

SUPPLEMENTARY DATA

The Influence of Cellulose-Type Formulants on Anti-*Candida* Activity of the Tyrocidines

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Peptide purification and characterization

Tyrocidine mixture (Trc mix) was extracted from commercial tyrothricin using an optimised diethyl ether (DEE) and acetone precipitation protocol [42] and subjected to RP-HPLC (Figure S1). The insolubility of tyrocidine analogues (Trcs) and solubility of the hydrophobic linear gramicidins (Grms) in DEE-acetone, due to their different hydrophobic properties, allowed for the separation of the two groups of peptides from the tyrothricin mixture. However, it has been observed that the DEE-acetone wash results in the loss of some Trcs analogues. Therefore, each complex (tyrothricin and the Trcs fraction) were subjected to UPLC-ESMS to compare the abundance of different peptides analogues present within the preparation (Figure S1). The UPLC-ESMS was performed based on an optimised method [42].

The DEE-acetone wash resulted in complete loss of minor analogues and significant loss of some of the major analogues present within the fraction despite the overall increase in purity with the removal of the Grms (Table S2).

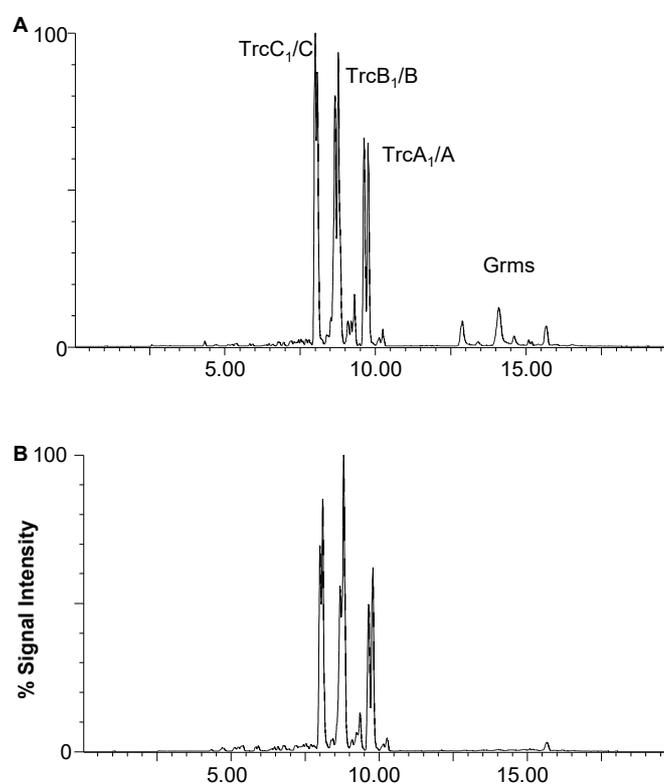


Figure S1: UPLC chromatograms of commercial tyrothricin complex with (a) the untreated tyrothricin, (b) the precipitate of the DEE:acetone wash containing the Trc mix.

Table S1: Percentage abundance and theoretical mass (mg) yield of the major Trc analogues present within the commercial tyrothricin complex before and after the wash with DEE: Acetone.

	%Abundance ^a		Theoretical mass (mg) yield of the major analogue ^c		%abundance Trc mix	
	Tyrothricin complex	Extracted Trc mix	Tyrothricin complex	Extracted Trc mix	Vosloo ²⁰	Troskie <i>et al.</i> ⁶
TrcC ₁	14.2	12.4	2.13	1.27	1.6	12.5
TrcC	13.6	15.0	2.04	1.53	10.7	14.8
TpcC ₁	1.1	1.2	0.17	0.12	0.3	-
TpcC	1.5	2.9	0.23	0.3	2.5	1.7
TrcB ₁	14.4	9.3	2.16	0.95	3.5	19.2
TrcB	15.2	22.9	2.28	2.34	9.4	18.2
TrcB'	3.9	-	0.59	-	8.5	4.0
TpcB	2.7	3.8	0.41	0.27	6.2	1.0
TrcA ₁	9.9	10.4	1.49	1.06	4.4	15.8
TrcA	9.6	13.0	1.44	1.32	21.1	12.9
TpcA	0.7	1.4	0.11	0.1	6.3	Trace
IGB	2.2	-	0.33	-	0.6	-
VGA	4.1	-	0.62	-	6.1	-
VGB	1.8	-	0.27	-	0.6	-
%Trcs ^b	86.8%	92.1%	13.05 mg	9.26 mg	81.8%	100.1%

^a % Abundance was calculated by expressing the peak area of each peptide as a percentage of the sum of the peak areas of all peptides present in the extract. It was assumed that the response factors of all peptides are similar due to their analogue structure.

^b % Trcs was determined by the sum of the peak areas of all the Trcs present in the tyrothricin complex and each of the extracts.

^c The mass (mg) of the predominant analogues present in the tyrothricin complex and each of the extracts was calculated by multiplying of % abundance by the total mass (mg) of in the tyrothricin complex and each of the extracts.

Activity of Trc mix and commercial antifungal compounds against *Candida albicans*

Table S2: Comparison of IC₅₀ and MIC values of selected antifungal compounds against planktonic cultures of *C. albicans* CAB1653. Tabulated IC₅₀ and MIC values represents the mean of 3–4 biological repeats and 12–32 technical repeats with SEM.

Drug or peptide	IC ₅₀ (μM)	MIC (μM)
Trc mix	11 ± 1.3	12.5–25
Caspofungin	4.8 ± 0.7	8.6 ± 1.1
Amphotericin B	13 ± 2.9	23 ± 4.2
Fluconazole	>325	>325

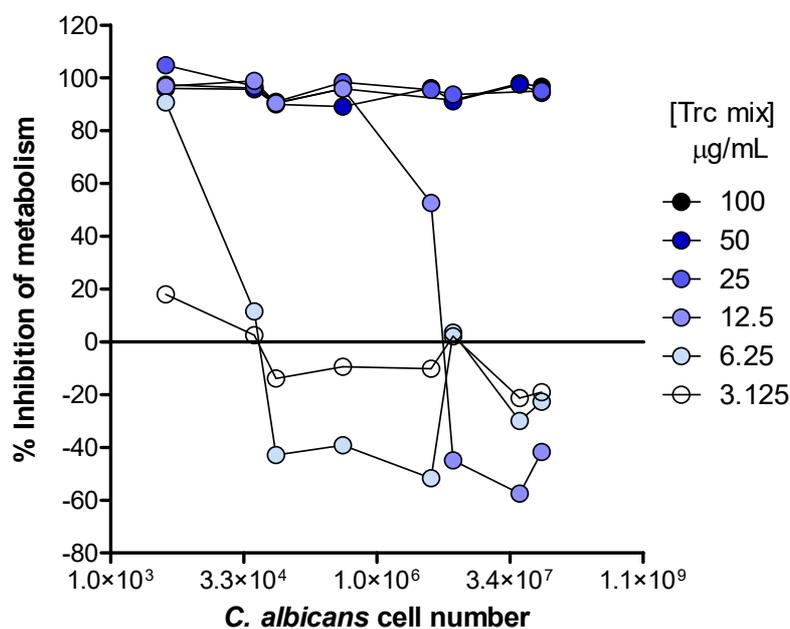


Figure S2: Correlation between metabolic inhibition of a range of *C. albicans* cell concentrations versus Trc mix concentrations. The values below 0% inhibition indicate higher conversion of the metabolic dye versus the control cultures. Each data point is the average determined for two cultures.

Statistical analysis of activity data

Table S6: Student t-test statistical comparison between different Trc mix of fresh (1 h) and matured (20 h) formulations of the observed inhibition parameters against planktonic cultures of *C. albicans* CAB1653. Tabulated IC₅₀ and MIC values (µg/mL) represents the mean of 3–4 biological repeats and 12–30 technical repeats with SEM. Unpaired Student t-test was done on each of the analysed pairs and only those with significant differences are shown.

Formulation at 1-hour vs 20-hours	IC ₅₀ ± SEM (µg/mL) 1 hour	IC ₅₀ ± SEM (µg/mL) 20 hours	P value
Control	11.4 ± 1.3	5.3 ± 0.4	0.0003
Trc mix:A4M (1:2)	8.9 ± 0.6	4.9 ± 0.2	<0.0001
Trc mix:A4M (1:4)	9.6 ± 1.3	5.6 ± 0.5	0.0196
Trc mix:E4M (1:2)	10.3 ± 0.8	6.8 ± 1.2	0.0143
Trc mix:E10M (1:2)	9.2 ± 0.3	6.2 ± 0.5	<0.0001
Trc mix:E10M (1:4)	7.0 ± 0.8	2.3 ± 0.7	0.0002
Trc mix:K15M (1:2)	8.9 ± 0.9	5.1 ± 0.3	0.0017

Formulation at 1-hour vs 20-hours	MIC ± SEM (µg/mL) 1 hour	MIC ± SEM (µg/mL) 20 hours	P value
Control	14.9 ± 1.5	7.5 ± 0.5	0.007
Trc mix:A4M (1:1)	14.0 ± 2.4	6.7 ± 0.5	0.0203
Trc mix:A4M (1:4)	14.8 ± 1.6	8.8 ± 0.9	0.0065
Trc mix:E10M (1:4)	10.1 ± 1.2	4.6 ± 0.8	0.0023

Table S7: One way Anova statistical comparison of IC₅₀ (µg/mL) value correlation between the different preparations of Trc mix (1 h vs 20 h). The analysed IC₅₀ values were the mean of 3–4 biological repeats and 12–30 technical repeats with SEM. One-way Anova with Bonferroni correlation test was done between each of the selected data sets.

		20 hours of maturation													
		1:4				1:2				1:1				Control	
		K15M	E10M	E4M	A4M	K15M	E10M	E4M	A4M	K15M	E10M	E4M	A4M		
1 hour of maturation	1:4	K15M	ns	<0.01	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns
		E10M	ns	<0.001	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns
		E4M	ns	<0.001	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns
		A4M	ns	<0.001	ns	ns	ns	ns	ns	<0.05	ns	ns	ns	<0.05	ns
	1:2	K15M	ns	<0.001	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns
		E10M	ns	<0.001	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns
		E4M	ns	<0.001	ns	ns	ns	ns	ns	<0.05	ns	<0.05	<0.05	<0.05	<0.05
		A4M	ns	<0.001	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns
	1:1	K15M	ns	<0.001	ns	ns	ns	ns	ns	ns	ns	<0.05	ns	ns	ns
		E10M	ns	<0.01	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns
		E4M	ns	<0.01	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns
		A4M	ns	<0.001	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns	ns
Control		ns	<0.001	ns	<0.01	<0.01	<0.001	ns	<0.01	ns	<0.01	<0.01	<0.001	<0.001	

