

Mechanism of action of naturally occurring antimicrobials against *Clostridium difficile*

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INTRODUCTION

- Clostridium difficile* causes infections of the gastrointestinal tract.¹
- Disease severity can range from self-limiting diarrhea to severe manifestations such as pseudomembranous colitis and toxic megacolon.²
- C. difficile* infection (CDI) has been managed with the conventional antimicrobials metronidazole (for mild to moderate CDI) and vancomycin (for severe CDI).³
- Several issues are associated with the use of conventional agents, including a high recurrence rate of 20-25% and reduced *in vitro* efficacy.³
- Natural compounds generally have broad-spectrum antimicrobial activity with several showing activity against *C. difficile*.⁴
- Compounds derived from plant extracts have great antimicrobial potential against drug-resistant microorganisms and, unlike conventional antimicrobials, they are less susceptible to the development of antimicrobial resistance.^{4,5}



Figure 1. Gram stain of *C. difficile*

OBJECTIVE

This study aimed to investigate the mechanism of action of natural occurring compounds and plant extracts against *C. difficile*.

METHODS

The mechanism of action of five compounds with bactericidal activity (cinnamon root powder, peppermint oil, *trans*-cinnamaldehyde, menthol and zingerone), and four with bacteriostatic activity (fresh garlic bulb extract, garlic clove powder, *Leptospermum* honey and allicin) against two *C. difficile* strains was investigated.

Assays:

- Bactericidal activity: Time kill kinetics and ATP-leakage assays.
- Alterations in cell permeability: determination of protein leakage using the Bradford assay and propidium iodide uptake assays.
- Inhibitory effect of compounds on prokaryotic translation/protein synthesis: An *in vitro* transcription/translation experiment using Promega's *E. coli* S30 extract system for circular DNA linked to the Steady-Glo®.
- Detection of antimicrobial cross-resistance: broth microdilution, comparing the MICs of products against a panel of antimicrobial *C. difficile* strains previously characterised for AMR phenotype and associated genotype.



Figure 2. Time-kill assay viable counts



Figure 3. Prokaryotic translation/protein synthesis



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RESULTS

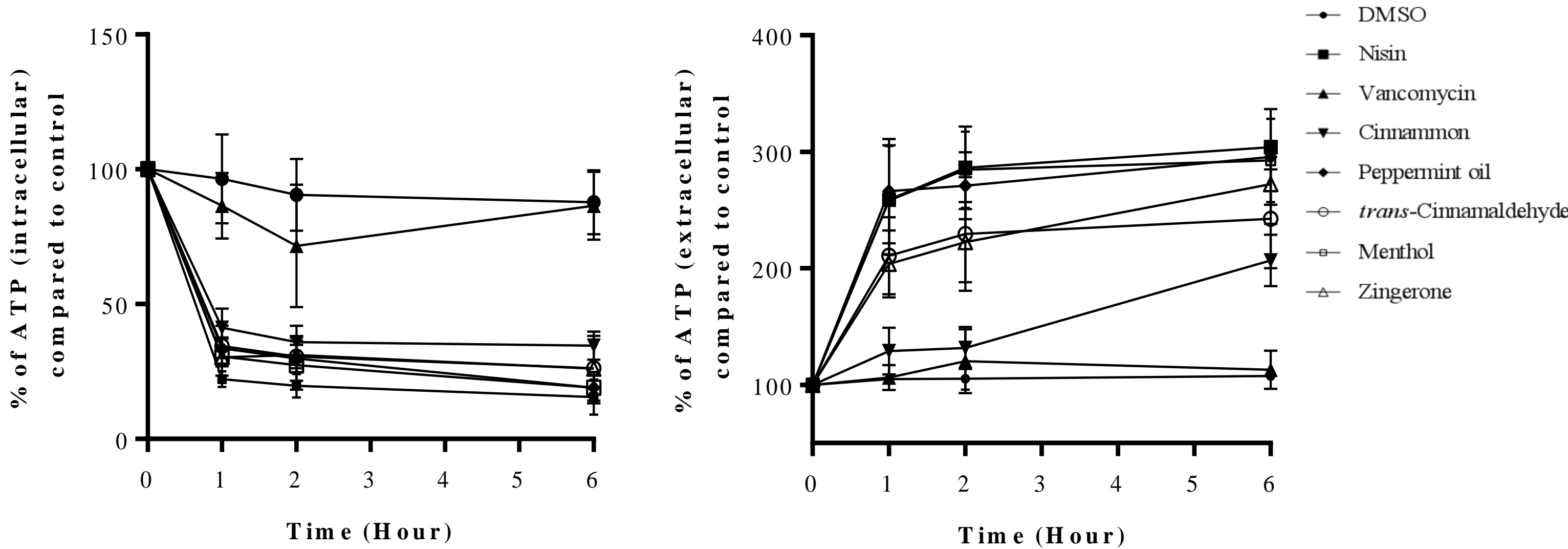
Time-kill kinetics

Table 1. Mean log10 cfu/ml reductions for *C. difficile* NCTC 13366 following of exposure to treatments.

Treatment	Conc	Clostridium difficile NCTC 13366									
		1h		2h		4h		6h		24h	
		EXP	STA	EXP	STA	EXP	STA	EXP	STA	EXP	STA
Cinnamon	2 × MIC	0.7 ± 0.3	1.6 ± 0.4	1.4 ± 0.2	2.4 ± 0.3	1.5 ± 0.2	2.6 ± 0.2	1.8 ± 0.5	2.8 ± 0.2	2.2 ± 0.6	3.6 ± 0.4
	4 × MIC	2.7 ± 0.3	NA	2.7 ± 0.4	NA	3.2 ± 0.7	NA	3.6 ± 0.5	NA	3.7 ± 0.4	NA
	8 × MIC	2.8 ± 0.4	NA	3.0 ± 0.7	NA	3.5 ± 0.6	NA	3.6 ± 0.5	NA	3.8 ± 0.4	NA
Peppermint oil	2 × MIC	2.0 ± 0.2	2.1 ± 0.5	2.2 ± 0.2	2.2 ± 0.5	2.6 ± 0.4	2.8 ± 0.3	3.1 ± 0.2	3.1 ± 0.3	3.6 ± 0.4	3.8 ± 0.7
	4 × MIC	2.5 ± 0.4	2.1 ± 0.4	3.0 ± 0.4	2.3 ± 0.5	3.0 ± 0.4	2.9 ± 0.3	3.6 ± 0.2	3.2 ± 0.3	3.9 ± 0.1	3.8 ± 0.6
	8 × MIC	2.5 ± 0.4	NA	3.1 ± 0.4	NA	3.2 ± 0.3	NA	3.7 ± 0.2	NA	3.9 ± 0.2	NA
<i>trans</i> -Cinnamaldehyde	2 × MIC	2.6 ± 0.8	1.1 ± 0.3	3.1 ± 0.5	1.7 ± 0.4	3.6 ± 0.6	2.0 ± 0.4	4.1 ± 0.2	2.6 ± 0.2	4.2 ± 0.2	3.0 ± 0.2
	4 × MIC	2.7 ± 0.6	1.2 ± 0.3	3.2 ± 0.2	2.3 ± 0.5	3.6 ± 0.7	2.8 ± 0.3	4.2 ± 0.2	3.1 ± 0.1	4.2 ± 0.1	3.3 ± 0.4
	8 × MIC	2.9 ± 0.6	1.7 ± 0.4	3.6 ± 0.4	2.8 ± 0.3	3.8 ± 0.7	3.2 ± 0.3	3.7 ± 0.6	3.3 ± 0.2	4.2 ± 0.2	3.3 ± 0.3
Menthol	2 × MIC	0.4 ± 0.4	1.8 ± 0.5	1.2 ± 0.3	2.2 ± 0.2	1.5 ± 0.5	2.5 ± 0.1	2.6 ± 0.9	2.5 ± 0.1	3.1 ± 0.9	3.6 ± 0.2
	4 × MIC	1.6 ± 0.7	2.2 ± 0.9	1.8 ± 0.8	2.5 ± 0.7	3.9 ± 0.2	2.9 ± 0.4	3.9 ± 0.2	3.1 ± 0.3	4.0 ± 0.1	3.7 ± 0.2
	8 × MIC	3.2 ± 0.5	2.2 ± 0.9	3.5 ± 0.6	2.7 ± 0.9	4.0 ± 0	3.2 ± 0.5	4.0 ± 0	3.2 ± 0.4	4.1 ± 0.2	3.9 ± 0.1
Zingerone	2 × MIC	2.1 ± 0.6	1.4 ± 0.6	2.9 ± 0.5	2.0 ± 0.6	2.9 ± 0.6	2.5 ± 0.4	2.9 ± 0.6	2.6 ± 0.2	3.0 ± 0.6	3.0 ± 0.1
	4 × MIC	2.5 ± 0.4	2.3 ± 0.4	3.1 ± 0.4	2.6 ± 0.1	3.3 ± 0.3	2.8 ± 0.1	3.4 ± 0.4	3.0 ± 0.2	4.2 ± 0.7	3.2 ± 0.4
	8 × MIC	3.1 ± 0.2	2.4 ± 0.6	3.4 ± 0.1	2.8 ± 0.3	3.7 ± 0.3	3.0 ± 0.3	4.3 ± 0.6	3.2 ± 0.2	4.2 ± 0.2	3.3 ± 0.5

EXP, exponential; STA, stationary; MIC, minimum inhibitory concentration; Conc: concentration; NA, not applicable; Sterile distilled water (SDW) and dimethyl sulfoxide (DMSO) (2.5% V/V) were used as negative controls.

ATP-leakage assay



Cell permeability assays

Table 2. Effect of products on *C. difficile* NCTC 13366, measured by protein leakage and propidium iodide uptake assays.

Treatment	Protein leakage assay			Propidium iodide uptake		
	1h	2h	6h	1h	2h	6h
Cinnamon root powder						
1 × MIC (75 mg/ml)	7.0 ± 2.3	3.0 ± 1.8	24.8 ± 5.7	324 ± 53	376 ± 136	602 ± 128
2 × MIC (150 mg/ml)	4.7 ± 5.0	24.4 ± 10.6	83.1 ± 10.6	446 ± 64	616 ± 168	1061 ± 228
Peppermint oil						
1 × MIC (8% v/v)	19.2 ± 6.8	23.6 ± 5.2	74.3 ± 3.9	241 ± 60	444 ± 108	1206 ± 186
2 × MIC (16% v/v)	29.6 ± 9.1	62.9 ± 2.7	90.7 ± 11.2	431 ± 156	548 ± 111	1387 ± 294
4 × MIC (32% v/v)	32.8 ± 11.0	74.1 ± 2.6	99.5 ± 8.7	256 ± 14	399 ± 142	1061 ± 228
<i>trans</i>-Cinnamaldehyde						
1 × MIC (0.02% v/v)	0.0 ± 0.0	0.2 ± 0.3	2.2 ± 3.8	548 ± 156	854 ± 181	523 ± 10.8
2 × MIC (0.04% v/v)	0.0 ± 0.0	3.9 ± 3.9	6.9 ± 5.9	724 ± 43	1831 ± 69	745 ± 14.5
4 × MIC (0.08% v/v)	0.0 ± 0.0	9.1 ± 7.4	18.2 ± 8.1	985 ± 22	1186 ± 37	618 ± 8.1
8 × MIC (0.16% v/v)	27.2 ± 14.9	31.0 ± 21.3	45.9 ± 12.7	1115 ± 46	1446 ± 70	1444 ± 49
Menthol						
1 × MIC (9.4 mg/ml)	0.0 ± 0.0	36.0 ± 6.9	42.2 ± 17.6	308 ± 12	315 ± 99	223 ± 21
2 × MIC (18.8 mg/ml)	24.6 ± 14.8	37.2 ± 12.6	51.1 ± 13.3	287 ± 21	287 ± 88	487 ± 99
4 × MIC (37.5 mg/ml)	37.7 ± 4.8	51.5 ± 16.7	62.3 ± 22.6	337 ± 8.9	463 ± 101	513 ± 88
8 × MIC (75 mg/ml)	24.2 ± 8.9	48.8 ± 20.1	76.6 ± 13.5	461 ± 84	499 ± 88	987 ± 44
Zingerone						
1 × MIC (9.4 mg/ml)	0.0 ± 0.0	0.0 ± 0.0	0.3 ± 0.4	698 ± 73	595 ± 20	990 ± 99
2 × MIC (18.8 mg/ml)	11.5 ± 5.4	5.2 ± 3.3	3.1 ± 3.3	739 ± 67	574 ± 19	1247 ± 29
4 × MIC (37.5 mg/ml)	5.7 ± 3.3	20.3 ± 9.6	35.2 ± 15.3	1263 ± 425	919 ± 21	1301 ± 12
8 × MIC (75 mg/ml)	38.9 ± 13.9	50.7 ± 13.4	41.7 ± 15.1	1101 ± 418	1601 ± 122	1214 ± 57

MIC, minimum inhibitory concentration; Values in bold differ significantly from controls (SDW and DMSO). Sterile distilled water (SDW) and dimethyl sulfoxide (DMSO) (2.5% V/V) were used as negative controls. Values in bold differ significantly from controls (SDW and DMSO).

Detection of antimicrobial cross-resistance

Table 3. MIC values for products against antimicrobial resistant and susceptible *C. difficile*.

Isolate ID	Phenotype	Treatment								
		Fresh garlic extract (% v/v)	Garlic clove powder (mg/ml)	Allicin (mg/ml)	Leptospermum honey (MGO 514+) (% w/v)	Peppermint oil (% v/v)	Cinnamon root powder (mg/ml)	Menthol (mg/ml)	Zingerone (mg/ml)	<i>trans</i> -cinnamaldehyde (% v/v)
ESP123	MLS _B /tetR	0.4	9.4	9.4	8	4	75	4.7	9.4	0.02
SQ383		0.8	9.4	9.4	8	8	75	9.4	9.4	0.03
SQ357	MLS _B	0.8	9.4	9.4	16	8	75	9.4	9.4	0.01
SQ463		0.8	9.4	9.4	16	8	75	9.4	9.4	0.02
SAP019	MLS _B /tetR	0.8	4.7	9.4	8	8	75	9.4	9.4	0.03
SQ317		0.8	9.4	9.4	16	8	75	9.4	9.4	0.03
ES0474	FQR/tetR	0.8	9.4	9.4	16	8	75	9.4	9.4	0.03
ES1046		0.8	9.4	9.4	16	8	75	9.4	9.4	0.03

R, resistant; S, susceptible; MLS_B, macrolide-lincosamide-streptogramin-B; tetR, tetracycline resistant; FQR, fluoroquinolone resistant.

CONCLUSIONS

Overall, this study provides a fundamental framework regarding the possible mechanism of action of natural occurring antimicrobials against *C. difficile*. The findings indicate that damage to the cytoplasmic membrane may contribute to the mechanism of action of several naturally occurring antimicrobials against *C. difficile*. Also, a lack of cross-over mechanisms of resistance between standard antibiotics and natural compounds are shown. Further studies are required to determine the efficacy of these compounds *in vivo*.



The time-kill assay showed a > 3 log₁₀ reduction in *C. difficile* counts by all five bactericidal compounds after 24 h. Peppermint oil at ≥ 1 × MIC resulted in a log₁₀ reduction of ≥ 3 against both log- and stationary-phase *C. difficile* after 24 h. A similar pattern of killing with a reduction in bacterial counts was observed with *trans*-cinnamaldehyde at almost all concentrations.

The ATP-leakage assay showed that all five bactericidal compounds at most concentrations significantly reduced the intracellular ATP after 1 h of incubation ($P \leq 0.01$). The extracellular ATP was increased significantly by all five antimicrobials at all concentrations after 2 h ($P < 0.001$).

Figure 2. Effect (mean ± SD) of five bactericidal products and comparators in the leakage of ATP from the *C. difficile* cells, strain NCTC 13366. Concentration of treatments used in this assay: Nisin 8 × MIC (4 µg/ml), vancomycin 8 × MIC (8 µg/ml), cinnamon root powder 8 × MIC (150 mg/ml), peppermint oil 8 × MIC (16% v/v), *trans*-cinnamaldehyde 8 × MIC (1.6% v/v), menthol 8 × MIC (18.8 mg/ml), zingerone 8 × MIC (37.5 mg/ml).

All five bactericidal compounds damaged the cell membrane in both cell permeability assays. Treatment with peppermint oil, *trans*-cinnamaldehyde, menthol and zingerone resulted in significant increases in OD₅₉₅ indicating protein leakage after 1 h of exposure compared to the untreated controls (SDW and DMSO) ($P \leq 0.04$). Treatment with all natural compounds at most tested concentrations resulted in a significant increase in propidium iodide fluorescence after 6 h of exposure ($P < 0.05$).

Inhibition of protein synthesis

Streptomycin sulfate and tetracycline were used as positive controls and both inhibited protein synthesis/prokaryotic translation. Other than streptomycin and tetracycline, *Leptospermum* honey (MGO 514+) was the only treatment that showed a reduction in RFU ratio compared to untreated controls (SDW and DMSO) with values ranging from 42.1% to 68.9%.

None of the compounds showed elevated MICs against antibiotic-resistant strains of *C. difficile* harbouring DNA gyrase mutations, or conjugative transposons carrying *ermB* and *tetM*, suggesting that antibiotic resistance mechanisms are not cross-protective for natural products.