of drugs incapable of crossing the blood-brain-barrier, and critical validation of efficacy and safety of this promising therapeutic principle is mandatory. Vice versa, since downregulated NEP expression or activity frequently accompanies acute or chronic inflammation and the malignancy of tumors, an artificial substitution seems to be desirable. Thus, a tissue-specific use of recombinant enzymes [94] or a local enhancement of NEP expression and activity may provide a novel mechanism-based therapeutic approach towards management of (neurogenic) inflammatory skin disorders. Unfortunately, despite a revived interest in NEP regulation due to the impact of NEP on the CNS A β level, only few data are available demonstrating an upregulation of NEP, that is, by glucocorticoids, thrombin, or cAMP [28,29,35]. Obviously, the future challenge remains to develop tools to control peptidase expression and activity in a highly selective cell and tissue-specific manner.

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