Supplementary Information

Solution NMR and racemic crystallography provide insights

into a novel structural class of cyclic plant peptides

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Supplementary Figure 1: Validation of L-PDP-23 dimer electron density map. The L-PDP-23 crystal structure is an excellent fit within a composite omit map (generated using PHENIX),¹ except for His5 and His6 in each monomer.



Supplementary Figure 2: Overlay of the 3D structures of PDP-23 in both monomeric and dimeric forms with PDP-24. (A) Overlay of one molecule of the symmetrical homodimeric form of PDP-23 (cyan) with PDP-24 (magenta). (B) Overlay of the monomeric form of PDP-23 (blue) with PDP-24 (magenta). Labels are supplied for orientation and are given in colour when there is a difference in residue between the two peptides. PDP-24 adopts a structure more akin to the structure of dimeric PDP-23 than monomeric PDP-23 but yet does not adopt a homodimer in aqueous solution.

Peptides	MIC (µM)			
	E. coli	S. aureus	C. albicans	
PDP-23	>80	>80	>80	
PDP-24	>80	>80	>80	
LL-37 (Positive control)	0.625	1.25	2.5	

Supplementary Table 1: Minimum inhibitory concentrations of PDP-23 and -24 against common microbes

Peptide	Average dose (µg/kg)	Average weight (g)	N	% Healthy
PDP-23				
	3897.1	2.57	5	100
	382.8	2.61	5	100
	37.0	2.71	5	100
	4.0	2.48	5	100
	0.4	2.41	5	100
α-1 (no HYP)				
	203.9	2.45	3*	0
	109.1	2.29	5	0
	23.1	2.17	5	20
	4.9	2.03	5	100
	0.8	2.56	5	100
	0.2	2.23	5	100
Control (H ₂ O)				
	0.0		5	100

Supplementary Table 2: Insecticidal assay dosages, insect weights, N values and % healthy specimens 24 h post injection

*Two animals were excluded from the assay due to failed injections.

References:

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