

1 Mechanism of antimicrobial activity of
2 honeybee (*Apis mellifera*) venom on Gram-
3 negative bacteria: *Escherichia coli* and
4 *Pseudomonas* spp.
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22 Table S1. Composition of dry honeybee venom, (Shipolini, 1984; Pucca et al., 2019)

Class of molecules	Compound	Amount (% of venom, dry weight)
Enzymes	Phospholipase A ₂ ^b	10-12
	Hyaluronidase ^a	1,5-2
Peptides	Melittin ^a	40-50
	Apamin	3
	Mast Cell Degranulating Peptide (MCDP)	2
Amines	Histamine ^c	0.6-1.6
	Dopamine	0.13-1
Carbohydrates	Glucose	0.7
	Fructose	0.9
Amino acids	γ -aminobutyric acid and β -aminoisobutyric acid	1
Lipids		5
Minerals	P, Ca, Mg	3-4
a: minor allergen / potent allergen, b: major allergen, c: biogenic amine		

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29 Table S2: Summary of studies on antimicrobial activity of honeybee venom against bacteria.

Component	Bacteria		Strains	Assay	MIC (µg/mL)	MBC (µg/mL)	References	
	Gram positive	Gram negative						
Bee venom		<i>Ac. baumannii</i> BAA	ATCC 747	MM	30	60	Al-Ani et al., 2015	
	<i>B. subtilis</i>		ATCC 6051	MM	8	10		
		<i>Bur. mallei</i>	RTCC 2375	ADM			Zolfagharian et al., 2016	
		<i>Bur. pseudomallei</i>	RTCC 2375	ADM				
		<i>E. coli</i>	ATCC 35218	BDM	ND		Hegazi et al., 2017	
			ATCC 25922	MM	60	60	Al-Ani et al., 2015	
			SR 200756	MM	100	200		
			KL 206272	MM	200	200		
					ADM	NA	NA	Perumal Samy et al., 2007
		<i>E. coli</i> O157:H7	ATCC 35150	MM	100	200	Al-Ani et al., 2015	
			ATCC 25923				Zolfagharian et al., 2016	
		<i>En. aerogenes</i>	-	ADM	NA	NA	Perumal Samy et al., 2007	
		<i>Ent. casseliflavus</i>		ATCC 700327	MM	10	30	Al-Ani et al., 2015
		<i>Ent. feacalis</i>			BDM	ND		Hegazi et al., 2017
			ATCC 29212	MM	100	200	Al-Ani et al., 2015	
			ATCC 4082	MM	20	-	Leandro et al., 2015	
	<i>Ent. feacalis</i> VanB		ATCC 51299	MM	200	500	Al-Ani et al., 2015	
	<i>Ent. VRE</i>		ATCC 902291	MM	200	500		
				ATCC 102105	MM	200	500	

		ATCC 902316	MM	200	500		
	<i>K. pneumoniae</i>	ATCC 27736	BDM	ND		Hegazi et al., 2017	
		ATCC 700603	MM	30	60	Al-Ani et al., 2015	
		ATCC 800877	MM	500	500		
		KL 206436	MM	500	500		
	<i>K. oxytoca</i>	ATCC 700324	MM	500	500		
<i>Lact. casei</i>		EMCC 1093 T	BDM	ND		Hegazi et al., 2017	
		ATCC 11578	MM	20	-	Leandro et al., 2015	
<i>Lis. monocytogene</i>			BDM	ND	-	Hegazi et al., 2017	
<i>Myc. smegmatis</i>		ATCC 19420	MM	200	200	Al-Ani et al., 2015	
<i>Myc. fortuitum</i>		ATCC 6841	MM	100	100		
<i>Myc. phlei</i>		ATCC 11758	MM	800	800		
	<i>Pr. mirabilis</i>	-	ADM	-	-	Perumal Samy et al., 2007	
	<i>Pr. vulgaris</i>	-	ADM	-	-		
	<i>P. aeruginosa</i>	ATCC 27853	BDM	ND	-	Hegazi et al., 2017	
			MM	500	>500	Al-Ani et al., 2015	
						Zolfagharian et al., 2016	
			KL 206466	MM	500	>500	Al-Ani et al., 2015
			KL 206444	MM	>500	>500	
			-	ADM	-	-	Perumal Samy et al., 2007
	<i>Sal. choleraesuis</i>	ATCC 554	MM	500	>500	Al-Ani et al., 2015	
	<i>Sal. typhimurium</i>	ATCC 25923				Zolfagharian et al., 2016	
	<i>Sh. flexneri</i>	ATCC 29903	MM	60	100	Al-Ani et al., 2015	
<i>Staph. aureus</i>		ATCC 25923	BDM	ND	-	Hegazi et al., 2017	

						Zolfagharian et al., 2016
			MM	10	30	Al-Ani et al., 2015
	ATCC 29213		MM	60	200	
	USA300 (LAC)		MM	0.78	ND	Choi et al., 2015
	Newman		MM	0.78	ND	
	MW2		MM	1.56	ND	
	MRSA1		MM	3.12	ND	
	MRSA2		MM	1.56	ND	
	ISP4790		MM	6.25	ND	
	MU50		MM	6.25	ND	
	ATCC 6538P			17	ND	Kokot et al., 2009
	-		ADM	23.2±1.09	-	Perumal Samy et al., 2007
<i>Staph. aureus</i> BAA	ATCC 977		MM	30	60	Al-Ani et al., 2015
<i>Staph. aureus</i> MRSA	ATCC 1042		MM	60	100	
	ATCC 106804		MM	60	200	
	ATCC 106188		MM	60	100	
	ATCC 106091		MM	60	200	
<i>Staph. epidermidis</i>	ATCC 14990		MM	60	100	
<i>Staph. saprophyticus</i>	ATCC 15305		MM	10	10	
<i>Strep. agalactiae</i>	ATCC 27956		MM	40	40	Choi et al., 2015
	CNCTC 10/84		MM	6.25	ND	
<i>Strep. bovis</i>	NEM 760		MM	1.56	ND	
<i>Strep. epidermidis</i>	RP62a		MM	0.78	ND	
<i>Strep. gordonii</i>	M99		MM	6.25	ND	
<i>Strep. mitis</i>	ATCC 49452		MM	40	-	Leandro et al., 2015
<i>Strep. mutans</i>	EMCC 1815T		BDM	ND	-	Hegazi et al., 2017
	ATCC 25175		MM	20	-	Leandro et al., 2015

	<i>Strep. oralis</i>		ATCC 35037	MM	100	100	Al-Ani et al., 2015
	<i>Strep. pneumonia</i>		TIGR4	MM	3.12	-	Choi et al., 2015
	<i>Strep. pyogenes</i>			BDM	ND	-	Hegazi et al., 2017
			ATCC 12344	MM	100	200	Al-Ani et al., 2015
	<i>Strep. salivarius</i>		ATCC 25975	MM	20	-	Leandro et al., 2015
	<i>Strep. sanguinis</i>		ATCC 10556	MM	30	-	
	<i>Strep. sobrinus</i>		ATCC 33478	MM	40	-	
	<i>Strep. thermophilus</i>		ATCC 19258	MM	30	60	Al-Ani et al., 2015
Melittin		<i>Ac. baumannii</i> BAA	ATCC 747	MM	30	30	Al-Ani et al., 2015
		<i>B. subtilis</i>	ATCC 6051	MM	6	30	
		<i>E. coli</i>	ATCC 25922	MM	30	60	
			SR 200756	MM	30	60	
			KL 206272	MM	30	60	
		<i>E. coli</i> O157:H7	ATCC 35150	MM	30	60	
		<i>Ent. casseliflavus</i>	ATCC 700327	MM	8	20	
		<i>Ent. faecalis</i>	ATCC 29212	MM	30	100	
			ATCC 4082	MM	6	-	
		<i>Ent. faecalis</i> VanB	ATCC 51299	MM	50	100	
		<i>Ent. VRE</i>	ATCC 902291	MM	50	100	Al-Ani et al., 2015
		<i>Lact. casei</i>	ATCC 11578	MM	4	-	Leandro et al., 2015
		<i>K. pneumoniae</i>	ATCC 700603	MM	100	300	Al-Ani et al., 2015
			ATCC 800877	MM	100	300	
	<i>K. oxytoca</i>	ATCC 700324	MM	60	500		

	<i>P. aeruginosa</i>	ATCC 27853	MM	100	100	
		KL 206466	MM	100	100	
		KL 206444	MM	100	100	
<i>Staph. aureus</i>		ATCC 25923	MM	10	30	
		ATCC 29213	MM	6	10	
<i>Staph. aureus</i> BAA		ATCC 977	MM	8	8	
<i>Staph. aureus</i> MRSA		ATCC 1042	MM	100	30	
		ATCC 106804	MM	30	50	
		ATCC 106188	MM	10	50	
		ATCC 106091	MM	30	50	
<i>Staph. epidermidis</i>		ATCC 14990	MM	10	30	
<i>Staph. saprophyticus</i>		ATCC 15305	MM	10	30	
<i>Strep. pyogenes</i>		ATCC 12344	MM	10	30	
<i>Strep. oralis</i>		ATCC 35037	MM	200	200	
<i>Strep. agalactia</i>		ATCC 27956	MM	30	30	
<i>Strep. mitis</i>		ATCC 49452	MM	10		Leandro et al., 2015
<i>Strep. mutans</i>		ATCC 25175	MM	40		
PLA ₂	<i>E. coli</i>	-	ADM	NA	NA	Perumal Samy et al., 2007
	<i>En. aerogenes</i>	-	ADM	NA	NA	
	<i>Ent. feacalis</i>	ATCC 4082	MM	NA	NA	Leandro et al., 2015
<i>Lact. casei</i>		ATCC 11578	MM	400	-	
	<i>Pr. mirabilis</i>	-	ADM	NA	NA	Perumal Samy et al., 2007
	<i>Pr. vulgaris</i>	-	ADM	NA	NA	
	<i>P. aeruginosa</i>	-	ADM	NA	NA	
<i>Staph. aureus</i>		-	ADM	13.3±0.83	-	
<i>Strep. mitis</i>		ATCC 49452	MM	NA	NA	Leandro et al., 2015

	<i>Strep. mutans</i>		ATCC 25175	MM	NA	NA	
Ac.: <i>Acinetobacter</i>		En.: Enterobacter		Lis.: <i>Listeria</i>		Sal.: <i>Salmonella</i>	
B.: <i>Bacillus</i>		Ent.: Enterococcus		Myc.: <i>Mycobacterium</i>		Sh.: <i>Shigella</i>	
Bur.: <i>Burkholderia</i>		K.: <i>Klebsiella</i>		Pr.: <i>Proteus</i>		Staph.: <i>Staphylococcus</i>	
E.: <i>Escherichia</i>		Lact.: <i>Lactobacillus</i>		P.: <i>Pseudomonas</i>		Strep.: <i>Streptococcus</i>	
ADM: Agar Dilution Method		BDM: Broth Dilution Method		MM: Microdilution Method		ND: Not determined	

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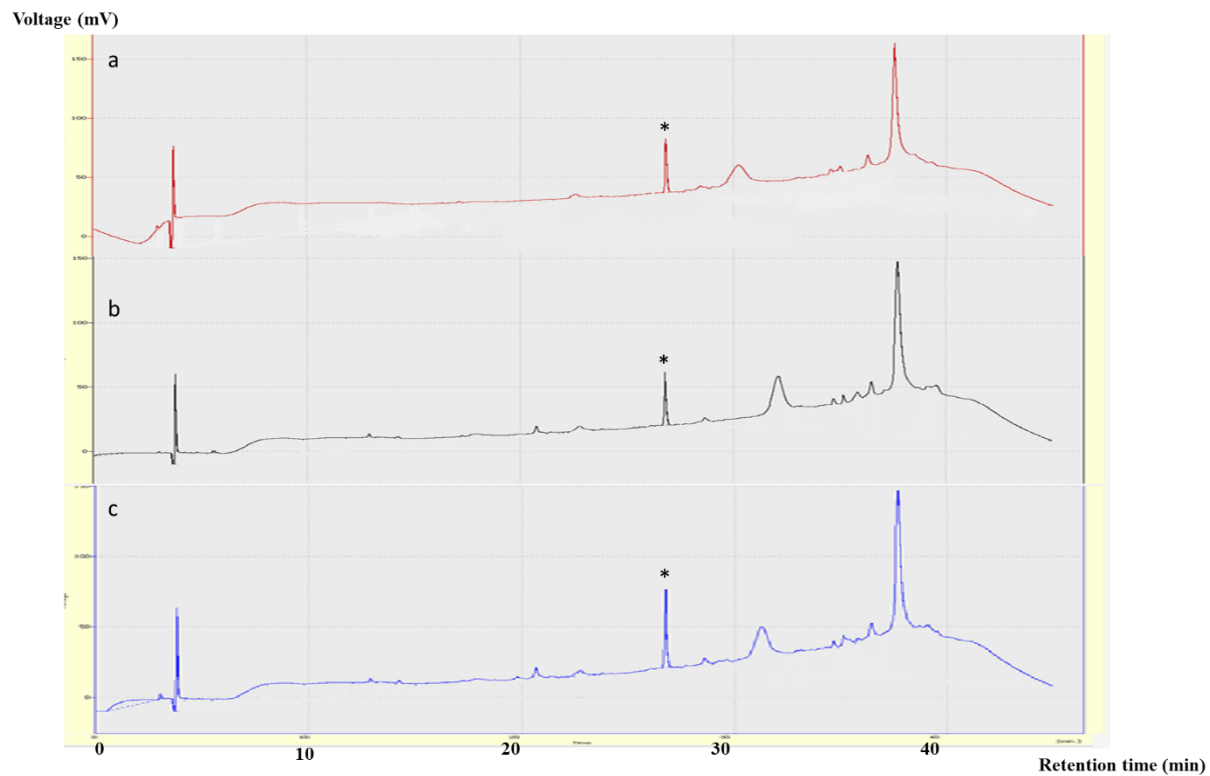
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45 Figure S1. HPLC chromatograms of melittin standard aqueous solution (50 μ g/mL) (a) BV-1 (b) and
46 BV-2 (c) aqueous solutions (150 μ g/mL) at 220nm. Peak (*): melittin. Detection was at 220nm
47 (Chromatographic conditions as in Materials and Methods section).



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