ARTICLE IN PRESS

Progress in Lipid Research

Progress in Lipid Research xxx (2015) xxx-xxx

Contents lists available at ScienceDirect

Progress in Lipid Research

journal homepage: www.elsevier.com/locate/plipres



Review

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The increasing role of phosphatidylethanolamine as a lipid receptor in the action of host defence peptides

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ARTICLE INFO

- 16 Article history: Received 27 November 2014 17
- 18 Received in revised form 26 February 2015
- Accepted 27 February 2015 19
- 20
- Available online xxxx
- 21 Keywords:
- 22 Amyloid-forming host defence peptides 23 Cyclotides
- 24 Host defence peptides
- 25 Lantibiotics 26
- Lipid receptors 27 Phosphatidylethanolamine (PE)
- 28
- PE-binding 29 Maximin H5
- 30

ABSTRACT

Host defence peptides (HDPs) are antimicrobial agents produced by organisms across the prokaryotic and eukaryotic kingdoms. Many prokaryotes produce HDPs, which utilise lipid and protein receptors in the membranes of bacterial competitors to facilitate their antibacterial action and thereby survive in their niche environment. As a major example, it is well established that cinnamycin and duramycins from Streptomyces have a high affinity for phosphatidylethanolamine (PE) and exhibit activity against other Gram-positive organisms, such as Bacillus. In contrast, although eukarvotic HDPs utilise membrane interactive mechanisms to facilitate their antimicrobial activity, the prevailing view has long been that these mechanisms do not involve membrane receptors. However, this view has been recently challenged by reports that a number of eukaryotic HDPs such as plant cyclotides also use PE as a receptor to promote their antimicrobial activities. Here, we review current understanding of the mechanisms that underpin the use of PE as a receptor in the antimicrobial and other biological actions of HDPs and describe medical and biotechnical uses of these peptides, which range from tumour imaging and detection to inclusion in topical microbicidal gels to prevent the sexual transmission of HIV.

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1. Introduction

Abbreviations: AD, Alzheimer's disease; DMPE, 1,2-dimyristoyl-sn-glycero-3phosphoethanolamine; HDPs, host defence peptides; HIV, human immunodeficiency virus; IAPPs, islet amyloid polypeptides; LPS, lipopolysaccharide; MD, molecular dynamic; PE, phosphatidylethanolamine.

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http://dx.doi.org/10.1016/j.plipres.2015.02.003 0163-7827/© 2015 Published by Elsevier Ltd.

The production of host defence peptides (HDPs) as a defence 62 and survival strategy is practiced by organisms across the prokary-63 otic [1,2] and eukaryotic kingdoms [3,4]. HDPs from bacteria, more 64 generally known as bacteriocins, inhibit the growth of similar or 65 closely related bacterial strains and it is well established that many 66 of these peptides utilise lipid and protein receptors in the 67

Please cite this article in press as: Phoenix DA et al. The increasing role of phosphatidylethanolamine as a lipid receptor in the action of host defence peptides. Prog Lipid Res (2015), http://dx.doi.org/10.1016/j.plipres.2015.02.003

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membranes of target bacteria to facilitate their antibacterial action [5–8]. For example, class I bacteriocins, or lantibiotics, such as nisin from *Lactococcus lactis*, are commonly used as food preservatives and are produced by a large number of Gram-positive bacteria to attack other Gram-positive bacteria [9,10]. These peptides commonly target lipid II, which is a major cell wall precursor, to exert their antimicrobial action through pore formation in target membranes and/or the inhibition of cell wall biosynthesis [11,12]. Other receptors, which promote the activity of bacterial HDPs are increasingly being reported [5–8] with most recent examples, including: the maltose ABC transporter for the class IIc bacteriocin, garvicin ML [13]; a Zn-dependent metallopeptidase for the class IId bacteriocin LsbB [14]; and an undecaprenyl pyrophosphate phosphatase for the class IIb bacteriocin lactococcin G [15].

82 HDPs produced by eukaryotic organisms are evolutionarily con-83 served, multi-functional components of innate immune systems 84 that show potent activity against a wide spectrum of microbes 85 [16–19] and cancer cells [20,21]. Numerous studies have shown 86 that in the vast majority of cases, this activity involves membrane 87 interactive mechanisms but in contrast to bacterial HDPs, the pre-88 vailing view has long been that these mechanisms do not require 89 the use of membrane receptors [20-23]. However, this view has been increasingly questioned in the light of recent studies [20] 90 91 and several investigations have suggested that the antibacterial 92 mechanisms of a number of HDPs may utilise protein receptors to 93 facilitate membrane translocation and interaction with cytoplasmic 94 targets [24–26]. There is also increasing evidence that eukaryotic 95 HDPs utilise lipid receptors in their antimicrobial mechanisms 96 [23]. For example, it has been recently shown that lipoprotein in 97 the outer membrane of Gram-negative bacteria serves as a cell sur-98 face receptor to promote the action of a number of HDPs against these organisms [27], including SMAP-29 from sheep [28], and 99 CAP-18, and LL-37 from humans [29,30]. It has also recently been 100 101 shown that interaction with microbe-specific lipid receptors is a 102 key step in the antimicrobial action of a number of defensins [31-103 33], which are HDPs found across eukaryotes [34–42]. For example, 104 lipid II has been shown to promote the inhibition of cell wall biosyn-105 thesis and antibacterial activity of some defensins [31–33], includ-106 ing: Cg-Defh1, Cg-Defh2, and Cg-Defm from the oyster, Crassostrea 107 gigas [43], and plectasin from the fungus, *Pseudoplectania nigrella* 108 [44]. However, when fungi are the targets of defensins, it has been 109 demonstrated that fungal-specific sphingolipids can promote the activity of a number of defensins [31-33], as in the case of RsAFP2 110 111 from the plant, Raphanus sativus, whose interaction with glucosylceramides promotes fungal death by a variety of mechanisms, 112 113 including membranolysis and apoptosis induction [45].

114 Currently, a major focus of research into the role of lipid receptors 115 in the antimicrobial action of HDPs is phosphatidylethanolamine 116 (PE), which is present at high levels in the membranes of Gram-neg-117 ative bacteria and is also found in some Gram-positive bacteria [46]. 118 PE is also ubiquitous in eukaryotic membranes, and is predominantly a constituent in the inner leaflet of the plasma membrane 119 [47,48]. However, emerging evidence has shown that in addition 120 to serving as a structural element in the bilayer, PE is translocated 121 122 or redistributed across the membrane in a number of biological events [49,50] such as apoptosis and malignant transformation 123 124 where membrane asymmetry is compromised and the lipid is exposed to the extracellular milieu [51]. Here, we review current 125 126 understanding of HDPs that utilise PE as a receptor in their antimi-127 crobial and other biological actions.

128 2. Amyloid-forming HDPs

129 It is well established that human Aβ40 and Aβ42 are the major 130 amyloid forming peptides associated with Alzheimer's disease (AD) but currently, their normal physiological functions are 131 unknown [52]. However, a major recent study presented strong 132 evidence to suggest that this function may be to serve as HDPs in 133 protecting the human brain against microbial infection [53]. 134 Several authors have proposed that the ability of A^β40 and A^β42 135 kill bacteria involves binding to this lipid and the induction of amy-136 loid-mediated membrane pore formation [54,55]. In addition to 137 Aβ40 and Aβ42, a number of major amyloid forming peptides 138 involved in neurodegenerative and other diseases are known to 139 possess antimicrobial activity [55]; it has been proposed that PE-140 binding may also be involved in membrane pore formation by at 141 least some of these peptides [56,57]. For example, the human islet 142 amyloid polypeptides (IAPPs), which are involved in type II dia-143 betes, have recently been shown to possess antibacterial activity 144 that is associated with the ability of these peptides to oligomerise 145 and thereby induce the disruption of bacterial membranes [58]. 146 IAPPs have been shown to induce leakage in bacterial membranes 147 using nucleation-dependent mechanisms of membrane pore for-148 mation that show many similarities to those of amyloidogenic 149 HDPs [55,59]. Moreover, IAPPs were found to exhibit full cross-co-150 operativity with magainin 2, which is an amphibian AMP shown to 151 form amyloid fibrils [60], in the induction of membrane leakage 152 and inhibition of bacterial growth [59]. Studies on the membrane 153 interactions of IAPPs found that their monomers had a weak affin-154 ity for membranes containing PE but amyloid fibres of the peptide 155 interacted strongly and specifically with this lipid to induce high 156 levels of membrane disruption *via* a pore-type mechanism [56]. 157

3. Lantibiotics

Amongst the first PE-binding HDPs to be described were cin-159 namycin, duramycin, duramycin B and duramycin C, which are 160 structurally related, class I, type B lantibiotics (Fig. 1A) [61,62]. 161 These peptides are produced by some strains of the Gram-positive 162 bacteria, Streptomyces, and exhibit weak activity against other 163 Gram-positive organisms, such as species of Bacillus, which 164 appears to involve the compromise of membrane integrity and/or 165 function [61,63–65]. It is well established that cinnamycin and 166 the duramycins have a high affinity and specificity for PE 167 [61,62,66–70] and there is evidence to suggest that this affinity 168 plays a role in the antibacterial activity of these peptides although 169 this role has, as yet, not been fully elucidated [61,63–65]. However, 170 consistent with this role for PE, it has been shown that the lipid is 171 absent from the membranes of species of Bacillus subtilis and 172 Bacillus firmus with resistance to duramycin [71,72]. One study 173 suggested that this resistance mechanism is underpinned by the 174 replacement of PE in bacterial membranes by its plasmalogen 175 form, plasmenylethanolamine [73] but more recent studies have 176 suggested that both duramycin and cinnamycin have a general 177 ability to bind PE-lipids, including its plasmalogen form [74]. In 178 addition, these HDPs also show a number of PE-mediated activities 179 against eukaryotic cells [70,75–78], which contrasts to most lan-180 tibiotics whose activity is generally limited to bacterial cells 181 [79,80], For example, cinnamycin possesses haemolytic ability 182 [81] and appears to promote this ability by inducing the transbi-183 layer movement of PE from the inner leaflet of erythrocyte mem-184 branes to be exposed on the outer surface of these cells [82]. 185 More recent studies have suggested that duramycin also promotes 186 the transbilayer movement of the lipid [74] and that binding of the 187 peptide to PE exposed on the surface of pancreatic tumour cells led 188 to the death of these cells via the disruption of the structure and/or 189 function of the cancer cell membrane [83,84]. 190 191

Cinnamycin and duramycin are by far the best characterised of the PE-binding, lantibiotics, in part, because they are one of the very few examples of small peptides with a three-dimensional

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